

**PROCEEDINGS OF
THE YOUNG SCIENTISTS FORUM SYMPOSIUM**

January 23, 2015

***National Science and Technology Commission
Young Scientists Forum***

The Young Scientists Forum Symposium

YSF SYMPOSIUM



JANUARY 23, 2015



Organized by

**National Science and Technology Commission
Young Scientists Forum**

EDITORS

Dr. LDB Suriyagoda

Dr. Indika Herath

Ms. Lohini Athiththan

Dr. Nilanthi Wijewardhana

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***Message from the Chairman
National Science and Technology Commission
Prof. Dhammika A. Tantrigoda***

It is with pleasure that I send this message on the occasion of the inauguration of the 4th Annual Research Session of the Young Scientists Forum (YSF) of the National Science and Technology Commission (NASTEC). The Annual Research Session of YSF provides a stage for presentation and discussion of latest scientific research findings of its membership. I am happy to note that 53 presentations covering a multitude of academic disciplines will be made at this year's YSF research sessions. I am sure YSF membership will make use of this opportunity to interact with each other and share their knowledge, experience and wisdom.

I would like to take this opportunity to show my gratitude to the hardworking YSF Steering Committee and to the dedicated NASTEC officials for organizing this important session. I hope the annual sessions will be a great success as in the last three years with active participation of all members of the YSF. I wish good luck for the proceedings of the sessions and all future endeavours of YSF.

***Message from the Director
National Science and Technology Commission
Dr. Muditha Liyanagedara***

I consider it as a privilege to give this message for the 4th YSF Symposium. I have been observing the development of this annual event since its inception in 2012. I am happy to say that its success is quite satisfactory. The demand for the symposium has been increasing every year and the standard of the symposium is also maintaining at a high level to give it to a due prestige.

NASTEC sponsors this event to promote a mutually beneficial research network among young scientists. General awareness of the research fields of each other, significance and relevance of them to one's research work may lead to collaborative research which would be more productive.

The message that I would like to give to the young scientists of this country is that it is high time for them to initiate research that brings direct economic benefits to the country. This should be started at the research planning stage. If the planning can be extended up to the identification of potential investors for the commercialization of their products/services, that would give more value to their research.

***Message from the Chairman
Young Scientist Forum
Dr. Chalinda Beneragama***

On behalf of the Organizing Committee, it is an honor and a privilege for me to welcome you to the Fourth Young Scientists Forum (YSF) Symposium of National Science and Technology Commission (NASTEC).

This symposium brings together researchers interested in the ever advancing state-of-the-art in the fields of Agricultural Sciences, Biochemistry, Biotechnology, Economics, Engineering, Environmental sciences, Health Sciences, Information Technology, Medicine, Microbiology, Molecular Biology, Physical Sciences and Social Sciences. High caliber meritorious contributions from researchers make this symposium a viable gathering platform of paramount importance for young scientists.

The committee received 82 submissions from all corners of the country which reflects the nationwide nature of this symposium. Each paper was peer reviewed typically by two specialized reviewers. Following the rigorous review process, you will find 53 extended abstracts appear in the symposium proceedings, corresponding to an acceptance rate of 64%.

The organization of the YSF Symposium is entirely voluntary. The review process required an enormous effort from the members of the YSF Steering Committee, and I would therefore like to thank all its members together with the reviewers for their contribution to the success of this symposium. I would like to express my sincere thanks and gratitude to Prof. Dhammika Tantrigoda, the Chairman of NASTEC and to Dr. Muditha Liyanagedara, the Director of NASTEC for their invaluable guidance and continual support for the symposium, being the main strength for the YSF. Ms. Asha Pitadeniya, Scientific Programme Manager of NASTEC and the Symposium Coordinator, deserves a very special word of appreciation for her tireless work with full of commitment. Moreover, I wish to thank the Editorial Board for their excellent team work in producing the Proceedings of the Symposium. The helping hand lent by all the staff members of the NASTEC is also gratefully acknowledged.

I strongly believe you will find the symposium productive, informative, and enjoyable. I wish the YSF Symposium all the very best in everything!

Forward by Editors

Dr. LDB Suriyagoda

Dr. Indika Herath

Ms. Lohini Aththithan

Dr. Nilanthi Wijewardhana

The Research Symposium of Young Scientists Forum (YSF) 2014 was intended to provide a platform for young scientists and professionals of various disciplines to exchange their knowledge and to discuss their new research findings. This symposium provides a great opportunity for them to present their work to a scientific audience and to obtain feedback. The YSF symposium is also an opportunity for networking among the scientists of different disciplines which is vital for future multidisciplinary research opportunities on current research needs.

For this year's Symposium, we received 82 extended abstracts from various disciplines. They were first screened by the editors, and 53 papers were selected through a double-blind review process. As per the decision of the steering committee of YSF 2014, the authors of all the selected papers were given an opportunity to present their findings at the symposium.

The editorial board wishes to thank all the authors for their valuable contribution to make this symposium a success. We are also thankful to the panel of reviewers for their valuable input through the review process. The editorial board would like to express their sincere gratitude to Professor D. A. Thanthirigoda, Chairman, National Science and Technology Commission (NASTEC) for his guidance extended to YSF throughout the year. We are very much thankful for Dr. Muditha Liyanagedara, The Director, NASTEC for his constant support. We would also like to express our sincere gratitude to other YSF steering committee members for their support and cooperation. Our very special thank is for Ms Asha Pitadeniya, and NASTEC staff members for all the support in organizing the symposium and preparation of proceedings. The symposium would not be a success without their dedication and hard work.

While congratulating all the authors who have been selected to present their research, we wish the YSF Symposium 2014 a great success.

**IN VITRO LIPASE, CHOLESTEROL ESTERASE AND CHOLESTEROL
MICELLIZATION INHIBITORY ACTIVITIES OF LEAF AND BARK OF
Cinnamomum zeylanicum Blume (CEYLON CINNAMON)**

W. P. K. M. Abeysekera¹, G. A. S. Premakumara^{1,*} and W. D. Ratnasooriya^{2,3}

¹*Industrial Technology Institute (ITI), 363, Baudhaloka Mawatha, Colombo 07*

²*Department of Zoology, Faculty of Science, University of Colombo*

³*Faculty of Allied Health Sciences, General Sir John Kothalawala Defence University,
Ratmalana, Sri Lanka*

Introduction

Cinnamomum zeylanicum Blume (Ceylon cinnamon) is indigenous to Sri Lanka and used as a spice in several countries. According to some Sri Lankan traditional physicians the bark and leaf of this plant is claimed to possess anti-lipidemic effects by inhibition of lipid digestion and/or absorption. Scientifically anti-lipidaemic properties of bark of cinnamon have shown in various *in vitro* and *in vivo* models. However, anti-lipidaemic properties of Ceylon cinnamon bark via lipase, cholesterol esterase and cholesterol micellization inhibitory activities were not previously reported. Moreover, anti-lipidaemic properties of leaf of Ceylon cinnamon is not Investigated to date worldwide. In this connection this study was initiated to investigate anti-lipidemic potential of bark and leaf of Ceylon cinnamon via *in vitro* lipase, cholesterol esterase and cholesterol micellization inhibitory activities.

Materials and Methods

Ethanollic and dichloromethane:methanol (DCM:M) extracts of bark and leaf Powdered 20 g of each alba grade bark and leaf were extracted into 200 mL of 95 % ethanol in a soxhlet extractor and 200 mL of DCM:M (1:1 v/v) at room temperature for 7 days. The extracts were filtered separately and evaporated under reduced pressure and freeze dried. Freeze dried ethanollic and dichloromethane:methanol (DCM:M) extracts of bark and leaves were used for analysis of anti-lipidemic properties.

Anti-lipidemic activity of bark and leaf extracts: Anti-lipidemic activity of bark and leaf extracts of Ceylon cinnamon were evaluated using *in vitro* lipase, cholesterol esterase and cholesterol micellization inhibitory assays.

Statistical analysis: Statistical analysis of the data were done using SAS version 6.12. One way analysis of variance (ANOVA) and the Duncan's Multiple Range Test (DMRT) were used to determine the differences among treatment means and $P < 0.05$ was considered as significant. Values are indicated as mean \pm SD.

* gasp@iti.lk

Results and Discussion

Significant differences were observed between bark and leaf extracts of Ceylon cinnamon for *in vitro* anti-lipase activity ($p < 0.05$) with, bark extracts having significantly high activity compared to leaf extracts ($p < 0.05$). Further, ethanol leaf and DCM:M leaf showed only 12.92 ± 1.37 and 19.71 ± 1.10 % inhibition at the highest studied concentration of $600 \mu\text{g mL}^{-1}$. The IC_{50} values of ethanol bark and DCM:M bark were 301.09 ± 5.73 and $297.57 \pm 11.78 \mu\text{g mL}^{-1}$ respectively. Moreover, the dose response relationship of ethanol bark and DCM:M bark are given in Table 1.

Table 1. Anti-lipase activity of ethanol and DCM:M bark extracts of Ceylon cinnamon

Concentration ($\mu\text{g mL}^{-1}$)	% Inhibition	
	Ethanol bark	DCM:M bark
37.5	55.27 ± 3.59	55.66 ± 3.07
75	49.54 ± 0.29	52.07 ± 1.96
150	27.35 ± 4.43	24.14 ± 3.11
300	5.30 ± 1.28	17.95 ± 5.72
600	5.74 ± 0.80	12.50 ± 1.32
$\text{IC}_{50} (\mu\text{g mL}^{-1})$	301.09 ± 5.73^a	297.57 ± 11.78^a

Data represented as mean \pm SD ($n=3$). Mean values in the column superscripted by different letters for bark extracts were significantly different at $p < 0.05$.

Both bark and leaf extracts of Ceylon cinnamon showed significant ($p < 0.05$) *in vitro* cholesterol esterase inhibitory activity in a dose dependent manner. However, bark extracts had significantly high activity compared to leaf extracts ($p < 0.05$). Moreover, ethanol extracts showed high activity than DCM:M extracts. The dose response relationship of bark extracts and leaf extracts are given in Figure 1 and Figure 2.

Both ethanol and DCM:M bark and leaf extracts of Ceylon cinnamon showed Cholesterol micellization inhibitory activity in a dose dependent manner. However, bark extracts had significantly high activity compared to leaf extracts ($p < 0.05$). The IC_{50} values of ethanol bark, ethanol leaf, DCM:M bark and DCM:M leaf were 0.23 ± 0.01 , 0.62 ± 0.01 , 0.48 ± 0.01 and $1.14 \pm 0.05 \text{ mg mL}^{-1}$ respectively. The dose response relationship of ethanol and DCM:M bark and leaf extracts for cholesterol micellization inhibitory activity is given in Table 2.

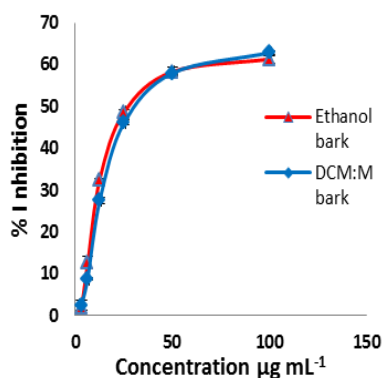


Figure 1. Cholesterol esterase inhibitory activity of ethanol and DCM:M bark extracts of Ceylon cinnamon. IC_{50} values: Ethanol bark $30.62 \pm 1.67^a \mu\text{g mL}^{-1}$; DCM:M bark: $34.39 \pm 0.91^b \mu\text{g mL}^{-1}$. IC_{50} values superscripted by different letters are significantly different at $p < 0.05$.

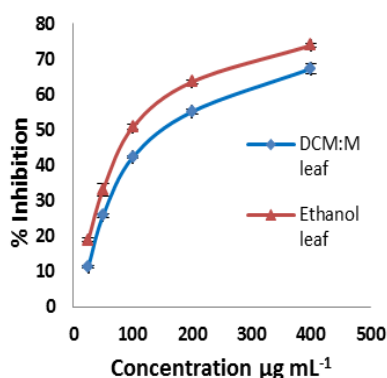


Figure 2. Cholesterol esterase inhibitory activity of ethanol and DCM:M leaf extracts of Ceylon cinnamon. IC_{50} values: Ethanol leaf $110.19 \pm 3.27^a \mu\text{g mL}^{-1}$; DCM:M leaf: $160.83 \pm 8.32^b \mu\text{g mL}^{-1}$. IC_{50} values superscripted by different letters are significantly different at $p < 0.05$.

Table 2. Cholesterol micellization inhibitory activity of bark extracts of Ceylon cinnamon

Concentration (mg mL ⁻¹)	% Inhibition of cholesterol solubility in micelles				
	Ethanol bark	DCM:M bark	Ethanol leaf	DCM:M leaf	EGCG
0.25	98.09 ± 0.56	73.94 ± 0.88	70.84 ± 0.80	43.59 ± 1.57	96.75 ± 1.08
0.5	69.48 ± 0.89	62.15 ± 1.06	44.54 ± 0.66	29.00 ± 0.61	69.78 ± 1.16
1	49.48 ± 0.85	19.36 ± 2.05	29.08 ± 1.11	12.83 ± 1.96	55.16 ± 0.58
IC_{50} (mg mL ⁻¹)	0.23 ± 0.01 ^a	0.48 ± 0.01 ^b	0.62 ± 0.01 ^c	1.14 ± 0.05 ^d	0.15 ± 0.01

Note: Data represented as mean ± SE (n=3). Mean values in the column superscripted by different letters for bark and leaf extracts were significantly different at $p < 0.05$.

We have previously reported anti-oxidant properties of leaf and bark extracts of Ceylon cinnamon. Leaf and bark extracts of Ceylon cinnamon had potent anti-oxidant activities via multiple mechanisms. Oxidative stress is now known to be involved in hyperlipidaemia; it is indeed an early event in the evolution of hyperlipidaemia. As free radicals are involved in lipid peroxidation and related hyperlipidaemic activities anti-oxidants can play a vital role in antilipidaemic activities. It has been reported that phenolic compounds show the ability to inhibit the formation of cholesterol micelles. Therefore, observed antilipidaemic activities of leaf and bark extracts of Ceylon cinnamon may be

due to the presence of anti-oxidative compounds. Further, experiments are in progress to isolate active compounds and efficacy in *in vivo* studies.

Conclusions and Recommendations

It is concluded that both bark and leaf extracts of Ceylon cinnamon possess lipase, cholesterol esterase and cholesterol micellization inhibitory activities. Bark had higher activity compared to leaf for all the above bioactivities. This is the first study to report lipase, cholesterol esterase and cholesterol micellization inhibitory activities of bark and leaf of Ceylon cinnamon worldwide.

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SEX HORMONE PROFILES AND ANTHROPOMETRIC PARAMETERS OF POST MENOPAUSAL BREAST CANCER PATIENTS

H.M.K Akalanka^{1*}, S. Ekanayake¹ and K. Samarasinghe²

¹*Department of Biochemistry, Faculty of Medical Sciences,
University of Sri Jayewardenepura*

²*Department of Pathology, Faculty of Medical Sciences,
University of Sri Jayewardenepura*

Introduction

Breast cancer (BC) is the most common carcinoma among women in Sri Lanka. According to the Cancer Registry 2007, the incidence of BC in Sri Lankan women is increasing since past few decades and the crude BC incidence is 19 per 100,000 population. It is found that mammary gland proliferation is increased with oestrogen, thus is reported to be associated with risk for BC (Enderson and Feigelson 2000; Russo et al 2000). Studies have shown that testosterone is found to enhance mammary tumor growth. Nevertheless progesterone and BC risk is grossly studied but results remain controversial. It is also reported that post-menopausal women being overweight and obese have the risk of BC due to adiposity related increase in levels of endogenous oestrogen concentrations (Bianchini et al. 2002; Lahmann et al. 2004).

Thus in the present study, serum sex hormone concentrations of newly diagnosed post-menopausal BC women were determined and data correlated with anthropometric parameters of each patient as no related data among Sri Lankan BC women exist. This study was conducted to (i) determine serum estrogen, testosterone and progesterone concentrations of newly diagnosed post-menopausal BC patients, (ii) measure/calculate anthropometric parameters (weight, height, body mass index (BMI), waist circumference (WC), hip circumference (HC), waist: hip ratio (WHR) mid upper arm circumference (MUC)) of BC patients and (iii) study possible correlations between serum estrogen, testosterone and progesterone concentrations and anthropometric measures.

Materials and Methods

Consent was obtained from newly diagnosed post-menopausal Sri Lankan BC patients (n=75) from National Cancer Institute Maharagama for the participation of the study. Serum total 17- β estradiol, total testosterone and progesterone concentrations were measured using an enzyme immunoassay

* kasuniakalanka@gmail.com

competition method with final fluorescent detection methodology with immunoanalyzer (VIDAS Progesterone Ref 30409 assay kits, VIDAS Testosterone Ref 30418 assay kits and VIDAS estradiol II Ref 30431 assay kits (Biomerieux, France).

Weight, height, BMI, WC, HC, MUC and WHR of each BC patient was measured/calculated. BMI ≥ 23 and >25 were considered as overweight and obese respectively. Over 80 cm of WC and WHR of ≥ 0.80 were considered as risk category and significances were analyzed.

Descriptive statistics, K independent sample test (Kruskal Wallis) and Spearman correlations were determined using the statistical software SPSS version 16 (Ethical approval Number- 651/12).

Results and Discussion

Mean age of menopause of the study population was 50 (± 3) years and mean (\pm SEM) testosterone, estrogen and progesterone concentrations of the study sample were 0.22 ng mL^{-1} (± 0.16), 27.8 pg mL^{-1} (± 0.52) and 0.57 ng mL^{-1} (± 0.15), respectively. The average hormone concentrations according to the age category are stated in Table 1. Accordingly, 43 %, 40 % and 17 % of BC women were present in each age category respectively.

Table 1. Testosterone, estrogen and progesterone concentrations of BC women according to the age

Age Category (Years)	Testosterone \pm SEM (ng mL^{-1})	Estrogen \pm SEM (pg mL^{-1})	Progesterone \pm SEM (ng mL^{-1})
51-60 n= 32	0.25 ± 0.03 (0.09-0.61)	29.69 ± 9.24 (0.89-210)	0.52 ± 0.13 (0.24-3.60)
61-70 n=30	0.17 ± 0.02 (0.09-0.64)	23.54 ± 7.19 (0.89-183)	0.27 ± 0.01 (0.24-0.63)
71-79 n= 13	0.19 ± 0.03 (0.09-0.43)	12.25 ± 1.24 (8.9-22.4)	0.51 ± 0.20 (0.24-2.87)
Reference range	0.1-0.9	< 58	< 0.41

Among the total group, 88 % had testosterone below half of the recommended upper value (0.9 ng mL^{-1}) and among them, 26 % had levels below the lower reference margin (0.1 ng mL^{-1}). None of the participants had testosterone above the recommended upper value. Twenty two percent of the study group had progesterone above the upper limit (0.41 ng mL^{-1}). From the total group, 47 % had progesterone below half of the recommended upper limit. Only 9 % had oestradiol II above 58 pg mL^{-1} (upper limit of normal). From the total group, 87 % had oestradiol II below half of the recommended upper

value. Significant differences in serum hormone concentrations with respect to age was not observed ($P>0.05$).

Among the patients, 64 % had BMI greater than 23 and 40 % among them were obese. According to WC, 67 % of women were in the risk category. With respect to WHR, 90 % belonged to the risk category. The mean values of anthropometric parameters are given in Table 2.

Table 2. Anthropometric parameters of BC patients

Anthropometric parameter (Unit)	Mean	Minimum	Maximum
Weight (kg)	56.0± 1.41	34.00	90.00
Height (m)	1.51± 0.01	1.38	1.70
BMI (kg m ⁻²)	24.8± 0.50	17.00	35.60
Waist (cm)	84.0± 1.50	33.00	116.00
Hip (cm)	91.0± 2.70	0.88	121.00
WHR	0.87± 0.60	0.76	0.99
MUC (cm)	28.4± 1.80	21.00	40.00

Oestrogen showed significant positive associations with BMI($r=0.3$, $p=0.02$), MUC ($r=0.4$, $p=0.00$) and weight ($r=0.4$, $p=0.00$). Progesterone showed significant positive associations with BMI($r=0.3$, $p=0.02$), MUC ($r=0.3$, $p=0.03$) and weight ($r=0.3$, $p=0.01$), even though oestradiol II was closer to lower limit of normal in most of the individuals. However, testosterone showed a significant association($r=0.28$, $p=0.02$) only with BMI. WC, HC or WHR were not significantly associated with oestrogen, progesterone or testosterone.

Even though testosterone and oestrogen are reported to be positively associated with BC, majority of participants had estrogen and testosterone below half of the recommended upper value while one fifth of the study population had elevated serum progesterone concentrations.

Conclusions and Recommendations

Majority of the patients were either overweight or obese and had low oestrogen which may have contributed to the high BMI, WC and WHR irrespective of the positive correlations observed with oestrogen.

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SOCIO-CULTURAL CHANGE IN MEEMURE VILLAGE IN KANDY DISTRICT: TWENTY YEARS AFTER THE PROHIBITION OF CHENA CULTIVATION

D.T.H. Ananda^{1*} and C.A.D. Nahallage¹

¹Department of Sociology and Anthropology, University of Sri Jayewardenepura

Introduction

In the recent past many cultures around the world have undergone extensive changes. Meemure village is an interesting example that has continuously undergone changes in the last two decades. Meemure is traditionally an agricultural village with a population of about 331 people, which has its own specific culture as a result of being isolated from main stream Sri Lankan society for a long time and some of the unique cultural traits associated with Meemure cannot be seen elsewhere in the country. Mythical stories related to king Rawana have dignified the pre-history of this village and have disclosed that the villagers are descendants of king Rawana and the Yaksha tribe. Meemure villagers have developed specific cultural traits from the beginning. Religious practises and rituals, food customs, Chena cultivation practises, paddy cultivation practises are among the unique traditional systems they developed and maintained. However they were confronted with a huge problem after the government prohibited their Chena cultivation practice in 1988. Various cultural practices related to kinship and marriage, economy, religion and traditional knowledge etc. are entwined together to form one fully integrated cultural system. When one or more traits get disrupted it affects the whole cultural system. Other factors that contributed to the change in the Meemure culture are the increase in the assimilation with outside society, the exodus of the young generation, a decline in the practise of the traditional knowledge system, innovation, devolution and forcible change: forced to adapt only to paddy cultivation. These factors cause, misery and community degradation which is colloquially known as a "Culture Crash". Some cultures cannot survive after being exposed to cultural changes. People of the Meemure village underwent this phase. Even though some of these cultural changes could be beneficial and adaptive, it might still be difficult for individuals within that particular culture to accept them. Thus cultural change is considered as a social problem even though it is a part of the necessary process of adaptation. The main objective of this study was to discover the consequences of the prohibition of Chena cultivation toward Meemure village and its unique culture.

Materials and Methods

* tharakaananda@hotmail.com

The study was carried out from April to November 2013. Questionnaire surveys were conducted to gather information about the cultural history and synchronic cultural traits on the Meemure village.

The population of the Meemure village consists of 331 people including 115 families; 40 families were randomly selected for the questionnaire with the aim of including individuals from families who have been living in the village for many generations. Five elderly individuals who were highly knowledgeable about their cultural heritage were selected for in-depth interviews. Observation and participant observation methods were used for comparison of cultural practises.

Study Area: Meemure village is situated in Kandy district; Ududumbara Secretariat Division, belonging to the Meemure Grama Niladari Division [N 07.433330 and E 08.833330]. It is a 5 km² village in the middle of the Knuckles Conservation Forest. From Colombo to Meemure village it is 229 km. There are no electricity facilities in Meemure village.

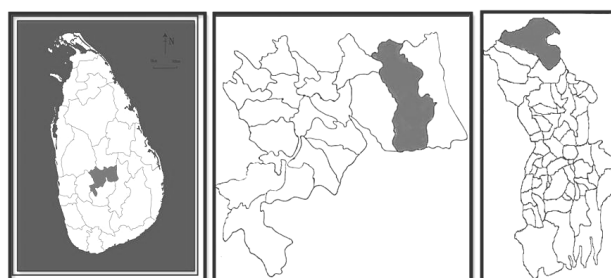


Figure 1. A) Kandy District, B) Ududumbara secretariat division, C) Meemure Grama Niladari Division (highlighted)

Results and Discussion

The single most significant factor that contributed to the change in their culture is the prohibition of Chena cultivation in 1988. The National Heritage Wilderness Area Act in 1988 defined areas above 3500 ft. as protected areas which included their hereditary Chena cultivating lands. It terminated their self-sustaining economy and consequently these people had to look outside their village to find necessary resources that are now not available in the village. When they were under the power of traditional political authorities they had not such limitations because the traditional leaders knew the link between man and the nature excessively.

During the time when villagers depended on Chena and Paddy cultivation they had enough food such as mun (green gram-*Phaseolus aureus*), thala (sesame), iringu (wheat), kurakkan (*Eleusine coracana*), undu (*Phaseolus mungo*) etc. Due to this self-sustaining life they hardly went out of the village for their daily

requirements. In those days they had to visit the nearest town of Kandy only about twice a year to get cloth, salt and other a few other basic requirements. In these visits they had taken their Chena harvest, orange (*Citrus sinensis*), honey, areca nuts (*Areca catechu*), and betel to exchange for other goods or to sell. At present they have to find other ways to get these daily requirements such as vegetables and grains. Therefore the villagers travel 38 km to reach the nearest town of Hunnasgiriya to meet their daily needs. It opened the doors to the outside world on a regular basis and villagers started to settle in these areas. As a result, more marriages occurred between Meemure people and people from outside villages. In addition villagers have sold their village lands for cheaper price and have settled in urban areas. Assimilation has increased by 52 % among three generations. From the sample size of 40 families including 148 children; only 43 % of them are remaining in the village and 57 % have migrated to urban areas for work, marriage or to educate their children.

Exodus of young has become a major problem in the village. Lacks of facilities in the village have made young people to migrate from the village for seeking job opportunities elsewhere. During the earlier times collective agrarian lifestyle supplied their daily needs. However at present they have to find everything individually for their own survival. Thus the decision of young people to migrate out of the village to support themselves cannot be halted.

The traditional knowledge system prevailing in the society is now limited only to the older generation as new generations do not practice or benefit from that knowledge. Thus their traditional cultural knowledge does not get passed down to the next generation through socialization. The traditional knowledge and community relations have been disturbed and these factors have produced individuals who know very little about their traditions or history.

As a result of the prohibition of Chena cultivation, Meemure villagers are practising only paddy cultivation and have absorbed modern farming methods. Many facets of Meemure life and their cultural traits have changed and been lost without replacement. Acceptance of new innovation leads to the loss of older ones. Some aspects of Chena cultivation such as traditional *chena pala* (shack), *pendi weta* (stockade around the chena), *vee atuwa* (traditional paddy storing system different from other areas), and food storing methods such as *iringu aduththa*, *iringu uga* (buck wheat storing systems) are unique cultural traits which are not being practiced by modern day Meemure villagers (Ananda and Nahallage 2014).

Religious practises such as Yakkama and Adukku Pujawa are specific characteristics in Meemure culture that are slowly diminishing. Meemure villagers continue to practice their religious ceremonies despite the fact that

the farming practices which they were built upon no longer exist. Also television and radio has replaced the traditional gathering of the villagers into a nearby house for discussion or entertainment at night or during other free time; 86 % of the families use radios, 44 % televisions and 35 % have telephones. Solar-cells are used by 88 % of the families in the village.

The traditional embalming methods that they used to postpone the putrefaction of the body such as honey, betel and arecanut mixture and mee (*Madhuca longifolia*) leaves, seeds, barks and tumeric etc. These practices are specific to them and are no longer practised since they rely on the modern methods used by the main society.

There is a new tendency among the villagers to let their school age children stay in urban temples until they finish schooling, in this way they can attend urban schools which have far more facilities than the village school does. Meemure primary school is older than 100 years. In 2011 there were only 12 children in the school with 3 teachers and classes up to grade 5. This represents 24 % of the school going children in the village. Seventy six percent of the children are going to nearby schools in Kaikawala (2 km from the village), Hunnasgiriya (38 km) and to schools in Kandy (116 km). As a consequence the majority of the people remaining in the village are of the older generation and the children below five years of age.

Conclusions and Recommendations

As well as outside influence, materialistic and technological factors have forced a number of cultural changes upon Meemure people. Many of these changes have arisen as a result of alterations in the Meemure traditional agrarian society. Most extreme cases of cultural collapse occur as a result of displacement of traditional political authority by the conquerors who know nothing about the culture they control. Eventually as anthropologists our duty is to document these specific cultural practices for future generations before it altogether diminishes from our society.

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AN ANALYSIS OF AIR POLLUTION IN SELECTED CITIES OF SRI LANKA USING MULTIVARIATE CONTROL CHARTS

D. N. S. Attanayake^{1*} and R. A. B. Abeygunawardana¹

¹*Department of Statistics, University of Colombo*

Introduction

Since the world began, decaying vegetable matters, dead animals and products of forest fires had produced noxious gaseous to the environment indicating there was no truly unpolluted environment ever. Thus, the matter of air pollution came into existence with the technological development of the human. Air pollution can be introduced as a silent hazard compared to the other hazards, since it is invisible and can be classified as Indoor and Outdoor pollution where in this study only outdoor air pollution was considered. Air pollutants are the chemical substances that are harmful to human and other living beings.

Even though Statistical Process Control (SPC) is a well practiced and widely spread tool of quality monitoring, application of SPC to monitor air quality is not frequent. On the other hand applications of SPC to monitor air quality in Sri Lanka are even harder to find. Two scientists Corbett and Pan emphasize that “applying SPC tools to environmental monitoring has substantial potential, whether for air, water or other emissions”. Additionally, many scientists and organizations have researched and revised the quality of the air surrounded Colombo city, the commercial capital of Sri Lanka, a few attempts have been made to have a look on the quality of air of the cities other than Colombo. Thus, through this research the eminence of the atmosphere of few other locations in addition to Colombo was investigated using SPC techniques.

Materials and Methods

The concentrations of the five criteria air pollutants; Carbon Monoxide (CO), Sulphur Dioxide (SO₂), Nitrogen Dioxide (NO₂), Ozone (O₃) and Particulate Matter (PM 10) were used to evaluate the quality of air surrounds the cities Colombo (Townhall area), Kandy, Maharagama and Kurunegala. Concentrations were measured in mg m⁻³ during 1.00 p.m. to 4.00 p.m. and the study period was October, 2010 to April, 2013.

The statistical approach used to evaluate the quality of air is control charts. A control chart is a graphical representation of a quality characteristic that has

* sandarekha89@yahoo.com

been measured over time. Main features of a control chart are the center line which represents the average value of the quality characteristic at in-control state, upper control limit and the lower control limit.

For the individual evaluations of the process level both Individual and Moving Range (I-MR) charts and Exponentially Weighted Moving Average (EWMA) charts were constructed. I-MR chart represents an individual chart together with a moving range chart enabling easier interpretation of both charts and is a control chart for variables. On individual chart each data point is plotted as a separate point and on moving range chart difference between two successive data points is plotted as they come from the process in a sequential order. The EWMA chart can be used instead of Shewhart control chart when the objective is to detect small shifts in the process level and variability.

Multivariate charts were employed as the air pollutants need simultaneous process monitoring. The multivariate counter parts of the above univariate charts, Hotelling T^2 and Generalized variance chart and Multivariate EWMA respectively were employed depending on the correlation of different air pollutants. Both univariate and multivariate EWMA (MEWMA) charts were used due to their sensitivity of notifying small process shifts and insensitivity to normality assumption of the variables.

Results and Discussion

According to I-MR charts CO concentrations at each location were in-control and SO₂ process was out-of-control only at Kurunegala.

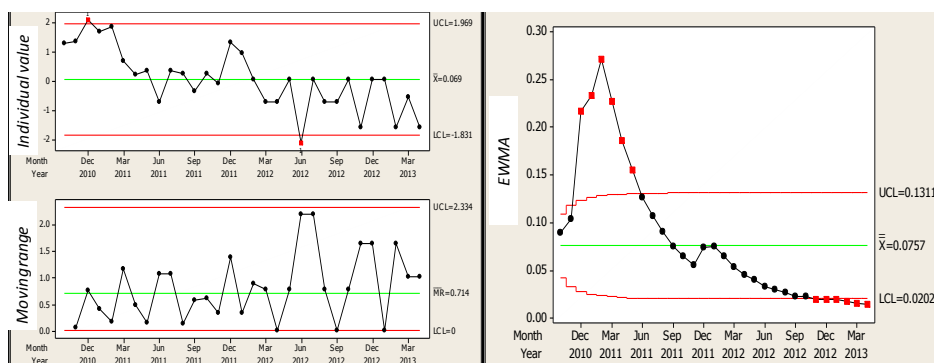


Figure 1. I-MR chart and EWMA chart of NO₂ at Colombo

Even though the I-MR chart of CO concentrations at Maharagama was in-control, EWMA chart depicted an out-of-control process. Contradictory to I-MR charts, EWMA charts of SO₂ depicted out-of-control processes at Colombo, Kandy and Kurunegala and a decline in the process level was practiced. A

decrement in the process mean was experienced in EWMA charts of NO₂ after February, 2011 at all the locations (Fig. 1). Both I-MR and EWMA charts illustrated PM10 concentration at Colombo were uncontrolled. Though I-MR charts were not applied to the O₃ data (since the data are not normal), considering the robustness to the normality assumption, EWMA charts were constructed. Based on the obtained figures it was concluded that all the processes are out-of-control and the out-of-control alarms appeared at the beginning of the process.

Foremost disadvantage of applying univariate control chart is the omission of relationships among the variables. The remedy used to overcome this problem was the application of multivariate process monitoring and controlling techniques.

Hotelling T² chart is the multivariate counterpart of Shewhart control charts. Application of the T² and variance charts to monitor the process shifts of CO and O₃ at Colombo revealed that the combined process is out-of-control and this result was proven by MEWMA chart (Fig. 2). Even though the Hotelling T² charts constructed to monitor the joint process of SO₂ and NO₂ concentrations at Kandy and Kurunegala were in-control, MEWMA charts depicted out-of-control processes. Hence it is advisable to use MEWMA chart over Hotelling T² chart.

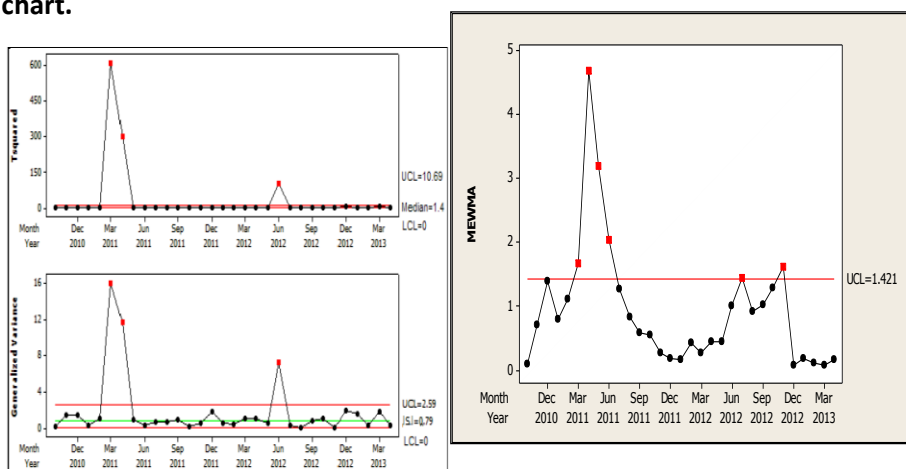


Figure 2. Hotelling T² chart and MEWMA chart for CO and O₃ at Colombo

Considering the correlations between the three gases CO, SO₂ and NO₂, Hotelling T² chart was applied to the three pollutants and the figure showed an in-control process. On the other hand MEWMA chart depicted an out-of-control process.

Conclusions and Recommendations

Based on the results it can be concluded that SO₂, NO₂ and O₃ concentrations were not at acceptable levels in the four locations selected due to the out-of-control processes observed under both I-MR and EWMA charts employed. PM10 concentration at Colombo and CO concentration at Maharagama were uncontrolled. Thus, the air at the four locations was polluted with respect to the five criteria air pollutants.

When appraising and controlling the quality of air environmental organizations should focus on the simultaneous process monitoring of SO₂ and NO₂; CO and O₃. Both multivariate and univariate EWMA control charts can be efficiently used in detecting moderate to small shifts in the process level and are successful as a phase II monitoring procedure. Shewhart type control charts (I-MR and Hotelling T² along with generalized variance chart) can be utilized in identifying large shifts in the process level. EWMA chart outperforms I-MR chart in appraising the quality of air due to the insensitivity to normality and efficiency in uncovering small shifts.

Concentrating on the sources of pollutants following suggestions can be proposed to reduce the concentrations in air.

- Implementation and maintenance of vehicle emission standards
- Quality of the fuel or gasoline should be continuously monitored and must be maintained in order to raise the quality of air
- Enhance the vehicle emission test facilities of CleanCo Lanka Limited
- Employment of a proper traffic controlling system
- Industries must pay attention on releasing the pollutants to the air in excessive amounts and government must play a role to secure the quality of air

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AN ALGORITHM TO FIND OPTIMAL ORDER QUANTITY WHEN PRICE DISCOUNTS ARE ALLOWED

M.M.R. Balasooriya^{1*} and D.M. Samarathunga¹

¹*Department of Mathematics, Faculty of Science, University of Peradeniya*

Introduction

Inventory control is one of the important areas in Production Management. The proper inventory control system helps in control of supply, storage and accessibility of items in order to ensure adequate supply without disproportionate over supply. It also aims to maximize the profit from the least amount of inventory without intruding upon customer satisfaction level.

Inventory control can facilitate the management to take advantages of the diverse discount offers as quantity discounts. Quantity discount are price reductions designed by suppliers to persuade bulk orders. Generally, in inventory control models, the cost function varies with ordered quantity. When there are price breaks, the cost function becomes a piecewise smooth, non linear, convex function over the quantity. Currently, there is a procedure to find optimal order quantity when price breaks are allowed. However it is not more efficient when there are several price breaks. Objective of this study is to develop an algorithm to solve such problems. First the inventory control problem is formulated with variable purchase price equivalently as a non linear, piecewise smooth, integer programming problem. Then an algorithm will be developed to find the optimal order quantity. Next computer program will be given to solve the mathematical model and finally to explore the applicability of the proposed method for a numerical example considered.

The model formulation

The unit purchase price depends on the quantity ordered as follows in a given inventory control model.

$$\begin{array}{ccc} \text{Quantity Purchased } (q) & \text{Unite Price } (p) & \\ q_{i-1} \leq q < q_i & p_i & \text{for } i = 1, 2, 3, \dots n. \end{array}$$

The total inventory cost = Purchasing cost + setup cost + holding cost.

$$TC(q) = \begin{cases} TC_1(q) = p_1 D + \frac{KD}{q} + \frac{Hq}{2} & \text{if } q_0 \leq q < q_1 \\ TC_2(q) = p_2 D + \frac{KD}{q} + \frac{Hq}{2} & \text{if } q_1 \leq q < q_2 \\ \dots & \dots \\ TC_n(q) = p_n D + \frac{KD}{q} + \frac{Hq}{2} & \text{if } q_{n-1} \leq q < q_n \end{cases},$$

* maduka.balasooriya@yahoo.com

where q = order quantity (number of units), D = demand rate (units per unit time), K = setup cost associated with the placement of an order (rupees per order), H = holding cost (rupees per inventory unit per unit time).

Results and Discussion

The above cost function is a piecewise smooth, non-linear, convex function. The problem (i.e., minimization of the total cost) can be equivalently written as follows:

$$\text{Min } z = \alpha_1 TC_1(q) + \alpha_2 TC_2(q) + \alpha_3 TC_3(q) + \dots + \alpha_n TC_n(q)$$

subject to

$$\alpha_1 + \alpha_2 + \alpha_3 + \dots + \alpha_n = 1$$

$$q \leq q_{i+1} \alpha_i + M y_i \quad \text{for } i = 1, 2, \dots, n.$$

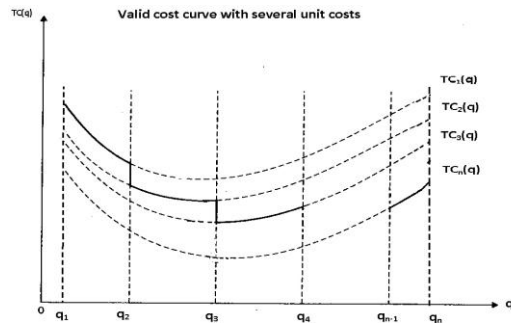
$$q \geq q_i \alpha_i - M y_i \quad \text{for } i = 1, 2, \dots, n.$$

$$y_1 + y_2 + y_3 + \dots + y_n = n - 1$$

$$\alpha_i + y_i = 1 \quad \text{for } i = 1, 2, \dots, n.$$

$$\alpha_i, y_i \in \{0, 1\} \quad \text{for } i = 1, 2, \dots, n.$$

$$q \geq 0, \quad M \text{ a big number.}$$



Since the above problem is a piecewise smooth, convex problem, the problem can be solved using the following algorithm.

Algorithm:

Step1: Set $\alpha_1 = 1$. Find $\left. \frac{\partial TC_1(q)}{\partial q} \right|_{q_1}$ and $\left. \frac{\partial TC_1(q)}{\partial q} \right|_{q_2}$. If both values are

positive, the optimal solution occurs at q_1 and find $TC_1(q)^* = TC_1(q_1)$. If both

$\left. \frac{\partial TC_1(q)}{\partial q} \right|_{q_1}$ and $\left. \frac{\partial TC_1(q)}{\partial q} \right|_{q_2}$ are negative, the optimal solution occurs at q_2

and find $TC_1(q)^* = TC_1(q_2)$. Else, the minimum cost occurs at \hat{q}_1 where

$\frac{\partial TC_1(\hat{q}_1)}{\partial q} = 0$. Calculate $TC_1(q)^* = TC_1(\hat{q}_1)$. Go to step 2.

Step 2: Repeat step 1 by fixing $\alpha_i = 1$ and find $TC_i(q)^*$ for each $i = 2, 3, \dots, n$.

Step 3: Among $TC_1(q)^*, TC_2(q)^*, \dots, TC_n(q)^*$ find the minimum value of $TC_k(q)^*$. The corresponding value q gives the optimal order quantity.

Computer programme:

Next, a computer programme has been developed for this algorithm using “R software”.

Case study:

```

minqfun <- function(q1,q2,K,D,p,l) # q1 < q2

{ A <- ((l*p)/2 -(K*D)/(q1^2)) B <- ((l*p)/2 -(K*D)/(q2^2))

  if(A> 0 & B >0) { return(q1)} else if(A< 0 & B <0)

{ return(q2)}Else {q <- sqrt((2*K*D)/(l*p)) return(q)}

cqfun <- function(q,p,D,K,l){ res <-p*D + (K*D)/q + (l*p*q)/2

  return(res)}

#Run from

q1 <- c(100,500,1000) q2 <- c(500,1000,1500) p <- c(1,0.8,0.6)

data <- cbind(q1,q2,p) D <- 2000 K <- 10 l <- 0.4 n <- 3

results <- matrix(NA,nrow = n,ncol=2) # q , cq for(i in 1:n){ results[i,1]
<- minqfun(data[i,1],data[i,2],K,D,data[i,3],l)

  results[i,2] <- cqfun(results[i,1],data[i,3],D,R,l)} alphai <-
which.min(results[,2]) minCq <- results[alphai,2] minq <-
results[alphai,1]minCq minq

```

In this example the optimal order quantity can be found when the annual demand is 2000 units and each order costs Rs. 1000 and annual holding cost is 40 % of unit cost. The unit costs depends on the quantity ordered as, Rs. 100 when quantities less than 500, Rs. 80 when quantities between 500 and 999 and Rs. 60 when quantities of 1000 or more.

Here, $D=2000$, K =Rs. 1000 an order, $H =0.4* p$ and p varies and is either Rs. 100, Rs.80, Rs.60.

When $\alpha_1 = 1 \Rightarrow y_1 = 0, y_2, y_3 = 1$, $q_1 = 100$ and $q_2 = 500$

Then $TC_1(q) = 2 \times 10^5 + \frac{2 \times 10^6}{q} + 20q$ and

$$\left. \frac{\partial TC_1(q)}{\partial q} \right|_{q_1} = 20 - \frac{2 \times 10^6}{100^2} < 0, \quad \left. \frac{\partial TC_1(q)}{\partial q} \right|_{q_2} = 20 - \frac{2 \times 10^6}{500^2} > 0.$$

Therefore minimum cost occurs when $\frac{\partial TC_1(q)}{\partial q} = 0$ at $\hat{q} = 316.23$ and

$$TC_1(\hat{q}) = 212,648.83.$$

When $\alpha_2 = 1 \Rightarrow y_2 = 0, y_1, y_3 = 1, q_2 = 500$ and $q_3 = 1000$.

Then $TC_2(q) = 1.6 \times 10^5 + \frac{2 \times 10^6}{q} + 16q$ and

$$\frac{\partial TC_2(q)}{\partial q} \Big|_{q_2} = 16 - \frac{2 \times 10^6}{500^2} > 0, \frac{\partial TC_2(q)}{\partial q} \Big|_{q_3} = 16 - \frac{2 \times 10^6}{1000^2} > 0.$$

Therefore minimum cost occurs at $q_2 = 500$ and $TC_2(q_2) = 172,000$.

When $\alpha_3 = 1 \Rightarrow y_3 = 0, y_1, y_2 = 1, q_3 = 1000$ and $q_4 = 1500$.

Then $TC_3(q) = 1.2 \times 10^5 + \frac{2 \times 10^6}{q} + 12q$ and

$$\frac{\partial TC_3(q)}{\partial q} \Big|_{q_3} = 12 - \frac{2 \times 10^6}{1000^2} > 0, \frac{\partial TC_3(q)}{\partial q} \Big|_{q_4} = 12 - \frac{2 \times 10^6}{1500^2} > 0. \text{Therefore}$$

minimum cost occurs at $q_3 = 1000$ and $TC_3(q_3) = 134,000$.

Then minimum costs at each category are $TC_1(q = 316.23) = \text{Rs. } 212,648.83$, $TC_2(q = 500) = \text{Rs. } 172,000$ and $TC_3(q = 1000) = \text{Rs. } 134,000$. Therefore, to achieve minimum cost, the company should buy 1000 items from the supplier with the cost of Rs. 134,000.

Conclusions and Recommendations

In this study a new method to find the optimal order quantity is proposed when there are price brakes. The propose model can be applied to solve inventory control model with variable purchase price. In general it can be applied to solve any unconstraint minimization problem with piecewise smooth, nonlinear convex integer programming problem.

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C-REACTIVE PROTEIN AND FIBRINOGEN LEVELS IN SMOKERS WITH CORONARY ARTERY DISEASE

E. M. S. Bandara^{1*}, S. Ekanayake¹, A. D. Kapuruge² and C. A. Wanigatunge³

¹*Department of Biochemistry, Faculty of Medical Sciences,
University of Sri Jayewardenepura, Nugegoda, Sri Lanka*

²*Cardio-thoracic Unit, Sri Jayewardenepura General Hospital, Nugegoda, Sri Lanka*

³*Department of Pharmacology, Faculty of Medical Sciences,
University of Sri Jayewardenepura, Nugegoda, Sri Lanka*

Introduction

Cigarette smoking is a major modifiable risk factor for the development of cardiovascular diseases. Smoking is associated with alteration in lipid profile, coronary vasomotor reactivity, platelet aggregation, and prothrombotic states. These are the main factors related to atherosclerosis. Cigarette smoking is also associated with increase in several inflammatory markers such as C-reactive protein (C-RP), interleukin -6 and tumor necrosis factors- α . These inflammatory responses are essential factors for initiation and the development of atherosclerosis. Since smoking has a fairly high prevalence in Sri Lanka, we assessed the CRP and fibrinogen concentrations of male patients awaiting Coronary Artery Bypass Graft surgery both smokers and non-smokers to study their association with disease severity score.

Materials and Methods

The study is a cross sectional descriptive study carried out at the Cardio-thoracic Unit of Sri Jayewardenepura General Hospital. The ethical approval was obtained from University of Sri Jayewardenepura (Approval No.635/12) and Sri Jayewardenepura General Hospital. There were 64 (age 56.9 ± 10) male patients in the study group. C-RP was determined using immuno-turbidometry assay and fibrinogen was determined by thrombin time. An interviewer administered questionnaire was used to collect data on smoking. Coronary Artery Disease (CAD) severity was evaluated by Gensini score (Gensini 1975). The Gensini score was computed by assigning a severity score according to the degree of luminal narrowing and geographical importance of each coronary stenosis as seen in the coronary angiogram. The assigned severity score according to the degree of luminal narrowing was multiplied by a factor considering the geographical importance of coronary artery (Table 1). The sum of all coronary arteries was expressed as the Genisini score. Mann-Whitney U test was used to analyse the result (SPSS version 16).

* ems.bandara@yahoo.com

Results and Discussion

The results of the study are presented in Table 2. Among the total study population more than 2/3 was smokers. Smokers had significantly high C-RP (U=320, $p < 0.05$) and fibrinogen (U=285, $p < 0.05$) level compared to non-smokers. The concentrations of both parameters of 20 -25 % of smokers were closer to the upper limit of the normal or higher. Previous studies have reported that increase in CRP and fibrinogen in smokers compared to non-smokers (Bazzano et al. 2003).

Table 1. Assigned score according to luminal narrowing and multiplying factor according to location of lesion

Degree of luminal narrowing %	Assigned severity score (a)	Geographical location of lesion	Multiplying factor (b)
30 – 50	1	Left main coronary artery (LMCA)	05
51 – 70	2	Proximal Left anterior descending (LAD) or proximal circumflex (LCX)	2.5
71 – 90	4	Mid LAD	1.5
91 – 99	8	Distal LAD, 1 st diagonal, mid LCX, distal LCX, obtuse marginal, proximal right coronary artery (RCA), mid RCA, distal RCA, posterior descending artery (PDA)	1.0
100	16	2 nd diagonal, posterolateral ventricular branch (PLV)	0.5

Note: Gensini score = $a \times b$

Previous prospective cohort studies revealed elevated CRP and fibrinogen associated positively with ischemic heart disease (Danesh et al. 1997). According to Tracy et al. (1997) Smoking is associated with dose-dependent and time-dependent increase in C-RP concentration in both men and women. However, there was no significant difference in C-RP and fibrinogen, irrespective of the number of cigarettes (> 5 per day or < 5 per day) smoked or the duration (> 10 years or < 10 years) in the present study. Also CRP and fibrinogen of smokers who have stopped smoking (>10 yrs) and who currently smoke were also not significantly different. This also emphasizes that duration and quantity had not affected C-RP and fibrinogen in this study group.

Although the association of cigarette smoking and development of cardiovascular diseases have been reported, the dose dependent correlation between risk of cardio vascular events and number of cigarettes smoked or the pack-years exposure were not confirmed (Price et al. 1999). We also noted that the individuals who had stopped smoking before 10 years also had comparatively lower values for the Gensini score than those who had stopped smoking < 10 years ago.

Table 2. Result of the CRP, fibrinogen and Gensini score of individuals

Smoking status	Frequency (%)	CRP concentration (mg L ⁻¹ , median-Inter quartile range)		Fibrinogen concentration (mgdL ⁻¹ , median-Inter quartile range)		Gensini score (mean ± SD)	
Smokers	67	5.2 (8.0)	P=0.0 2	302 (53)	P=0.0 2	48 (23)	P=0.9
Non smokers	33	2.8 (3.2)		273 (65)		52 (27)	
<i>From the individuals of smoking (n = 41), according to number of cigarette smoke per day</i>							
>5 per day	60	5.2 (8.0)	P=0.9	302 (60)	P=0.5	54 (24)	P=0.1
<5 per day	40	4.9 (9.6)		290 (84)		41 (22)	
<i>From the individuals of smoking (n = 41), according to number of years smoke</i>							
>10 years	79	5.2 (7.7)	P=0.9	309 (46)	P=0.1	50 (23)	P=0.5
<10 years	21	5.4 (9.4)		290 (72)		41 (26)	
<i>Individuals of smoked previously (n = 34), according to number of years cessation of smoking</i>							
>10 years	40	4.5 (6.7)	P=0.0 5	296 (131)	P=0.	35 (44)	P=0.07
<10 years	60	5.6 (20.4)		300 (30)		55 (42)	

Note: *The normal range for CRP and fibrinogen is < 6mg L⁻¹ and 150 – 400 mg dL⁻¹, respectively.

Conclusions and Recommendations

Smokers had significantly increased CRP and fibrinogen concentrations compared to non-smokers. There was no significant difference in CRP and fibrinogen, irrespective of the number of cigarettes (> 5 per day or < 5 per day)

smoked or the duration (> 10 years or < 10 years). Even though not significant those who had smoked for more than 10 years had higher Gensini score compared to those who smoked for <10 year. This indicate that the severity of the CAD is less in the individuals who stopped smoking before 10 years compared to individuals stopped smoking < 10 years ago. Even though this is not significant it indicates the severity of CAD is reduced if the period of smoking is less.

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**POTENTIAL ANTI-TUMOR COMPOUNDS FROM
Asparagus officinalis L.**

P.K.G.S.S. Bandara^{1*}, K. Takahashi² and K. Nabeta²

¹ *Department of Genetics and Plant Breeding,
Rubber Research Institute of Sri Lanka*

² *Division of Applied Bioscience, Research Faculty of Agriculture, Hokkaido University,
Kita-9, Nishi-9, Kita-ku, Sapporo 060-8589, Japan*

Introduction

Of the genus *Asparagus*, *Asparagus officinalis* is the most commercially important species and it has been used for a long time as an anti-cancer herbal medicine. Pharmacological studies on this plant have shown that a variety of biological activities such as, antitumor, anti-inflammatory, antifungal, antiviral and antioxidant. Among commonly consumed vegetables, antioxidant activity of asparagus, based on dry weight, has been ranked as the highest. Cancer is one of the most detrimental diseases and currently massive research works have been conducted all over the world to discover new anti-cancer agents from natural product sources. The potential of using natural products as anti-cancer agents was recognized in the 1950s by the U.S. National Cancer Institute (NCI) and has since made major contributions to the discovery of new naturally occurring anti-cancer agents. According to the international agency for research on cancer (IARC), which is a part of the world health organization (WHO), by the year 2030, there will be 20 to 25 million incident cases of cancer and 13 to 16 million cancer deaths annually. Currently, over 50 % of anti-cancer drugs have been isolated from natural sources. It is of paramount importance to explore novel naturally occurring anti-cancer agents from plant kingdom and therefore, this study was started with the objective of identifying potential anti-tumor active compounds from *A. officinalis*.

Materials and Methods

Cladophylls of *A. officinalis* were used for the extraction of antitumor active compounds. In the extraction process, air dried and powdered 600 g of cladophyll of *A. officinalis* was extracted with 500 mL of 70 % aqueous ethanol and evaporated to dryness. The resultant aqueous residue was thrice extracted with an equal volume of ethyl acetate. The organic layer and the aqueous layers were collected separately and the organic layer was dehydrated with Na₂SO₄ and concentrated under reduced pressure to afford fraction A (3.6 g). The aqueous layer was extracted with *n*-butanol to afford fraction B.

* sarathbandara2003@yahoo.co.uk

The fraction A was subjected to a silica gel column chromatography (SGC) with CHCl_3 - CH_3OH in increasing order of polarity to obtain 10 fractions (A1 to A10). The fraction A3 with comparatively higher anti-tumor activity was re-chromatographed over silica gel with *n*-hexane-EtOAc (80:20) to afford 9 fractions (A3.1 to A3.9). From fraction A3.3, compound 1 (18.2 mg) was purified by PTLC with *n*-hexane-EtOAc (70:30). Compound 2 (14.4 mg) was purified by PTLC with *n*-hexane-EtOAc (50:50) from fraction A3.6. The fraction A3.5 was further subjected to reversed phase HPLC (C_{18} silica column with 250 mm x 20 mm i.d.) with 90 % aqueous acetonitrile as the isocratic mobile phase at a flow rate of 5 mL min^{-1} to afford compound 3 (10.2 mg). The fraction B was thrice extracted with an equal volume of *n*-butanol and concentrated under reduced pressure to afford brown residue (4 g). It was then subjected to SGC with CH_3OH - CHCl_3 in increasing order of polarity to obtain 3 fractions (B1 to B3). The active fractions B1 and B2 were further separated by the reverse phase HPLC (C_{18} silica column with 250 mm x 20 mm i.d.) using 40 % and 70 % aqueous CH_3OH , respectively as the isocratic mobile phase at a flow rate of 5 mL min^{-1} to afford compounds 4 (7.4 mg) and 5 (40 mg), respectively. The structures of extracted compounds were elucidated based on one and two dimensional NMR and MS spectroscopy. Additionally, NMR data were compared with those reported in literature. The ^1H -NMR and ^{13}C -NMR spectra (500 MHz for ^1H -NMR and 125 MHz for ^{13}C -NMR) were generated using CD_3OD or CDCl_3 as solvents. Anti-tumor activity of each isolated compound was screened against HL-60 cells using MTT bioassay and IC_{50} values of isolated compounds were calculated as described previously. The statistical significance of antitumor activity of all the compounds was analyzed by *t*-test at $p < 0.05$.

Results and Discussion

Based on the results of the present study, compound 1 (phytol), 2 (phytene-1,2-diol) and 3 (betulin) were found to have a significantly higher anti-tumor activity compared to that of compound 5 (methyl (25S) protodioscin) (Fig. 1), which is a well-known anti-tumor active compound (Table 1). The inhibitory activity of phytol isolated from *Perilla* leaves on HT-29 human colon cancer cells, MG-63 osteosarcoma cells and AZ-521 gastric cancer cells have been described and as an anti-tumor component in *Scutellaria barbata* has been found to be phytol. However, it seems that studies on anti-tumor activity of phytol have not yet been carried out up to a satisfactory level. The compound 2 has slightly lower IC_{50} value compared to that of compound 1 (Table 1). This activity difference is likely to be due to the additional OH group present in the compound 2. This speculation was further supported by the results obtained by Hibasami et al. (2002) which explain that a higher apoptotic activity has been observed in diol and triol type of phytol against lymphoid leukemia Molt 4B cells indicating the number of OH groups might be an important criterion in

determining anti-tumor activity. During the extraction process of the present study, phytol has been obtained as the major component in quantity wise leading to determine that *A. officinalis* is a rich source for phytol. Other than phytol, *A. officinalis* might be considered as a rich source for methyl protodioscin since the amount of methyl protodioscin also seemed to be significantly high. The compound 3 (betulin) (Fig. 1) has also shown comparatively higher activity against HL-60 cells. Betulenic acid, a derivative of betulin, is a well-known compound which has various biological activities including anti-tumor activity as well.

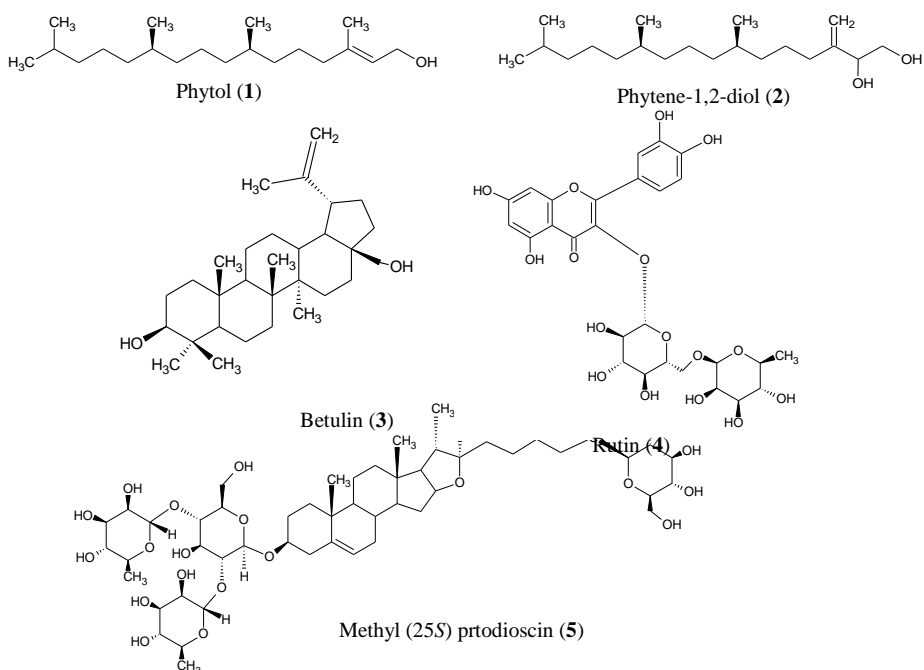


Figure 1. Structures of purified compounds.

The only difference between betulin and betulenic acid is the presence of carboxylic acid group at C-28 of betulenic acid. *A. officinalis* is well known as a rich source of flavonoids mainly rutin which is a strong antioxidant. Compound 4 (rutin) (Fig. 1), isolated from *A. officinalis* has not shown significant anti-tumor activity against HL-60 cells suggesting that it might not be a potent anti-tumor active compounds. However, it is a well-known flavonoid with high antioxidant activity. Among spirostanol type and furostanol type steroidal saponins isolated from *A. acutifolius*, only spirostanol type steroidal saponins have shown strong antifungal activity against pathogenic yeast. In the present study, the compounds 5 was identified to be furostanol type steroidal

saponins (methyl (25S) protodioscin) and found to have no significant antitumor activity against HL-60 cells (IC_{50} values $>100 \mu\text{g mL}^{-1}$) (Table 1).

Table 1. IC_{50} values of isolated compounds against HL-60 cells

Compound	IC_{50} value ($\mu\text{g mL}^{-1}$)
1	15.7
2	12.3
3	15.1
4	>100.0
5	>100.0

Conclusions and Recommendations

The compound 1 (phytol), 2 (phytene-1,2-diol) and 3 (betulin) are the potential anti-tumor active compounds available in *A. officinalis*. Further investigation should be performed to elucidate the specificity of anti-tumor activity of each compound against different cancer cell lines. Finally, *Asparagus* species available in Sri Lanka should be thoroughly investigated to identify potential novel anti-tumor active agents.

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WOUND HEALING ACTIVE CONSTITUENT FROM *Ficus racemosa* Linn. BARK

**N. S. Bopage¹, K. H. Jayawardena², S. C. Wijeyaratne³, A. M. Abeysekera⁴
and G. M.K.B. Gunaherath^{1*}**

¹*Department of Chemistry, the Open University of Sri Lanka, Nawala, Nugegoda*

²*Department of Zoology, the Open University of Sri Lanka, Nawala, Nugegoda*

³*Department of Botany, University of Sri Jayewardenepura, Nugegoda*

⁴*Department of Chemistry, University of Sri Jayewardenepura, Nugegoda*

Introduction

Ficus racemosa Linn. (*Ficus glomerata* Roxb.) is a large, deciduous tree which belongs to family Moraceae and exhibits wound healing activity. Wound is an injury that results in opening or breaking of the skin. The classical model of wound healing is divided into three sequential, yet overlapping phases: (1) inflammatory, (2) proliferate and (3) remodeling. Immediately after an injury, a fibrin clot is formed to limit the active bleeding (homeostasis). In the inflammatory phase, bacteria and debris are removed by phagocytic action. Proliferate phase is characterized by angiogenesis, collagen deposition, granulation tissue formation, epithelialization, and wound contraction. In the remodeling phase, collagen is remodeled and realigned along tension lines and cells. From ancient times plant preparations have been used to enhance wound healing though few of them have been either evaluated scientifically for their efficacy or studied for their mechanism of action. The present study was undertaken with the view of identifying potential wound healing constituent(s) of hexanes extract of stem bark of *F. racemosa* through bioactivity directed fractionation using wound healing assay (WHA).

Materials and Methods

Extraction and Fractionation of the Plant material: Dried powdered stem bark of *F. racemosa* Linn. (500 g) was sequentially extracted with hot hexanes, dichloromethane (CH₂Cl₂), ethyl acetate (EtOAc) and methanol (MeOH) and the solvents were removed under vacuum at 40 °C to obtain the respective extracts and were assayed for the wound healing potential. Among these four extracts, the hexanes extract was found to be the most active on wound healing assay (Table 1). The hexanes extract was fractionated using column chromatography on silica gel to give 11 major fractions. These fractions were subjected to wound healing assay and those showed significant wound healing activity (>75 %) were further fractionated/ investigated.

Establishment of cell cultures: Baby Hamster Kidney (BHK) cell cultures were established in the laboratory using standard *in-vitro* methods. At the

* kbgun@ou.ac.lk

confluence stage of the growth, cells were trypsinized and transferred to 12 well plates. Cells were maintained at 37 °C in a 5 % CO₂ humidified incubator until a monolayer is formed in Dulbecco's Modified Eagles Medium (DMEM) containing 5 % Fetal Bovine Serum [5 % growth medium (5 %GM)].

Wound Healing Assay (WHA): A scratch (wound) was performed along the vertical axis of each well under the microscope. The monolayer with wound was washed with 750 µL of phosphate buffer saline (PBS). Each test well was filled with 990 µL of DMEM and added 10 µL of DMSO containing appropriate amount of test sample. Two controls, 1 % DMSO in DMEM and 100 % DMEM were used in this experiment. In addition, asiaticoside (0.01 mg dm⁻³) was used as a positive control when pure compounds were assayed. Plates were incubated for 24 hours at 37 °C with 5 % CO₂. The widths of the wounds at different time intervals (0 h, 12 h, 18 h, and 24 h) were measured and the percentage wound closure was calculated. All the experiments were carried out in three replicates and three measurements were taken for each wound. Data were statistically analyzed using SPSS version 20.0. P-values of less than 0.05 were considered to be significant.

Results and Discussion

Hexanes and CH₂Cl₂ extracts of *F. racemosa* showed wound healing activity against BHK cells as shown in Table 1. Hexanes extract was selected for further fractionation in the present study.

Table 1. Weights of the extracts and activity of the different bark extracts of *F. racemosa* in WHA

Sample ^a	Weight of extract (g)	% Closure of the wound ^b at t = 24 h
Hexane extract	9.32	90.9(± 4.7)
CH ₂ Cl ₂ extract	1.35	78.1(± 7.0)
EtOAc extract	1.25	8.7(± 6.2)
MeOH extract	1.21	17.4(± 7.8)
1 % DMSO (Control 1)		15.5(± 8.5)
100 % DMEM (Control 2)		16.5(± 7.8)

Note: ^aSample concentration at a 0.1 mg dm⁻³

^bResults are given as mean values of three replicates with 95 % confidence level

Fractionation of hexane extract (2.5 g) by column chromatography over SiO₂ using solvent gradients containing Hexanes, Hexanes/CH₂Cl₂, CH₂Cl₂, and CH₂Cl₂/MeOH afforded 52 fractions which were combined according their thin layer chromatographic (TLC) patterns to give 11 fractions and subjected to WHA at a concentration of 0.05 mg dm⁻³ as described previously. The rate of

healing was calculated at 24 h from initial stage. Among these, fraction 4 showed highest activity (93.3 % wound closure at 24 h) while fractions 2, 3, 5 and 6 also showed high (>75 %) wound healing activity, which compelled further fractionation of these column fractions (Table 2).

Table 2. Weights of the column fractions and activity of the column fractions of the hexane extract of stem bark of *F. racemosa* in WHA

Sample ^a	Weight of fraction (mg)	% Closure of the wound ^b at t = 24 h
01	233.8	45.9 (± 5.6)
02	1314.8	90.0 (± 3.4)
03	19.7	87.6 (± 4.0)
04	5.1	93.1 (± 3.1)
05	20.5	84.1 (± 1.5)
06	8.8	76.7 (± 2.6)
07	103.6	12.7 (± 3.1)
08	157.7	17.0 (± 5.4)
09	94.5	10.0 (± 2.4)
10	289.3	13.0 (± 4.0)
11	31.5	21.5 (± 4.2)
1 % DMSO (Control 1)		7.9 (± 4.3)
100 % DMEM (Control 2)		9.4 (± 3.2)

Note: ^a Sample concentration at a 0.05 mg dm⁻³

^b Results are given as mean values of three replicates with 95% confidence level

Fraction 4 was found to contain one major compound which was isolated by preparative TLC and identified as lupeol m.p. 213 – 215 °C (lit. 215 °C) by comparison with an authentic sample (TLC, Co-TLC and mixed m.p.). Fraction 2 was found as oily fraction and contained five major compounds. One of the major compounds was identified as lupeol on TLC. This fraction (500 mg) was chromatographed over a column of silica gel solvent gradients containing hexanes, hexanes/CH₂Cl₂, CH₂Cl₂, and CH₂Cl₂/MeOH afforded yielding 59 fractions. The column fractions were pooled according their TLC patterns to give 13 major sub fractions F1–F13. The sub fraction F2 (25 mg) was contained three constituents. About 10 mg of this fraction further purification by repeated preparative TLC, α-amyrin acetate was isolated (2 mg), identity was confirmed by TLC, Co-TLC, m.p. (m.p. 224 – 225 °C) (Lit. 227 °C) and mixed m.p. with an authentic sample. The sub fraction F6 (32 mg) on further purification yielded white crystalline compound (29 mg), which was identified as lupeol acetate by TLC, Co-TLC. Its identity was confirmed by m. p. (m. p. 216 – 217 °C) (lit. 218 °C) and mixed m. p. with an authentic sample. These isolated compounds were subjected to WHA at a concentration of 0.01 mg dm⁻³ and

the percentage closure of wound at 24 h was obtained (Table 3). These results indicate that wound healing activity of lupeol is slightly higher than that asiaticoside. Investigation of fractions 3, 5, and 6 is underway.

Table 3. Activity of the isolated compounds of stem bark of *F. racemosa* in WHA

Sample ^a	% Closure of the wound ^b at t = 24 h
Lupeol	83.0(±9.4)
α- Amyrin acetate	22.8(±9.3)
Lupeol acetate	34.5(±8.5)
Asiaticoside (Positive control)	76.2(±7.5)
1 % DMSO (Negative control 1)	11.5(±10.0)
100 % DMEM (Negative control 2)	10.8(±10.7)

Note: ^aSample concentration at a 0.01 mg dm⁻³

^bResults are given as mean values of three replicates with 95% confidence level

Conclusions and Recommendations

In this study we have developed a rapid assay technique to study the potential of wound healing property of natural products and extracts. It is evident that compounds contained in the hexanes extract of stem bark of *F. racemosa* have potential wound healing activity. Among the compounds isolated from the hexanes extract of *F. racemosa*, lupeol is responsible for the enhancement of wound healing on BHK cells.

Acknowledgement

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IDENTIFICATION OF RESISTANT ACCESSIONS OF *Capsicum* spp. FOR BREEDING MEALY BUG RESISTANT CHILI: A GREEN HOUSE TRIAL

**M. D. M. Chamikara¹, M. Ishan¹, S. S. Karunadasa¹, V. D. W. Kasthuriarachchi¹,
R. V. Lelwala¹, M. K. D. I. Perera¹, P. I. Rajapaksha¹
and S. D. S. S. Sooriyapathirana^{1*}**

¹*Department of Molecular Biology and Biotechnology, Faculty of Science,
University of Peradeniya, Sri Lanka*

Introduction

Chili (*Capsicum* spp.), a vegetable crop grown in Sri Lanka, is an important flavouring constituent in meals. Approximately 750 million Rupees of GDP is contributed by chili annually. Mealy bug belongs to family Pseudococcidae is an insect pest in various crops such as chili, papaya, guava, banana and citrus which is a huge threat to Sri Lankan cultivations. Mealy bug infestations can be typically observed as cotton-like masses on the above part of the plants especially on fruits and flowers. Adult mealy bugs suck plant sap and inject toxic substances that lead to crooked fruits, chlorosis, plant stunting and early leaf and fruit drop. Infestation of this pest, leads to excessive application of pesticides by the farmers. Consequently these chemical pesticides cause health risks and environmental pollution. Breeding of resistant chili varieties seems the most promising solution to reduce excessive usage of chemicals for a developing country like Sri Lanka. Therefore the current study was conducted to assess the variation of a set of *Capsicum* spp. germ plasm of Sri Lanka towards the resistance to mealy bug infestation.

Materials and Methods

Seeds of 44 wild chili selections (accessions) were collected from a farm at Viskamgama in Rathnapura District, Sri Lanka (6° 32' 31.7" N 80° 22' 16.6" E) and grown in a greenhouse at University of Peradeniya (Yala season, 2014) with four replicates each. Five commercial chili varieties, MI-hot, MI-2, *Varaniya miris*, CA-8 and Bell pepper, were also grown along with wild accessions in four replicates. After three months of establishment, severe natural mealy bug infestation was observed at a varying degree on the greenhouse grown chili accessions and commercial varieties. Thus we took the advantage of characterizing these accessions for mealy bug resistance before applying control measures for the continuity of the project for other objectives of the study. A categorical scale (Category 1=No infestation, Category 2=Slight infestation, Category 3= Moderate infestation, Category 4= Severe infestation) (Fig. 1) was used to rank the individual plants for the degree of infestation by

* sunethss09@gmail.com

mealy bugs. The data were analyzed using ANOVA procedure in statistical package SAS 9.3.

Results and Discussion

There was a significant effect of the chili accession on the degree of infestation of mealy bugs ($P < 0.0001$) and $R^2 = 98.91\%$. Out of 49 accessions, seven (16 %) displayed no infestation (Class 1), 31 (63 %) displayed slight infestation (Class 2). Four (8%) were with moderate infestation (Class 3) and 6 (12 %) were with severe infestation (Class 4). All accessions that showed least infestation (Class 1 and 2) were wild collected ones. Out of five commercial varieties (MI-hot, MI-2, CA-8, Bell Pepper, *Varaniya miris*), MI-hot and MI-2, which are popular Sri Lankan chili varieties were in Category 4 undergone severe infestation (Fig. 2). Moreover bell pepper was in Category 3 and CA-8, a popular Sri Lankan pepper (curry chili) and *Varaniya miris* were in Category 2. This demonstrated a critical situation of the chili cultivation as most of the popular varieties are vulnerable to the mealy bug attack. Seven wild collected accessions showed complete resistance to the mealy bug attack and others showed partial resistance. These resistant accessions could be used as parents in future to develop mealy bug resistant chili varieties in breeding.

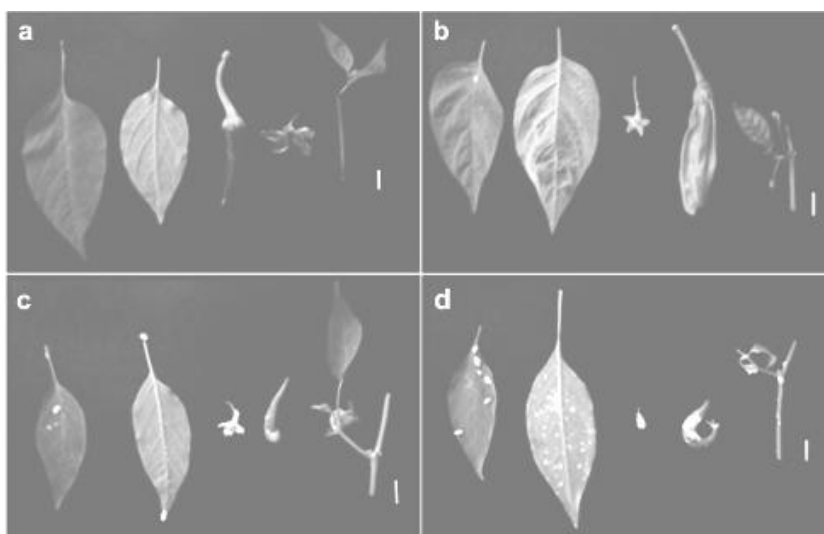


Figure 1. Severity classes of the *Pseudococcidae* sp (mealy bug) attack on the chili plant. a: Not infested under heavy presence in the surrounding (Severity Class 1); b: Slight infestation (Severity Class 2); c: Moderate infestation (Severity Class 3); d: Severe Infestation (Severity Class 4)(Scale bar= 1 cm).

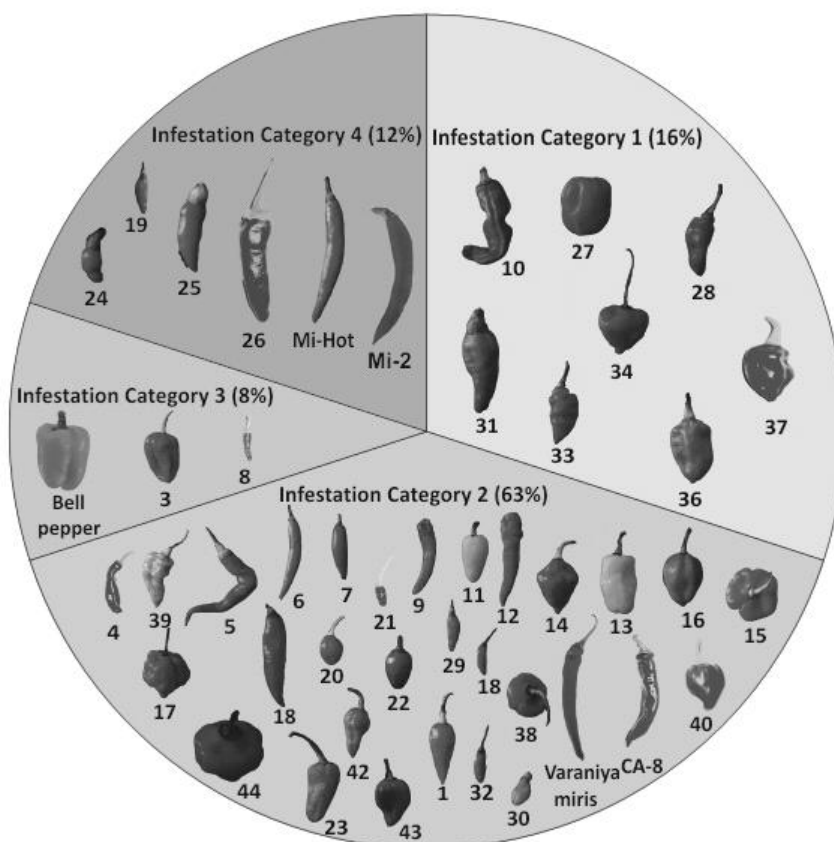


Figure 2. The pie chart showing the percentages of *Capsicum* accessions in each class of infestation severity (Fig. 1).

The total number of accessions was 44 and additionally five commercial chili varieties were also included. Inside the slices, a typical mature pod of each accession is shown. For the wild accessions collected, the numerical labels are assigned (indicated under each pod) for analytical purposes. The pods are not shown according to the scale.

Conclusions and Recommendations

Most of the commercial chili varieties grown in Sri Lanka are vulnerable to mealy bug attack. Some of the wild chili accessions are a good source of resistance and can be used in future as parents to develop mealy bug resistant chili varieties in breeding. Plans are also underway to screen the studied germplasm for mealy bug resistance under diverse field conditions. Moreover,

the segregating populations can be produced by crossing mealy bug resistant and sensitive accessions, to dissect the underlying genetics of the tolerance. Detailed Quantitative Trait Loci (QTL) mapping procedure can be carried out to tag the candidate genes and linked molecular markers to the resistance. These molecular markers can be used to expedite the chili breeding in Sri Lanka. Further to the discovering of the genetic basis of mealy bug resistance, the same set of germplasm is also currently being used by our group to characterize the genes related to pungency, fruit size and shape as they are the most important quality traits in chili.

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**EXTRACTS OF *Alpinia calcarata* (LESSER GALANGAL) INHIBITS
PRODUCTION OF INFLAMMATORY MEDIATOR-NITRIC OXIDE IN RAW
264.7 MURINE MACROPHAGES**

M. Chandhrakanthan¹, S. Kathirgamanathan^{1*}, S. M. Handunnetti²
and G. A. S. Premakumara¹

¹*Industrial Technology Institute, Colombo, Sri Lanka*

²*Institute of Biochemistry, Molecular Biology and Biotechnology,
University of Colombo, Sri Lanka*

Introduction

Inflammatory diseases such as rheumatoid arthritis (RA) are associated with increased production of nitric oxide (NO), due to its activation of the inducible nitric oxide synthase pathway (Farrell et al. 1993). NO plays a critical role in the pathogenesis of joint inflammation and tissue damage, since the severity of RA can be reduced by the administration of NOS inhibitors in mouse model. Many cell types especially macrophages can be induced to produce NO by bacterial cell wall lipopolysaccharides (LPS). Therefore, inhibition of NO production in LPS stimulated RAW 264.7 cells is one of the possible ways to screen for anti-inflammatory drugs. *Alpinia calcarata* Roscoe (Family: Zingiberaceae) commonly known as lesser galangal in English, heen araththa in Sinhala, nattaratta in Tamil is widely used in Sri Lankan traditional medicine as a remedy for bronchitis, cough, respiratory ailments, diabetics, asthma and arthritis. Maharasnadhi Quathar (MRQ) is a polyherbal formulation used to treat RA which contains mainly 80 % of *A. calcarata* and 25 other plant materials. Studies on MRQ has proven that it can significantly and dose-dependently inhibit carrageenan-induced rat paw oedema and in RA patients, after 3 months of MRQ treatment, there was a marked improvement in the pain and inflammation as well as in the mobility of the affected joints (Thabrew et al. 2008). Previous studies on anti-inflammatory activity of hot water and ethanolic extracts of *A. calcarata* have shown significant inhibition of inflammation in carrageenan-induced rat paw oedema (Arawwawala et al. 2011). Based on the highly acclaimed properties of *A. calcarata*, the present study investigated the *in vitro* NO inhibition of LPS induced RAW 264.7 murine macrophages.

Materials and Methods

Reagents and chemicals: Mouse macrophage cell line RAW 264.7 was obtained from American Type Culture Collection (ATCC) and grown in RPMI medium supplemented with 10 % fetal bovine serum (FBS), 100 units mL⁻¹ penicillin and streptomycin. All the cell culture reagents and lipopolysaccharide (LPS),

* selvaluxmy@iti.lk

Sulphanilamide, N-(naphthyl) ethylenediamine dihydrochloride, sodium nitrite, N-monomethyl-L-arginine acetate salt (NMMA) and 3-(4,5-dimethyl-2-thiazolyl)-2,5-diphenyl tetrazolium bromide (MTT) were purchased from Sigma Aldrich, USA.

Plant material and extraction: The shade dried and powdered separate plant parts (rhizome, leaf, stem and root) of *A. calcarata* were extracted with petroleum ether in hot and cold conditions to analyze the best way of extraction depends on the activity of extract. Hot extraction was performed in Soxhlet apparatus whereas the cold extraction was performed by maceration under cold conditions. Extracts were further filtered and solvents were evaporated in rotary vacuum evaporator under reduced pressure. Dried and powdered rhizome and leaves (500 g) were hydro distilled in Clevenger apparatus for 8 h to distill the essential oils.

MTT assay for cell viability: Mouse macrophage cell line RAW 264.7 were cultured in 96 well flat bottom plate at concentration of 1×10^5 cells/well at 37 °C and 5 % CO₂. After 14 h of pre-conditioning, cells were treated with various concentrations of extracts and oil ($7.8\text{--}500 \mu\text{g mL}^{-1}$) and ($0.1\text{--}50 \mu\text{g mL}^{-1}$) respectively for 30 min. Thereafter, culture media was removed and MTT dye (5 mg mL^{-1}) was added to the cultures and further incubated for 3.5 h at 37 °C. The formazan crystals made due to dye reduction by viable cells were dissolved in acidified isopropanol (1N HCl). Index of cell viability was calculated by measuring the optical density of colour produced by MTT dye reduction at 570 nm (Scudiero et al. 1988).

Nitric oxide inhibition assay: The RAW cells were cultured in RPMI 1640 at cell concentration of 1×10^5 cells/well for 6 h in 96 well plates. Cells were then treated with LPS ($1 \mu\text{g mL}^{-1}$) with $7.8\text{--}250 \mu\text{g mL}^{-1}$ and $0.1\text{--}2.5 \mu\text{g mL}^{-1}$ of extracts and oil respectively for 30 min. Cells were washed with culture medium, resuspended in LPS containing medium for 24 h at 37 °C and 5 % CO₂. The culture supernatants were collected and they were used to assess nitrite concentration as a measure of NO production. Equal volume of Griess reagent was mixed with the cell supernatant (100 μL) and the absorbance was measured at 540 nm (Yadav et al. 2003). Cells were treated with 1 mM NMMA as a positive control. The concentration of nitrite was calculated from standard curve drawn with known concentration of sodium nitrite. The results were expressed as mean \pm SEM.

Results and Discussion

Effect of extracts and oils on RAW 264.7 cell viability: MTT is a pale yellow substrate reduced by living cells to yield a dark blue formazan product. This process requires active mitochondria to produce the product. MTT assay was performed to find out the possible cytotoxicity of *A. calcarata* extracts and

oils, when treated for 30 min. Cells showed comparable viability, >80 % in extracts treated with less than 500 $\mu\text{g mL}^{-1}$ and 5 $\mu\text{g mL}^{-1}$ for oil (Fig. 1a).

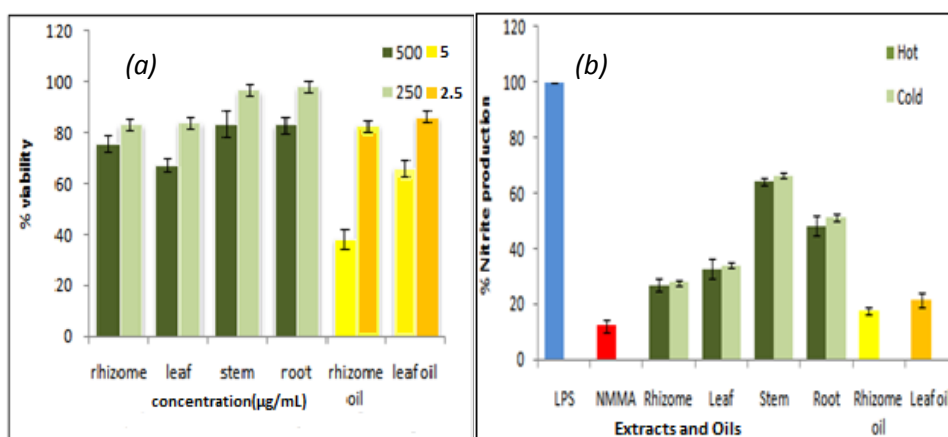


Figure 1. (a) Viability of RAW 264.7 macrophages in the presence of extracts at 500 and 250 $\mu\text{g mL}^{-1}$ and oils at 5 and 2.5 $\mu\text{g mL}^{-1}$. (b) Percentage of NO production by RAW 264.7 after treatment with extracts, oils and NMMA (+ control) with LPS stimulation. Data are expressed as mean \pm SEM of two independent experiments performed in duplicates.

Hence the concentration of (250-7.8 $\mu\text{g mL}^{-1}$) of extracts and (2.5-0.1 $\mu\text{g mL}^{-1}$) of oil were determined as non toxic and suitable for NO inhibition assay. In the present study, hot and cold solvent extracts of rhizome and leaf and the oils exhibited toxicity at higher concentrations where as stem and root had no toxicity and they have increased the macrophage proliferation in some concentrations.

Inhibition of LPS induced NO production by A. calcarata extracts and oils:

Murine macrophage RAW 264.7 cells were stimulated *in vitro* with LPS to produce NO and the effect of *A. calcarata* extracts on NO inhibition was assessed. All *A. calcarata* extracts, inhibited NO production in a dose-dependent manner. The unstimulated cells secreted NO at the basal level of $3.39 \pm 0.01 \mu\text{M}$, while the untreated LPS-stimulated cells showed an increase in NO production ($67.20 \pm 3.31 \mu\text{M}$). NMMA, a standard NOS inhibitor was used as the positive control and it inhibited NO ($12.91 \pm 0.01 \mu\text{M}$; 88.1 % inhibition) at 1mM. Among the *A. calcarata* extracts tested, hot extracts and oils of rhizome and leaf significantly inhibited NO production at all concentrations tested and the IC_{50} for petroleum ether extracts of rhizome and leaf are 124.2 ± 0.1 , $156.1 \pm 0.7 \mu\text{g mL}^{-1}$ (Fig. 1b). The stem and root extracts also inhibited NO production at 250 $\mu\text{g mL}^{-1}$ (35.0 ± 1.5 and 51.8 ± 3.5) which is

comparatively less than rhizome and leaf (73.2 ± 2.2 and 67.0 ± 3.5). Many studies have demonstrated that the NO inhibitory effect in RAW264.7 cells is due to the down regulation of iNOS. Inducible nitric oxide synthase (iNOS) is a soluble enzyme that catalyses the production of NO and the generation of high concentrations of NO through the activation of iNOS by immune stimulating cytokines and the activation of inducible nuclear factors such as NF- κ B may predispose an individual to arthritic conditions. Thus, the inhibition of iNOS exerts a beneficial anti-inflammatory effect on inflammatory disorders. This study demonstrated that the hot solvent extract and oils of *A. calcarata* rhizome and leaf significantly inhibited NO production in LPS-stimulated murine macrophage cells and scavenged NO radicals effectively.

Conclusions and Recommendations

In conclusion, this study has shown promising NO inhibition by *A. calcarata* with less or no cytotoxicity. Among them hot solvents extracts and essential oils of rhizome and leaf showed highest inhibition patterns than the other plant parts. We can conclude that the active compounds in rhizome and leaf are associated with this activity and further isolation and identification of compounds is necessary to validate the findings and the present study also validate the traditional use of *A. calcarata* for the treatment of RA in Sri Lanka.

Acknowledgement

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CURRENT STATUS OF POSTHARVEST HANDLING OF CUT FLOWERS AND CUT FOLIAGE AT PRODUCTION SITES IN THE UP COUNTRY OF SRI LANKA

G.D.R.K. Chandrasena¹, R.H.M.K. Ratnayake^{1*} and J.C. Edirisinghe²

¹*Department of Horticulture and Landscaping Gardening and*

²*Department of Agribusiness Management,*

Faculty of Agriculture and Plantation Management, Wayamba University of Sri Lanka

Introduction

The floriculture industry in Sri Lanka earns significant foreign exchange and generates employment, thereby, empowering rural communities. Although reliable statistics are rare on present status of the domestic market, a gradual increase in the local use of cut flowers and foliage is partly reflected by the recent expansion and distribution of retail flower shops in many urban and suburban areas of the Island. Retail outlets scattered throughout the production areas and in Western (WP), North Western (NWP) and Central (CP) Provinces are the popular centres where the cut flowers are sold for local consumption.

A range of tropical and temperate cut flower and foliage species are supplied to retail sites through various channels. The reliable supply of high quality cut flowers is important in maintaining customer satisfaction. Insufficient quality management or improper actions at different stages of handling cause a quality loss of products. Postharvest quality and vase life of cut ornamentals could be maintained throughout the distribution process by adopting correct handling practices, including proper postharvest treatments. Postharvest handling chains have been well characterised and standard treatments have been adopted elsewhere in the world. In Sri Lanka, it is essential to conduct a systematic study to clearly recognize the handling chains involved in cut flower supply channels for the domestic market. With a view to identify the shortcomings throughout the cut flower distribution chains, we have characterized the retail handling and major problems in retail sites revealed. This study was, therefore, conducted to identify the current handling practices in growers' and intermediaries' sites located in the up country of Sri Lanka. The outcome will help recognize the drawbacks in current handling systems.

Materials and Methods

Sampling and data collection: The study was conducted from January to June 2014. A sample of 34 cut flower growers and intermediaries in Uva and Central Provinces were selected for the survey. This sample also included the suppliers to retail sellers in WP and NWP. Information was gathered using a structured,

* kamani.ratnayake@uqconnect.edu.au

pre-tested questionnaire with face-to-face interviews. The information collected included: main products, methods of cultivation, harvesting details, quality of supplies, pre-treatments given to harvested produce, packaging and modes of transport, storage conditions, sanitation, postharvest problems, and financial and marketing aspects.

Data analysis: Data were descriptively analyzed using STATA (Stata Crop, United States) software package.

Results and Discussion

Species composition at production sites: Of the producers for the local market in the up country region, 98 % engaged in cut flower production instead of cut foliage. The species distribution among the respondents was: roses (77 %), gerbera (53 %), chrysanthemum (27 %), alstroemeria (24 %), white daisy (21 %), anthurium (6 %), orchid (3 %), zinnia (3 %), and others (< 3 %). The majority of growers (59 %) produced cut flowers and foliage in open field conditions while 41 % used protected houses such as poly-tunnels and net-houses.

Harvesting and postharvest handling at production and distribution sites: Table 1 presents the handling operations at production sites along with identified drawbacks. Lack of use of pre-treatments after harvesting, such as pulses with flower food and preservative compounds, could have undesirable consequences along the marketing channel. Temperature is the single most important factor in the keeping quality of stored floriculture products. Respiration, sprouting, water loss, change of chemical composition and the development of storage diseases are all influenced by temperature. Maintenance of low temperatures during transportation is, therefore, crucial to obtain maximum end-user life in cut ornamentals. Cut flowers and foliage produced in up country were sent to retail destinations mainly in Colombo (77 %) and Gampaha (74 %; WP), Kandy (74 %) and Nuwara Eliya (65 %; CP), and in other districts (35 %). Although the effects are unseen at the producer level, the implications given in Table 1 could decrease the potential post-harvest life of cut flowers.

The standard corrugated fiberboard boxes (CFB) were used by a limited proportion of respondents to pack relatively high valued species such as roses. About 10 – 25 flower stems were bundled together, covered with newspapers and arranged in the CFB by placing cypress leaves and newspapers in between. Dry transport in traditional bamboo cages causes serious transport damages to products arriving at the retail centres. Our previous research reported that in retail sites, cut flowers and foliage supplies were occasionally (by 56 %) or rarely (by 31 %) rejected mainly due to damages occurred during transport; 37 % for cut flowers and 40 % for cut foliage. Around 35 % of cut flower rejections and 39 % of cut foliage rejections were caused by wilting.

Table 1. The drawbacks in handling operations at production and distribution sites of cut ornamentals in the up country region

Operation	Current practice	Potential implications
Harvesting	Harvest between 6.00 a.m. - 12.00 noon.	Flowers harvested closer to noon have early senescence.
	Cutting tools not sterilized.	A potential source of bacteria for cut stems and handling solutions.
Pre-treatments	Harvested stems kept in buckets of water, at room temperature for 2 - 3 h.	No pre-treatments with carbohydrate sources, anti-bacterial and anti-ethylene compounds. No pre-cooling is done to remove field heat.
	Sent to transport agents in non-refrigerated vehicles	No cold chain.
Packaging	15 % respondents used standard corrugated fibre board boxes. 62 % used sub-standard cardboard boxes. 24 % used bamboo cages.	Bamboo cages cause compression damage to cut flowers.
Modes of transport to retail sites	59 % by train; 25 % refrigerated vehicles at 10 – 20 °C for high valued species (e.g. Roses); 8 % by other non-AC vehicles; 8 % by bus.	No cooling system when transported by train, bus or other non-AC vehicles.
Treatments during transit	82 % used dry transport (i.e. no continuous water supply for cut stems); 17 % used wet transport for selected species (e.g. Alstroemeria, roses, orchids).	Prolonged dry transport causes water deficit stress in cut flowers and foliage leading to wilting. Transport buckets not disinfected.
	Duration in transit: 6 - 12 h	

There are essential pre-treatments that should be applied to cut flowers before transport, such as anti-ethylene treatments. However, no such treatments have been applied prior to arrival at the retail sites. Moreover both ethylene sensitive and insensitive species are packed together in cartons during transport. Major problems reported while on retail display period of 3 - 4 days include flower drop, petal discolouration, petal drop, petal wilting, failure to open buds and bent neck in cut flowers; leaf wilting, yellowing and leaf drop in cut foliage. Some of these problems observed during retail display

could also be a result of poor ethylene management. Thus, the handling practices adopted at production and distribution sites could have a direct relationship with postharvest problems reported in retail sites.

Conclusions and Recommendations

This field survey on qualitative aspects revealed that, least precautions have been taken at production and distribution sites to preserve the quality of cut flower products. The drawbacks include: lack of cold chain management during storage and transport to slow down the inherent physiological processes, non-application of appropriate pre-treatments such as pulsing with food sources, preservatives and anti-ethylene treatments (for ethylene sensitive species), and sub-standard packaging. These factors could significantly contribute towards the decreased quality of products at retail and consumer sites.

Acknowledgement

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FACTORS AFFECTING PREFERENCE FOR GENETICALLY MODIFIED FOODS: A PILOT STUDY AMONG UNDERGRADUATES

Y.A.P.K. Dayasena¹, I.C. Hettiarachchi² and P. Sivashankar^{3*}

¹*Postgraduate Institute of Science, University of Peradeniya*

^{2,3}*Department of Agribusiness Management,
Sabaragamuwa University of Sri Lanka*

Introduction

At present, there is a growing demand for Genetically Modified foods (GMF) as well as a growing public concern for GMF due to health risks. GMF are available in many countries. GMF are agricultural and livestock products where genes are artificially manipulated to get favorable traits. With rising living standards, public is much concerned about health, environmental and ethical issues arising from the consumption of GMF. Though there are very few studies to prove the negative effects of GMF, the skepticism within the consumers is high. To alleviate these studies on GMF preferences are needed. Though, a study in 2012 has looked into the preference of GMF and GMF labeling in Sri Lanka, it reveals that most of the consumers are in fact not heard about GMF. This study attempts to assess the preference for GM foods by university students and what factors shape their decisions to purchase GMF. This study differs from previous studies as this includes more variables on environmental and health concerns, geographic location, income, and awareness. Further, it is reasonable to get the respondents who have a fair level of education, or who know about GM foods, to look in to other variables.

Materials and Methods

A sample of 120 university students was randomly interviewed during the months of July and August, 2014 at the faculty of agricultural sciences of the Sabaragamuwa University of Sri Lanka. University Students represent all the differences in terms of location, and socio-economic differences from around the country. GM foods are not popular in Sri Lanka. Thus, going for a sample which has some understanding about the topic than going for the general public is justifiable. Pre tested questionnaire captured information related to Socio-economic details, Knowledge on GM foods, awareness and perceptions on GM foods and their level of willingness to consume GM foods if available in the market using likert scale.

Probit regression and simple descriptive statistics were used to analyze the data. Probit regression is suitable as the dependent variable is a limited binary response variable. Many studies have used probit regression for numerous

* sivashankar.p@hotmail.com

applications in the field of applied economics and more specifically on hypothetical markets to estimate the market demand. Demand is a function of price (P), product attributes (A) and income (I). This is extended with the inclusion of psychological factors and perceptions of respondents (C). Socio-demographic factors and knowledge and awareness about the product are the key things which make the consumer to make a decision regarding a new product. The general theoretical model becomes: $Y = f(P, A, I, C)$.

Accordingly, decision to consume GM Foods is based on the attributes of the GM food (environmental, health risks, and toxicity) and the individual's psychological characteristics like (perceptions, awareness, and knowledge) about the use of GM foods, food security, as well as attitudes towards new technologies and products and its price. The following empirical equation is used in the probit regression (Table 1).

$$WTP = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \beta_3 X_3 + \beta_4 X_4 + \beta_5 X_5 + \beta_6 X_6 + \beta_7 X_7 + \beta_8 X_8 + \varepsilon$$

Results and Discussion

Out of the total 64.4 % of the respondents were female and rest were (35.6 %) males. About 52 % of them were final year students followed by 32.5 % of third year students. Most of them come from Semi urban areas (56.8 %) followed by 27 % of rural residents. Among them 43.2 % of their households were earning a monthly income of between Rs. 25,000 to Rs. 40,000. Around 21 % of them were getting a monthly house hold income above Rs. 40,000. Respondents can get information on GM foods from multiple sources. Most of the students have got to know about GM foods from the classes (59.5 %) and from the Internet (56 %). About 46.7 % are familiar with plant derived GM products and 38 % are familiar with plant and animal derived GM foods.

Nearly 56 % of respondents think that GM foods are beneficial. Majority of the respondents are skeptic to buy nutritionally enhanced GM foods. Finally, 30 % say that they will buy GM foods if they are introduced in Sri Lanka, 56 % say that they will not buy GM food and 22 % are not decided. Though 47 % of respondent agree to GM crops are pest resistant, 51 % of the students do not even like to use GM foods in animal feed to improve meat quality. The main reasons to reject GM foods are thinking they are unhealthy, artificial and are a threat to the environment.

According to the probit model only gender, year of study, indices for environmental risk and health risk are significantly affecting the decision to consume GM foods at the price similar to Non GM foods. The pseudo R^2 is around 47 %. Probability value for the whole model is below 0.05, the whole

model is significant and is a good fitted model. This type of regressions using survey data yields usually low R^2 values. When the income increases respondents are more likely to go for GM foods. Males are 57 % less likely to consume GM foods, and First years are 48 % less likely to consume GMF than final years. Second and third years are indifferent with final years.

Table 1. Probit Regression results for factors affecting WTP

Var	DV=WTP(1/0)	Coefficient	p- value	Marginal effect (%)
X_1	Gender: Male, Ref: <i>Female</i>	-0.960**	0.035	0.5734
X_2	Year of Study (Ref:4 th Year)			
	1 st Year	-1.3787*	0.059	0.4881
	2 nd Year	-0.0696	0.921	0.7372
	3 rd Year	-0.6041	0.161	0.6412
X_3	Residence (Ref: <i>Rural</i>)			
	Urban	0.5836	0.341	0.7332
	Semi Urban	0.4147	0.368	0.7027
X_4	Income (Ref:More than Rs. 60,000)			
	Less than Rs. 15,000	1.2686	0.150	0.6725
	Between Rs. 15000 and 25,000	1.6729**	0.036	0.7477
	Between Rs. 25,000 and 40,000	1.6813**	0.029	0.7491
	Between Rs. 40,000 and 60,000	0.3729	0.670	0.4906
X_5	GMO Familiarity (Ref: Never)			
	Frequently heard	-7.4949	0.987	0.5163
	Occasionally heard	-6.7072	0.988	0.6778
X_6	GMO Benefits (Ref:Not sure)			
	Yes	-0.3245	0.592	0.7216
	No	-1.0256	0.120	0.57958
X_7	Index for Environmental Risk	-	0.031	
		0.8414**		
X_8	Index for health risk	-	0.003	
		0.9250**		
	Constant	10.4078	0.982	

Note: Probit regression $LR \chi^2 (16) = 67.81$, $Prob > \chi^2 = 0.00$

Log likelihood = -37.94, Pseudo $R^2 = 0.47$, $N = 116$

***, **, * Significant at 1 %, 5 % and 10 % significance level respectively.

For dummy variables the last level is chosen as the reference level.

Average income household's students are 74 % more likely to consume GM foods if available at same price compared to the high income household students. Respondents, who think that GM foods may affect the environment

and health of consumers, are less likely to consume GM foods at the price similar to non GM counterparts.

Conclusions and Recommendations

This paper studied the university students' awareness and perceptions on GMF and factors affecting their preference for GMF. Still the sample showed awareness regarding GM foods is quite low. Especially due to the nature of the subject many respondents were skeptical about the GM foods. More than half of the respondents know about the benefits of the GM products, Still 56 % of the respondents are not willing to buy GM foods if they are introduced in to Sri Lankan market. Mostly the perceptions regarding GM foods are through internet and classes, yet with the absence of scientific proof for those perceptions, a considerable amount of respondents are neutral on the given statements and undecided on purchasing GM foods. Our findings are in line with the general acceptance of GM foods in Europe and Some East Asian countries. Female students and final year students are more likely to consume GMF. Average income household student are more likely to consume GMF at the same price of their conventional counterparts. Negative beliefs on environment and health, negatively affects the consumption of GMF. Since the GM products are not available in the local market and consumers are not aware of it, it is hard to measure the consumer awareness and willingness to consume GMF. It is expected that in future people might adapt to GM foods and GM products with more market penetrations and marketing. A comprehensive study on actual consumers is important to make policy decisions and for industries.

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SERUM INSULIN LEVELS – IS IT A GOOD INDICATOR IN CHRONIC TYPE-2 DIABETES MELLITUS SUBJECTS?

Y.S.M. Dissanayake¹ and L.V. Athiththan^{1*}

¹Department of Biochemistry, University of Sri Jayawardenepura, Sri Lanka

Introduction

Type 2 diabetes (T2D) is characterized by hyperglycaemia due to defects in secretion of insulin and its action. Insulin resistance (IR) and hyperinsulinaemia are considered as key features of T2D. Worldwide prevalence of T2D has been increasing significantly. About 347 million people worldwide have diabetes and 90 % of them have T2D. South East Asian region accounts for the 2nd highest number of T2D cases and in Sri Lanka prevalence of diabetes was 10.3 % in 2008. The IR is a condition where insulin levels are higher than expected and also relatively higher to the level of glucose in the blood. Therefore hyperinsulinaemia is often caused by IR where genetic and environmental factors influence hyper secretion of insulin hormone from β -cells which ultimately leads to hyperinsulinemia. IR is the major underlying causative factor of metabolic syndrome (MS) and is one of the risk factors for several non-communicable diseases (NCD) such as type 2 diabetes, dyslipidaemia and cardiovascular diseases (CVD).

Several studies have introduced IR as a powerful predictor of T2D. Fasting serum insulin (FSI) level of normal healthy subjects in Sri Lanka was, 93.42 ± 54.17 (21.53-376.42) pmol L⁻¹, while FSI among Sri Lankan diabetics was 150.71 ± 87.37 (41.67-390.31) pmol L⁻¹ (Senevirathne et al. 2009). IR in urban and rural adult population in Sri Lanka was 22.3 %. T2D is linked with basal hyperinsulinaemia, reduced sensitivity to insulin, and disturbances in insulin release. Even though FSI is currently being measured in diabetes subjects some studies state that chronic T2D subjects with FPG exceeding 7.8 mmol L⁻¹ secrete insulin levels at a low level which is similar to healthy non-diabetic individual or even lower. Therefore in chronic diabetics, basal insulin secretion and FSI can be within the normal range or even lower. Even though studies have indicated serum insulin level as a marker of IR. Eventually, C-peptide has gained attention as an alternative marker for IR, which is a protein that is co-secreted with insulin on an equimolar basis from pancreatic beta cells. Unlike insulin, it doesn't undergo hepatic first pass metabolism, has a longer half-life, and has been recognized as a more stable and accurate marker of endogenous insulin secretion and the pancreatic health.

* lohinia@gmail.com

Hence this study was carried out to identify insulin resistance, hyperinsulinaemia and its association with FPG in a selected Sri Lankan T2D population and also to assess the clinical relevance of FSI in chronic diabetes subjects.

Materials and Methods

Diabetic subjects (121) aged 27-80 years who were already diagnosed as having diabetes mellitus by a registered medical practitioner or government hospitals and attend the clinic at Family Practice Centre, Faculty of Medical Sciences, University of Sri Jayewardenepura were enrolled, after obtaining ethical approval from the Ethics Review Committee of USJP. Informed written consent was obtained prior to the study. An interviewer administered questionnaire was used to collect information about socio-demographic factors, family history of diabetes, lifestyle patterns and physical activity.

About 3 mL of 8 - 10 hours overnight fasting venous blood samples were obtained from each subject to analyze FPG and FSI. Glucose oxidase method was used to measure plasma glucose level and solid phase Enzyme-Linked Immunosorbent Assay (ELISA) method was used for serum insulin assay. IR was estimated by the HOMA-IR, using the equation $\text{HOMA-IR} = [\text{FSI } (\mu\text{U mL}^{-1}) \times \text{FPG } (\text{mmol L}^{-1})] / 22.5$ (Schianca et al. 2003). FPG level was classified based on World Health Organization criteria [WHO, 2006]. Hyperinsulinaemia was measured when FSI is $\geq 75^{\text{th}}$ percentile of its distribution (Freiberg et al. 2004). As a measure of IR, subjects having $\text{HOMA-IR} \geq 2.6$ was taken as insulin resistant (Schianca et al. 2003). Results were analyzed using SPSS version 16 and Microsoft Excel 2010. Correlations and significant differences were determined and $p < 0.05$ was considered as significant.

Results and Discussion

Among the 121 study subjects, 64.5 % ($n = 78$) were females. Mean age of the study population was 59 ± 11 years. Most of the subjects were residing in Colombo (66.9 %) and majority had a monthly income $< \text{Rs. } 20000$ (72.7 %). Majority of the study population (62.8 %) had diabetes ≥ 5 years. Among the study population, 58.7 % had a family history of diabetes. Those who had $\text{FPG} \geq 7 \text{ mmol L}^{-1}$ ($n=59$), 66.1 % had a known family history for diabetes. These findings were supported by several prospective studies where first-degree family history had a strong correlation with twofold increased risk of future T2D. This denotes that genetic predisposition is a risk factor in development of T2D and those with family history should have proper screening time to time and ones who are already diagnosed should have a proper follow up and monitoring.

Study subjects had an average FPG value of $8.0 \pm 3.45 \text{ mmol L}^{-1}$ which indicates a poor glycaemic control among the subjects. Large number of the study population (82.7 %) had FPG value above the normal range ($\geq 4.0\text{--}5.6 \text{ mmol L}^{-1}$) suggestive of poor glycaemic control among the subjects. In this study population, good percentage (56.2 %) of subjects monitors FPG levels monthly. Blood glucose monitoring at least once in every 3 months would enable to see the effect of treatment and aids in assessing the patients' compliance. It is recommended that subjects with high risk and those who have complications should monitor their FPG levels at least every 3 months.

Geometric mean of FSI among the subjects was $34.67 \pm 2.3 \text{ pmol L}^{-1}$ which was below the given normal FSI value for normoglycaemic subjects (Senevirathne et al. 2009). Among the hyperinsulinaemic subjects geometric mean of FSI was $89.12 \pm 2.3 \text{ pmol L}^{-1}$. Though mean FSI of these subjects were within the normal range ($18\text{--}172 \text{ pmol L}^{-1}$) (Sultan et al. 2010) mean FSI of hyperinsulinaemic subjects was lesser than the given mean FSI for diabetics. Subjects whose FPG was $\geq 7 \text{ mmol L}^{-1}$, had a mean serum insulin level of $39.8 \pm 1.8 \text{ pmol L}^{-1}$ which indicates that FSI is normal in these T2D subjects. Concerning the duration of having diabetes among the T2D subjects even with high FPG, both groups with the duration less than and greater than 5 years had (less) geometric mean of insulin levels but similar in both groups (FPG 8.3 and 7.8 mmol L^{-1} and insulin 33.1 and 36.3). This may be due to deterioration of beta cells and are unable to maintain insulin secretion hence FSI decline precipitously. Yet 38 % of the study subjects were insulin resistant according to HOMA-IR values and their mean FPG and FSI levels were $9.7 \pm 4.3 \text{ mmol L}^{-1}$ and $69.18 \pm 1.58 \text{ pmol L}^{-1}$, respectively. Out of the subjects who had FPG $\geq 7 \text{ mmol L}^{-1}$, 54.2 % had HOMA-IR ≥ 2.6 but only 20.3 % of them had hyperinsulinaemia. Further FPG and FSI had a significant positive correlation with IR ($p=0.000$). Significantly higher mean FPG levels ($p<0.05$) were observed with insulin resistant subjects (HOMA-IR ≥ 2.6) and lesser frequency of screening ($<$ monthly) for FPG. Despite monthly income $\geq \text{Rs. } 20\,000$, lesser frequency of screening was observed in these study subjects and also they had FSI & HOMA-IR ≥ 2.6 which was significantly higher ($p<0.05$).

These findings were supported by American Diabetes Association (2010), indicating that T2D is strongly linked with insulin resistance but diabetics are not necessarily to become hyperinsulinaemic. Thus chronic T2D subjects can present with normal FSI hence low FSI in chronic diabetics is not an indicator of good control. Yet, hyperinsulinaemia can be a strong predictor of development of T2D and metabolic syndrome.

Conclusions and Recommendations

Though it is assumed that poorly controlled chronic diabetic subjects have high FSI, this study demonstrates that, FSI is not a good indicator to assess glycaemic control in chronic T2D subjects. FPG and FSI have a significant correlation with IR. Furthermore, frequency of screening of FPG levels and first degree family history plays a major role in the FPG levels in T2D subjects. Therefore it is recommended to increase the FPG screening frequency at least once in 3 months in subjects with a known family history and not to assess the FSI values in chronic T2D.

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ISOLATION, CHARACTERIZATION AND IDENTIFICATION OF LACTIC ACID BACTERIA FROM FERMENTED INDIGENOUS RICE FLOUR

D.M.W.D. Divisekera^{1*}, W.A.A.I. Warunasinghe² and I.G.N. Hewajulige¹

¹*Food Technology Section, Industrial Technology Institute, Colombo 7*

²*Faculty of Agricultural Sciences, Sabaragamuwa University of Sri Lanka*

Introduction

Rice (*Oryza sativa* L.) is the largest cereal prominently cultivated in Sri Lanka. Sri Lanka has indigenous rice varieties with high nutritional value. The research evidences regarding Lactic Acid Bacterial fermentation of indigenous rice in Sri Lanka is very rare. Lactic acid bacteria (LAB) from food sources are being extensively studied due to its diversified role in human health. Their function extends from normalization of intestinal microbiota composition in the gut and immunomodulation to metabolic effects such as reduce serum cholesterol effects, reduce toxic effects and lactose hydrolysis. Non-digestible foods provide nutritional substrate for the growth of lactic acid bacteria. Lactic acid bacteria (LAB) in traditional fermented foods are beneficial for flavoring foods and inhibiting pathogenic as well as spoilage bacteria in the products. According to the previous research evidences, rice can be used to develop functional foods with enhancing gastrointestinal health. The objective of this study is to isolate, characterize and identify the Lactic Acid Bacteria associated with fermented indigenous rice flour.

Materials and Methods

Four indigenous rice varieties namely *Kaluheenati*, *Kuruluthuda*, *Suwadhel* and *Madathawalu* were purchased from known farmer fields, milled to passed through a 0.5 mm sieve, mixed with sterilized distilled water (1:2 ratio, w/v) and allowed to ferment at 30 °C for 24 h. Serial dilutions were prepared up to 10⁻³ with sterilized saline from fermented sample, 0.1 ml from each dilution were spread on sterile MRS agar plates (Hi-Media, India) and incubated at 37 °C for 24 h (Suthan et al. 2010). Each colony forming unit isolated was streaked on fresh MRS plates to examine the purity, followed by gram staining. All purified isolates were inoculated in MRS broth and incubated (37 °C, 24h). Biochemical tests, indole, methyl red, vogus-prosper, citrate utilization, gelatin liquefaction, H₂S production, starch hydrolysis, urease and catalase were performed. DNA of the isolates was extracted using an in-house optimized SDS proteinaseK DNA extraction method. For the PCR, primers 1492R (5'TACGGYTACCTTGTTACGACTT - 3') and 27F (5'-AGAGTTTGATCMTGGCTC AG-3') were selected and for the sequencing, primers 518F (5' CCAGCAGCCGCGTAATACG 3') and 800R (5' TACCAGGGTATCTAATCC

* mb.wasu@gmail.com

3') were selected. The 16S rDNA sequencing was carried out at MacroGen-South Korea and sequence alignment was carried out by Basic Local Alignment Search Tool.

Results and Discussion

Seven organisms were isolated from fermented rice flour. Gram staining and colony characterization of the isolates from fermented rice flour is represented in Table 1, and Table 2 represents the biochemical and molecular level identification of the isolates.

Table 1. Gram staining and colony morphology of the isolates from fermented indigenous rice flour

Rice variety	Isolate number	Gram stain	Size	Surface	Texture	Color	Elevation	Margin
<i>Kaluheenati</i>	1	positive cocci	Punctiform	Glistening	Mucoid	Creamy white	Raised	Entire
<i>Kaluheenati</i>	2	positive cocci	Punctiform	Glistening	Mucoid	Creamy white	Flat	Entire
<i>Kaluheenati</i>	3	positive cocci	Punctiform	Glistening	Mucoid	Creamy white	Raised	Entire
<i>Suwandel</i>	4	positive cocci	Medium	Glistening	Mucoid	White	Raised	Entire
<i>Suwandel</i>	5	positive cocci	Medium	Glistening	Mucoid	White	Pulvinate	Entire
<i>Madathawalu</i>	6	positive cocci	Medium	Glistening	Mucoid	White	Pulvinate	Entire
<i>Kuruluthuda</i>	7	positive cocci	Punctiform	Glistening	Mucoid	White	Raised	Entire

Among the seven isolates, three organisms were able to isolate from fermented flour of *Kaluheenati*. Two organisms were able to isolate from fermented flour of *Suwadel*, where *as one organism was able* to isolate from fermented flour of *Madathawalu* and *Kuruluthuda*. All the seven isolates were observed to be Gram positive, morphologically observed to be cocci.

Table 2. Biochemical and molecular level identification of the isolates from fermented indigenous rice flour

Isolate no.	Indole	Methyl red	VogesProskauer	Citrate utilization	Gelatin liquefaction	Urease test	H ₂ S production	Catalase	Starch hydrolysis	Molecular level identification
1	-	+	-	-	+	-	-	-	-	<i>Enterococcus durans</i> 98D
2	-	+	-	-	-	-	-	-	-	<i>Enterococcus faecium</i> Aus 0004
3	-	+	-	-	-	-	-	-	-	<i>Enterococcus faecium</i> strain LMG 11423
4	-	+	-	-	+	-	-	-	-	<i>Staphylococcus warneri</i> SG1
5	-	+	-	-	-	-	-	-	-	<i>Staphylococcus pasteurii</i> strain ATCC51129
6	-	+	-	-	+	+	-	-	-	<i>Staphylococcus epidermidis</i> RP62A
7	-	+	-	-	+	-	-	+	-	<i>Enterococcus spp</i>

All seven isolates were negative for citrate utilization test, indole test, starch hydrolysis test, H₂S production test and vogesproskauer test. However, all the isolates were positive for methyl red test. Isolates namely no.1, 4, 6, and 7 were observed to be positive for gelatin liquefaction. Whereas only isolate no. 6 was positive for urease test. Except isolate no.7, all the other isolates were observed to be catalase negative. By molecular level identification, the isolates were identified as *Enterococcus durans*98D (Isolate 1), *Enterococcus faecium* Aus 0004 (isolate 2), *Enterococcus faecium* strain LMG 11423 (isolate 3), *Staphylococcus warneri*SG1 (isolate 4), *Staphylococcus pasteurii* strain ATCC51129 (isolate 5), *Staphylococcus epidermidis*RP62A (isolate 6) and *Enterococcus spp* (isolate 7). This is the first time reporting of *Enterococcus spp.* and *Staphylococcus spp.* from fermented indigenous rice in Sri Lanka.

Staphylococcus spp was reported to be isolated from fermented shrimp in Malaysia; there has been very little report on this genus in fermented food for the past decade (Probst et al. 1998). *Enterococcus spp* is the most controversial group of lactic acid bacteria. Studies on the microbiota of many traditional cheeses in the mediterranean countries have indicated that *Enterococcus spp*

play an important role in the ripening of these cheeses, probably through proteolysis, lipolysis, and citrate breakdown, hence contributing to their typical taste and flavor. *Enterococcus spp* are also present in other fermented foods, such as sausages and olives. However, their role in these products has not been fully elucidated. Production of bacteriocins by *Enterococcusspp* is also well documented. *Enterococcus spp* are nowadays used as probiotics even though these have been associated with a number of human infections. *E. faecium* have been applied in human as probiotic supplements. Since 2004, ten preparations (9 different strains of *E. faecium*) are authorized as additives in food in the European Union that are proposed to be clinically effective in the prevention and treatment of antibiotic associated diarrhea in children. *Enterococcus hirae* (*Streptococcus faecalis*) ATCC 9790 is a Gram positive lactic acid bacterium and reported to be isolated from fermented fish. *E. durans* use as co-cultures in the production of Feta cheese and a white-brined cheese. Among the seven isolates in this study, *Enterococcus faecium* is the most frequently reported species among the *Enterococcus spp.* that literature reveals as a starter culture for fermentation (Moreno et al. 2006). There is no data available in the literature on isolation of *Enterococcus spp.* from fermented indigenous rice flour. Therefore, this is the first time reporting of *Enterococcus spp.* from fermented indigenous rice in Sri Lanka.

Conclusions and Recommendations

Seven potential probiotic isolates namely; *Enterococcus durans* 98D, *Enterococcus faecium* Aus 0004, *Enterococcus faecium* strain LMG 11423, *Staphylococcus warneri* SG1, *Staphylococcus pasteurii* strain ATCC51129, *Staphylococcus epidermidis* RP62 and *Enterococcus spp* were isolated from fermented indigenous rice flour. The contribution of potential probiotics to the organoleptic and functional properties of fermented food products are important characteristics for their application in food technology. Therefore, a proper toxicological study should be conducted to assess the safety of the isolates for use as a starter; also functional properties of the isolates should be investigated.

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INVESTIGATION OF ANTIBACTERIAL PROPERTIES OF ENDOPHYTIC FUNGI ISOLATED FROM *Nymphaea nouchali* ENDEMIC TO SRI LANKA AND THE ISOLATION OF CHAETOGLOBOSIN C FROM THE ENDOPHYTIC *Chaetomium* sp.

D.M.R.K. Dissanayake¹, P. B. Ratnaweera^{1,2}, E. D. De Silva^{1*}, D. E. Williams², C.D. Wijayarathne¹, R.L.C. Wijesundera³ and R. J. Andersen²

¹ *Department of Chemistry, University of Colombo, Sri Lanka*

² *Department of Earth, Ocean and Atmospheric Sciences, University of British Columbia, Canada*

³ *Department of Plant Sciences, University of Colombo, Sri Lanka*

Introduction

Endophytic fungi which reside symbiotically inside plants are considered to be an outstanding source of novel bioactive secondary metabolites. However, only a few studies have been carried out so far to evaluate the potential of endophytic fungi inhabiting Sri Lankan biota (Ratnaweera et al. 2014). The rich biodiversity and the high degree of endemism among Sri Lankan plants suggest the possibility that the endophytes contained in them may have unique biosynthetic capabilities leading to the production of unique metabolites with varied biological activities. Therefore, the investigation of the endophytic fungi of Sri Lankan endemic plants for the presence of antimicrobial substances becomes especially attractive. The present study was carried out to isolate and characterize the antimicrobial substances produced by laboratory cultures of endophytic fungi isolated from the aquatic endemic plant *Nymphaea nouchali* (Nil Manel in Sinhalese).

Materials and Methods

Specimens of *N. nouchali* were collected from Udugampola, Gampaha District, Sri Lanka. Healthy leaves, stems, sepals and petals of the plant were surface sterilized, within five hours of collection, by sequentially washing with sterile distilled water (2 min), 70 % Ethanol (30 sec) and 5 % NaOCl solution (1.5 min). Next sterilized plant parts were dried, cut into 0.5 cm² segments and placed on dilute Malt Yeast Agar (dMYA) plates under aseptic conditions and left at room temperature up to seven days, until endophytic fungi emerged. The emergent fungi were sub-cultured on Potato Dextrose Agar (PDA) to obtain 24 pure cultures (10 from leaves, 05 from stem, 05 from sepals and 04 from petals) of morphologically distinct endophytic fungi. Next each distinct fungal species was cultured on 06 PDA plates (120 mm × 20 mm) for 03 - 05 weeks and the culture media together with the fungal mycelia were extracted in to

* edilip.desilva@gmail.com

ethyl acetate and concentrated under reduced pressure to obtain 24 crude organic extracts.

Antibacterial activity of the resulting 24 organic extracts were evaluated using disc diffusion method against *Staphylococcus aureus* (ATCC 25923), *Pseudomonas aeruginosa* (ATCC 9027), *Escherichia coli* (ATCC 35218), and *Bacillus cereus* (ATCC 11778) at 300 and 50 $\mu\text{g disc}^{-1}$ (National Committee for Clinical Laboratory Standards 2003). The obtained results were statistically analyzed by one way ANOVA using Minitab.

The fungus RDNM-04, which was active at 50 $\mu\text{g disc}^{-1}$ against *S. aureus* and *B. cereus*, was selected for compound isolation and molecularly identified by analysis of the nucleotide sequence of the ITS region using BLAST analysis. In order to isolate active metabolite(s) the fungus was grown large scale on PDA (250 Petri dishes, 120 mm \times 20 mm) for 18 days until sporulation. The culture medium together with the fungal mycelium was next extracted with EtOAc (3 \times 1.5 L) and the extract was filtered and concentrated at reduced pressure to obtain the crude organic extract.

The organic extract (2.2 g) was subjected to bioassay guided fractionation, first by normal phase silica gel column chromatography (2 cm \times 25 cm column; gradient elution starting with hexane, then mixtures of hexane/EtOAc to EtOAc) to give 58 mg of active material which was further purified by re-chromatographing on silica gel (1.5 cm \times 35 cm column; gradient elution from hexane/EtOAc, 30:70 to EtOAc/MeOH, 30:70) to give 20 mg of the active fraction. The final purification was achieved by size-exclusion chromatography on Sephadex LH 20 (1.5 cm \times 60 cm column with MeOH:CHCl₃, 1:1) to obtain 7.5 mg of a white amorphous solid which gave a single spot on thin layer chromatography. The ¹H, ¹³C and 2D NMR (COSY, HSQC, HMBC) spectral data of the pure active compound were obtained using a Bruker ADVANCE 600-MHz spectrophotometer (DMSO-d₆ used as a solvent) while mass spectroscopic (MS) data were obtained using a Bruker Esquire-LC electrospray spectrophotometer.

Results and Discussion

The results of antibacterial bioassays showed that of the 24 fungal extracts, 15 were inactive against all 04 bacterial species tested. Of the 09 active extracts, all inhibited the growth of *S. aureus* and *B. cereus* at 300 $\mu\text{g disc}^{-1}$ while only 02 extracts (RDNM-01 and RDNM-18) were active against *P. aeruginosa* and *E. coli* at this concentration. At 50 $\mu\text{g disc}^{-1}$ RDNM-01, RDNM-04, RDNM-06, RDNM-18 and RDNM-22 were active against *S. aureus* and *B. cereus* while only RDNM-1 and RDNM-18 were active against *P. aeruginosa* and *E. coli* (Table 1). A GenBank search for similar sequences showed that the 18S rRNA gene

sequence of RDNM-04 contain 99 % sequence identity to that of *Chaetomium* sp.

Table 1. Antibacterial activity of the organic extracts of endophytic fungi isolated from *Nymphaea nouchali* (Mean±SE)

Identification No. of Fungi	Origin	Antibacterial activity (µg disc ⁻¹) (Inhibition zone diameter - mm)							
		<i>S. aureus</i>		<i>B. cereus</i>		<i>P. aeruginosa</i>		<i>E. coli</i>	
		300	50	300	50	300	50	300	50
RDNM-01	Sepals	17±0.2	12±1.2	22±0.3	12±0.8	8±0.5	7±0.3	15±1.4	8±0.9
RDNM-04	Leaves	16±1.3	13±0.5	19±0.6	10±0.2	-	-	-	-
RDNM-06	Leaves	14±0.1	10±0.9	16±0.5	14±1.6	-	-	-	-
RDNM-13	Stem	11±0.3	-	11±0.8	-	-	-	-	-
RDNM-18	Leaves	25±0.2	15±1.4	27±1.6	17±1.1	8±0.6	8±0.1	20±2.3	10±0.3
RDNM-20	Leaves	10±0.4	-	12±0.1	-	-	-	-	-
RDNM-21	Stem	08±0.2	-	09±0.5	-	-	-	-	-
RDNM-22	Sepals	12±0.4	07±0.6	14±0.9	08±1.0	-	-	-	-
RDNM-23	Leaves	09±0.2	-	08±0.7	-	-	-	-	-
+ve		23±1.0	21±0.9	21±0.8	22±2.6	20±2	21±1.1	22±1.5	22±1.9

Large scale culturing of the fungus RDNM-04 yielded 2.2 g of crude extract and bioassay guided fractions yielded 7.5 mg of a white pure active compound. High resolution mass spectral data (M^+ , 528.6429 g mol⁻¹) indicated a molecular formula of C₃₂H₃₆N₂O₅. Interpretation of ¹H, ¹³C and ¹⁵N NMR together with 2D (COSY, HSQC, HMBC, HSQC) NMR data identified the structure of active compound as the known cytochalasan, Chaetoglobosin C (Fig. 1). A comparison of ¹³C NMR values obtained in the present study for chaetoglobosin C with the reported data (Sekita et al. 1976) is given in Table 2.

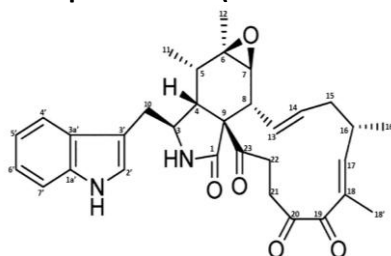


Figure 1. Chemical Structure of the Chaetoglobosin C. Chaetoglobosin C, a member of the cytochalasan family of compounds, was first isolated from the fungus *Chaetomium globosum* from Japan in 1976 together with the Chaetoglobosin A, D, E and F.

Table 2. Comparison of ^{13}C NMR data of chaetoglobosin C from the present study with published data in DMSO-d₆

^{13}C (ppm)								
C#	Present study	Published data	C#	Present study	Published data	C#	Present study	Published data
1	173.96	173.79	13	127.08	127.04	22	37.17	37.04
3	52.38	52.25	14	133.14	133.11	23	196.28	196.04
4	48.50	48.35	15	39.15	39.47	2'	121.03	120.86
5	36.19	36.10	16	32.63	32.50	3'	108.07	108.02
6	56.75	56.87	16'	19.43	19.28	3 α '	127.80	127.63
7	60.31	60.28	17	155.92	155.56	4'	118.42	118.25
8	48.19	48.24	18	131.05	130.95	5'	125.58	125.02
9	62.40	62.25	18'	10.08	9.93	6'	118.78	118.25
10	31.73	31.88	19	208.29	208.00	7'	111.35	111.22
11	12.42	12.30	20	205.48	205.15	1 α '	135.88	135.81
12	19.13	18.99	21	31.98	31.88			

Conclusions and Recommendations

N. nouchali harbors many endophytic fungi which are capable of producing antibacterial substances. Although the antibacterial compound isolated from the fungal extract RDNM-4 proved to be known, the possibility of isolating new biologically active compounds from the remaining active extracts still remains. Thus it can be concluded that the endophytic fungi of Sri Lankan endemic flora are a valuable potential source for the isolation of bioactive metabolites and more investigations should be performed to realize its true potential.

Acknowledgement

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DETERMINATION OF ANTIBACTERIAL ACTIVITY OF SRI LANKAN TEA BREWS BY BROTH MICRODILUTION ASSAY

T. M. Erandika¹, P.D. Abeysinghe^{1*}, R.N. Pathirana² and S. Abeysinghe³

¹*Department of Botany, University of Ruhuna, Matara, Sri Lanka*

²*Kotalawala Defence Academy, Ratmalana, Sri Lanka*

³*Tea Research Institute, Thalawakele, Sri Lanka*

Introduction

Even though pharmacological industries have produced a number of new antibiotics in the last three decades, the development and the rapid spread of strains resistant to the synthetic antibiotics by micro-organisms have become a global problem. In general, bacteria have the genetic ability to transmit and acquire resistance to drugs which are utilized as therapeutic agents. Some 'hospital-acquired' infections may be resistant to a number of clinically efficacious antibiotics and become less effective even to the high dosage of antibiotics. Therefore, measures such as controlling the use of antibiotics or search for new drugs or natural products have to be carried out to overcome this problem.

The antibacterial effect of many herbs and other materials of plant origin are widely known. Studies have been conducted on the possibility of using plant extracts in treating diseases caused by microbial strains resistant to antibiotics. These investigations also include tea (*Camellia sinensis*) extracts. Some investigations have found inhibitive properties of black, green and pu-erh teas in relation to *Staphylococcus aureus*, and *Staphylococcus epidermidis*, and of black and green teas in relation to *Vibrio cholerae*, *Vibrio mimicus*, *Plesiomonas shigelloides*.

Sri Lanka is the third largest tea producer and the exporter in the world. Although only few researches have been carried out on the antibacterial properties of teas produced within the island. Therefore, objective of the present study was to investigate antibacterial properties of different brands of tea manufactured in Sri Lanka using microdilution assay which is simple, sensitive, rapid, robust and a reliable method.

Materials and Methods

Methicillin-resistant *Staphylococcus aureus* (MRSA) and *Mycobacterium smegmatis* (MS) were used for investigation. Fourteen available black tea samples were obtained from Tea Research Institute (TRI), Sri Lanka. Ten green

* pushpa@bot.ruh.ac.lk

tea samples were obtained from Melfort Green Teas (Pvt) Ltd, Peliyagoda, Sri Lanka (Table 1).

Each tea sample (2 g) was brewed with 100 ml of boiled water in a covered container for 6 minutes, 1 ml of filtrate was transferred into sterilized centrifuge tubes. The dry matter content of tea was measured using oven method (at 110 ± 5 °C temperature for constant mass). Triplicates of brews were prepared from each tea sample and triplicates of assays were carried out using each brew to have nine replicates.

Determination of Minimum Inhibitory Concentration (MIC): The minimum inhibitory concentration (MIC) of each tea brew was determined using the previously described broth microdilution colorimetric assay with slight modifications. Main modification was the adding resazurin indicator after the incubation of the plates (previous method added the indicator before incubation).

The microtitre plates including eight rows and twelve columns with 96 wells were prepared for the assay in aseptic conditions. A volume of 100 µl of tea brews were pipetted into each well in the first row. To all wells in other rows 50 µl of sterilized distilled water was added. Serial dilutions were performed from first to last row in each column (in serially descending concentrations) using a micro pipette to make 50 µL of the tea brew in each well. The excess volume in the last row was discarded.

Using a micropipette 40 µl of $2.5 \times$ strength LB broth was added to each well to ensure that the final concentration of nutrient broth was single strength. Finally, 10 µl of bacterial suspension (5×10^6 CFU mL⁻¹) was added to each well to achieve a concentration of 5×10^5 CFU mL⁻¹ in the well. Amoxillin (1mg mL⁻¹) and distilled water were used as positive and negative controls respectively in each plate. As a sterility control one tea sample was used without bacteria. Sealed plates were incubated for 18 hours at 37 °C and then 30 µl of 0.25 mg mL⁻¹ resazurin solution as a bacterial growth indicator was added to all microplate wells. The plates were incubated further for 30 minutes. The color change was then assessed visually. Any color changes from purple to pink or colorless were recorded as positive. The lowest concentration at which color change occurred was taken as the MIC value which was the MIC for the test material and bacterial strain. All analyses were conducted in nine replicates, and the results were expressed as mean \pm standard deviation (SD). The differences between means were first analyzed by ANOVA test and then Duncan's multiple range tests ($p < 0.05$) using the SAS version 9.1.3 software.

Results and Discussion

Dry matter content of tea brews was within the wide range from 5.34 (Pekoe (a)) to 8.25 mg mL⁻¹ (BOPF (a)). Therefore, boiled water extractable compounds of tea are highly depends on the tea grades.

Table 1. Minimum Inhibitory Concentration (MIC) of brews from Sri Lankan black and green tea against *Mycobacterium smegmatis* (MS) and Methicillin-resistant *Staphylococcus aureus* (MRSA)

Tea type	Sample	Dry weight of brew (mg mL ⁻¹)	MIC (mg mL ⁻¹)	
			MS	MRSA
Black tea	BOP	6.84	1.43 ± 0.40 ^{ef}	0.38 ± 0.09 ^f
	BOPF (a)	8.25	2.78 ± 0.89 ^a	0.94 ± 0.33 ^{bcd}
	BOPF (b)	5.77	1.95 ± 0.32 ^{bcdef}	1.72 ± 0.49 ^a
	BOPF (c)	7.15	1.92 ± 0.68 ^{bcdef}	0.56 ± 0.18 ^{ef}
	BOPI (a)	6.15	2.05 ± 0.72 ^{abcde}	1.03 ± 0.36 ^{cb}
	BOPI (b)	5.58	2.48 ± 0.58 ^{abc}	0.85 ± 0.29 ^{bcde}
	Dust (I) (a)	5.58	1.24 ± 0.64 ^f	0.70 ± 0.28 ^{cdef}
	Dust (I) (b)	6.2	1.64 ± 0.85 ^{def}	0.86 ± 0.24 ^{bcde}
	Dust (a)	6.35	1.68 ± 0.48 ^{def}	0.63 ± 0.18 ^{def}
	Dust (b)	6.06	2.65 ± 0.75 ^{ab}	1.15 ± 0.39 ^b
	OPI (a)	5.38	2.24 ± 0.63 ^{abcd}	0.82 ± 0.28 ^{bcde}
	OPI (b)	5.64	2.04 ± 0.70 ^{abcde}	0.78 ± 0.36 ^{cde}
	Pekoe (a)	5.34	1.93 ± 0.66 ^{bcdef}	0.96 ± 0.33 ^{bcd}
	Pekoe (b)	6.24	1.73 ± 0.80 ^{cdef}	0.65 ± 0.18 ^{def}
Green tea	Chunmee (I)	6.19	1.20 ± 0.38 ^q	0.28 ± 0.10 ^{stu}
	Chunmee (II)	7.59	2.00 ± 0.70 ^p	0.63 ± 0.22 ^{pq}
	Fanning's	7.15	0.55 ± 0.19 ^r	0.79 ± 0.19 ^p
	Gun Powder	7.15	0.70 ± 0.22 ^r	0.40 ± 0.09 ^{rs}
	Hyson	6.36	0.66 ± 0.19 ^r	0.42 ± 0.15 ^{rs}
	OPA	7.14	0.69 ± 0.43 ^r	0.37 ± 0.11 ^{rs}
	Sp. Hyson	6.62	1.20 ± 0.41 ^q	0.32 ± 0.20 ^{stu}
	Ceylon Sen.OPA	6.58	0.55 ± 0.19 ^r	0.59 ± 0.20 ^q
	Ceylon Tencha	7.72	0.38 ± 0.12 ^r	0.25 ± 0.09 ^s
	Cut twist curl	7.14	0.35 ± 0.11 ^r	0.52 ± 0.21 ^{qr}

Note: Values are mean (n=9) ± standard deviation (SD). Values with the same superscript letter within each column and within each black and green tea categories are not significantly different (p≤0.05). Bold letters are for the best tea samples with lowest MIC values in each of black and green tea for each bacterium. Black and green tea samples were analyzed separately.

In our study, MRSA was more sensitive than that of MS for all tea brews of all tea grades. From tested black tea samples Dust (I) a and BOP grade had the

best antibacterial activity with the 1.24 mg mL⁻¹ and 0.38 mg mL⁻¹ MIC values for MS and MRSA respectively. BOPF (a) tea sample had the lowest antibacterial activity for MS from the tested black teas with 2.78 mg mL⁻¹ MIC value. Although dry matter content of BOPF (a) was highest, it may consist of less amount of active ingredients of BOPF (a) sample. Therefore, its antibacterial activity of BOPF (a) may be low. BOPF (b) was the sample with the lowest antibacterial activity for MRSA with the 1.72 mg mL⁻¹ MIC value. The properties of tea which inhibit bacterial growth are mainly related to their polyphenolic components. *In vivo* and *in vitro* investigations have observed the beneficial effect of green tea catechins in the treatment of periodontal disease which is caused by mainly *Porphyromonas gingivalis* and *Tannerella forsythia* and *Actinobacillus actinomycetemcomitans*.

The best antibacterial activity was shown by the CTC green tea and Ceylon Tencya green tea with 0.35 and 0.25 mg mL⁻¹ MIC values for MS and MRSA species respectively. Almost all green tea samples had higher antibacterial activity for both bacteria strains than those of black tea. The lowest inhibition activity measured by the MIC for green teas for MS and MRSA were shown by the Chunmee(II) green tea (2.00 mg mL⁻¹) and Fanning's green tea (0.79 mg mL⁻¹), respectively.

Conclusions and Recommendations

Both Sri Lankan green and black tea brews have good antibacterial activity against MS and MRSA bacteria species. However, a detailed biological and phytochemical studies are needed to find out the chemical constituents responsible for their activities. The antibacterial compounds in tea should be separated and purified. Antibacterial activity of purified compounds should be further tested for MRSA and MS bacteria.

Acknowledgement

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PREPARATION OF MICROWAVE IRRADIATED CELLULOSE BASED BIODEGRADABLE SUPPER ABSORBENT POLYMER (SAP) FOR AGRICULTURAL APPLCAITIONS

T.N. Fernando¹, A.G.B. Aruggoda^{1*}, C. K. Disanayaka² and S. Kulathunge²

¹*Department of Agricultural and Plantation Engineering,
Faculty of Engineering Technology, the Open University of Sri Lanka, Nawala,
Nugegoda*

²*Atomic Energy Authority, No 60/460, Baseline road, Orugodawatta, Wellampitiya,
Colombo, Sri Lanka*

Introduction

Superabsorbent polymers (SAPs) are highly swollen, hydrophilic polymer networks, capable of absorbing 400-1500 g of water per dry gram of SAP is having the possibility of applying on wide range of industries such as agricultural, medicinal, environmental, horticultural etc. Cellulose based biomaterials could be used as raw materials for developing SAP with high biodegradability, high strength after absorbing water, less water soluble components, strong water retaining ability and mould proofing ability (Czaja et al. 2007). Therefore, SAP from cellulose is cheap and superior in performance. Most agricultural residues such as corn stove, wheat straw, rice straw and bagasse which referred as lignocellulosic materials, are rich in cellulose fibers. Microwave technology is a green method for chemical synthesis of cellulose because of its high efficiency and homogenous nature of heating. The present study focused on synthesizing SAP by graft copolymerization of cellulose from bagasses and acrylic acid, using N, N-MethyleneBisAcrylamide (MBA) as a cross linker and Potassium persulfate ($K_2S_2O_8$ /KPS) as an initiator in an aqueous solution under microwave irradiation.

Materials and Methods

Cellulose was prepared by using alkaline pretreatment method as described by Feng et al. (2010) and it's percentage was determined using chlorinated method as described in Google Book (2014). SAP was prepared using a set up as described in Figure 01. Certain amount of MBA, Acrylic acid and 5 mol L⁻¹ NaOH solutions were mixed and kept in an ice bath until use. 1.0 g of bagasse cellulose and 50 mL of de- ionized water were added to the vacuum flask. Temperature was adjusted to 150 °C and content inside was stirred continuously. Simultaneously oxygen free nitrogen gas was bubbled in to the solution for 30 minutes. Thereafter, certain amount of KSP was added and stirred continuously for 15 minutes by reducing the temperature to 90 °C. Pre

* agbar@ou.ac.lk

cooled solution mixture was added to the above vacuum flask and stirred. Final mixture was subjected to microwave irradiation under the power of 340 W for 7 minutes. The product was cooled to room temperature, cut in to small pieces and immersed in absolute ethanol for 30 minutes followed by drying at 60 °C until gaining a constant weight. Un-grafted cellulose sample as well as grafted SAP was FT-IR analyzed and grafted SAP was subjected to Scanning Electronic Microscope (SEM) to analyze the morphological characteristics. Swelling ratio was determined by placing and immersing 1 g of SAP in a 200 mesh sieve pouch in 250 ml of distilled water over night. The swelling ratio was calculated as follows,

$$\text{Swelling ratio} - Q \text{ (g/g)} = (m_1 - m_0) / m_0$$

(Where m_0 and m_1 are the weight of the dry and swollen SAP respectively) Biodegradability was measured by embedding nylon cloth wrapped 100 g of water absorbed SAP to a depth of 10 cm, in pots filled with sandy loam soil. This was repeated three times and each sample was weighted at 14 days intervals until attain total degradation. The percentage of biodegradability was calculated as follow,

$$\text{Percentage of biodegradability} = (m_s - m_d) / m_s \times 100\%$$

(Where m_s the weight of the un- degraded SAP and m_d is the weight of the biodegraded SAP).

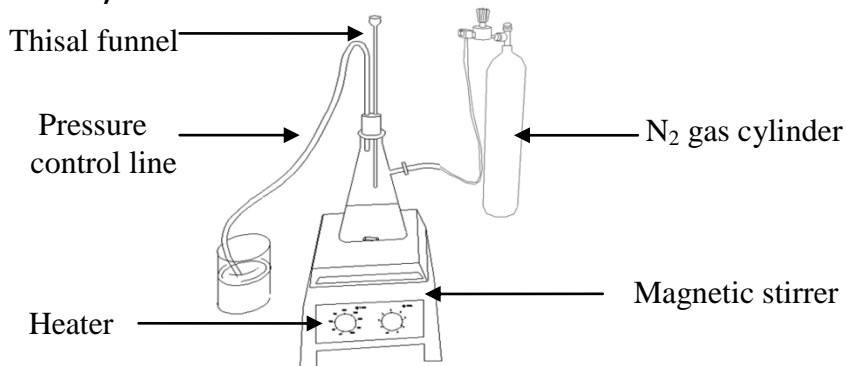


Figure 1. Schematic diagram of used apparatus for processing of SAP

Results and Discussion

83 % of cellulose was resulted from bagasse from alkaline pretreatment. Swelling ratio was 1:401. The IR spectra of the bagasse cellulose and grafted polymer prepared are shown in Figs. 2 and 3. The main characteristic peaks of cellulose are at 1162.1 cm⁻¹, 1070.7 cm⁻¹ (pyran structure) 3483.2 cm⁻¹ and 3427.1 cm⁻¹ (O-H Structure) and 2927 cm⁻¹ (C-H stretch). The small peaks at 1637.6 cm⁻¹ 1458.4 cm⁻¹ result from -C=O stretching and amorphous cellulose, respectively. The absorption band at 1378.7 cm⁻¹ and 897.8 cm⁻¹ are described to C-H bending vibration (Fig. 2). Some characteristic absorption peaks from acrylic acid and polyacrylate appear In addition to the cellulose peaks in grafted

cellulose (Fig. 3). The peak at 1718.1 cm^{-1} corresponded to the carboxyl absorption from grafted Poly Acrylic Acid (PAA). Furthermore, the bands at 1577.2 cm^{-1} , 1399.6 cm^{-1} and 1419 cm^{-1} corresponded to the sodium carboxyl group indicating the PAA (Na) grafting to cellulose. SEM indicates clear inter connected porous surface in SLSAP (Fig. 1). This surface morphology may accelerate the penetration of water into the polymer network. Biodegradability of SAP under average soil temperature of $29\text{ }^{\circ}\text{C}$ and at 10 cm depth in soil and with the average environment temperature $32\text{ }^{\circ}\text{C}$ was shown in Fig. 4. The SAP was degraded by about 81.45% at incorporated sandy loam soil in 56 days.

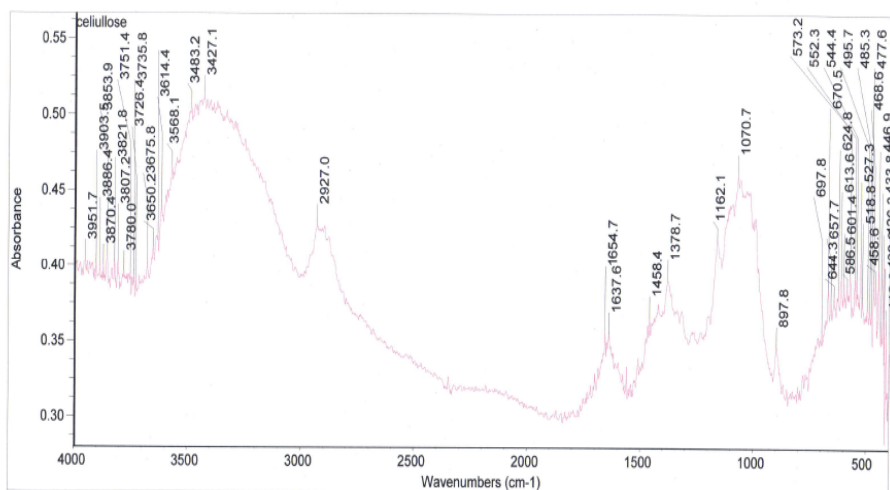


Figure 2. FT-IR spectra of un-graft cellulose

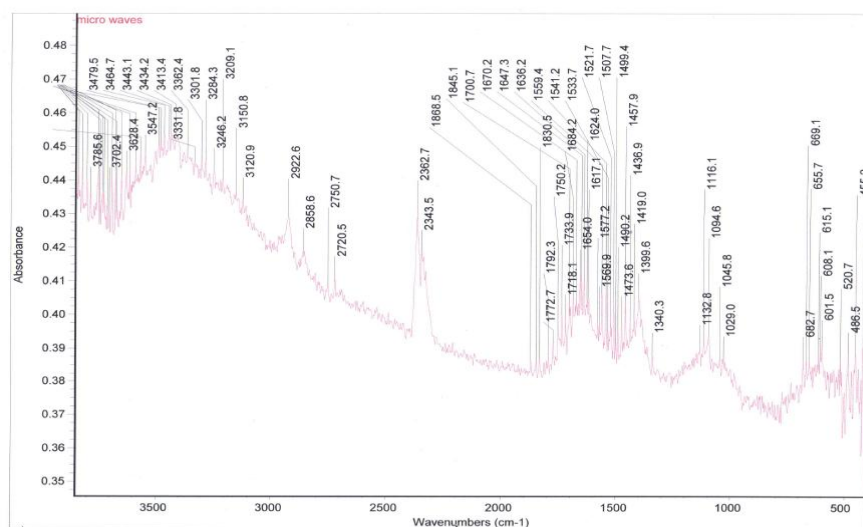


Figure 3. FT-IR spectra of graft SAP

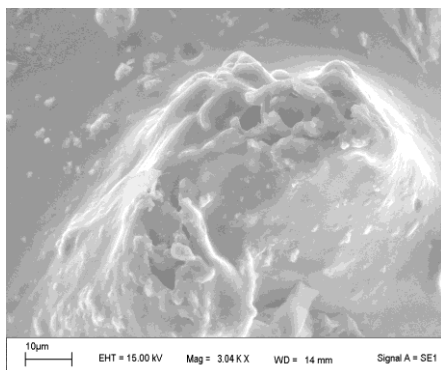


Plate 1. SEM diagram of grafted SAP

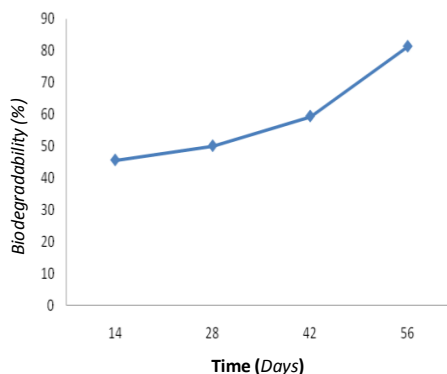


Figure 4. Biodegradability of Cellulose grafted SAP with time

Modified microwave oven was used to produce SAP Feng et al. (2010). The whole procedure had done under the microwave power of 160 W. In the present experiment, conventional heating as well as microwave irradiation were used for SAP development. 340 W microwave power was used for irradiation process according to results of Jing et al. (2011). According to the definitions of SAP, 400-1500 g of swelling recommended by it's own dry weight. Swelling ratio of the SAP has reached this super absorbent range. Further, due to high biodegradability within 56 days it can be recommended as environment friendly material. According to Feng et al. (2010) high temperature may improve the activation of some microorganisms and accelerate the biodegradation and furthermore, the porous structure allows microorganisms to go easily inside the network, which also favors its biodegradation.

Conclusions and Recommendations

SAP prepared by graft polymerization of acrylic acid onto the chain of cellulose from bagasse under microwave irradiation. The grafted product was biodegradable, had a porous structure and swelled 401 times on its own dry weight with distilled water.

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**AMPLIFIED FRAGMENT LENGTH POLYMORPHISM GENOME SCAN TO
REVEAL SELECTION SIGNATURES IN ASIAN TIGER MOSQUITO
(*Aedes albopictus*)**

G. H. Galhena^{1*}, M. A. Shaw² and R. Butlin³

¹*Department of Zoology, University of Colombo, Sri Lanka*

²*Institute of Integrative and Comparative Biology, University of Leeds, UK*

³*Department of Animal and Plant Sciences, University of Sheffield, UK*

Introduction

Aedes albopictus (Skuse), commonly known as the Asian tiger mosquito is recognized worldwide for its medical importance. It shows an aggressive daytime human-biting behavior and transmits many viruses, including dengue, LaCrosse, and West Nile. In Sri Lanka, the mosquito is largely responsible for the dengue epidemics in both peri-urban and rural settings. Since it was first recorded from India, it has undergone a dramatic global expansion facilitated by human activities. Due to its successive colonization of various habitat types, it is now listed as one of the top 100 invasive species by the Invasive Species Specialist Group, and is considered to be the most invasive mosquito species. This successful invasion of *A. albopictus* is mainly attributed to its ecological plasticity and strong competitive aptitude. Genomic scans are useful in identifying potential adaptive loci under selection at the genomic level. All loci across the genome are anticipated to possess similar demography and neutral evolution history of populations, including genetic drift and gene dispersal. If variation of a locus is beyond the genomic pattern with an unusual frame of higher genetic differentiation, it is deemed an “outlier locus” under natural selection. The outlier locus can be identified explicitly in the genes under selection and also in neutral flanking regions due to hitchhiking effects. Amplified fragment length polymorphisms (AFLPs) are reported to be the most efficient molecular approach to identify candidate genomic regions under selection. As such current study was aimed to identify the candidate loci under selection in *A. albopictus* in Sri Lanka, using an AFLP genome scan to understand the extent of its adaptive evolution in relation to current and historical mosquito control strategies.

Materials and Methods

Fourth stage larvae of *A. albopictus* were collected (n = 512) using standard dipping method in dengue endemic areas. Collections were made in a hierarchical manner on a logarithmic scale ranging from 1 m to 100 km. Accordingly 16 villages belonging to four regions in Sri Lanka (i.e. Colombo, Kandy, Anuradhapura and Pollonnaruwa) were sampled covering eight sites

* gayanihg@gmail.com

per village. Only four randomly selected larvae were taken from each container after identification up to the species level with the aid of morphological identification keys of *Aedes* larvae. Samples were preserved in absolute alcohol and were stored at 4 °C. Genomic DNA was extracted using phenol chloroform method. The AFLP procedure was carried out based on Vos et al. (1995) with following modifications: Genomic DNA was digested with *EcoRI* and *MseI* for 3 hours at 37 °C and immediately followed by ligation with *EcoRI* and *MseI* adapters overnight at 16 °C. Diluted ligated products were used as a template to conduct the pre-amplification. Each ligated product was amplified twice with two different primer combinations (Combination-1; *EcoRI*+A and *MseI*+C, Combination-2; *EcoRI*+T and *MseI*+C). The pre-amplification reaction was carried out for 20 cycles of 30 seconds (s) at 94 °C, 30 s at 56 °C and 1 min at 72 °C with a final extension of 10 min at 72 °C. Diluted (10x) pre-amplified products were used as templates for the selective PCR amplifications conducted with five primer combinations. Pre-amplification primer combination-1 was subjected to three selective amplifications (*MseI*_CAC, *MseI*_CGA and *MseI*_CTC with *EcoRI*_TCT), combination-2 was subjected to two selective amplifications (*MseI*_CTG and *MseI*_CAT with *EcoRI*_ATC). Each selective *EcoRI* primer was fluorescent-labeled at 5' end. Thermal cycling program was 94 °C for 2 min followed by 10 cycles of 94 °C for 20 s, 66 °C for 30 s with temperature decrease of 1 °C per cycle and 72 °C for 2 min. Another 25 cycles were carried out in the sequence of 94 °C for 30 s, 56 °C for 30 s and 72 °C for 3 min before a final extension of 30 min at 60 °C. All selective PCR products were immediately transferred and kept at -20 °C until they were run in the DNA sequencer within the course of 5 days. The selective PCR products were separated by electrophoresis on an ABI 3130 XL automated sequencer (Applied Biosystems) with a ROX-500-labeled internal size standard (Applied Biosystems).

Software GeneMapper 3.7 (Applied Biosystems) was utilized to collect and score fluorescent AFLP data. Altogether 557 polymorphic AFLP fragments were scored for presence (1) or absence (0) to establish the matrix of genetic identity of the sampled individuals. Outlier loci were tracked using the program Dfdist. Samples were run assuming a total of 32 demes representing the subpopulations occupying the 16 villages. The null distribution was generated, based on 10,000 simulated loci, with a mean *F_{st}* similar to the trimmed mean *F_{st}* calculated from the empirical distribution: Trimmed mean *F_{st}* is computed by removing the 30% highest and lowest *F_{st}* values observed in the empirical data set and thus is an estimate of the average "neutral" *F_{st}* value uninfluenced by outlier loci. Analyses were performed at 99 % confidence levels. Heterozygosity for each village was calculated using HICKORY v 1.0.4.

Results and Discussion

Forty nine loci were identified as potential outliers having F_{st} values outside the simulated 0.99 quantiles. The 10,000 simulated loci were built up in 10 different runs, since the program did not allow simulating them together for all 16 villages. Consequently, during each run 1000 loci were simulated and the empirical data set was compared with each set of simulated loci to detect outliers. An example plot corresponding to a single run is provided in Fig. 1. The simulated mean F_{st} value varied from 0.0503 to 0.0541 during the ten runs. The observed mean F_{st} from the Dfdist program for all 16 populations was 0.04876. Eighteen loci were consistently observed outside of the 0.99 quantiles across all runs and were regarded as true outliers corresponding to real selection signatures. They represent 3.2 % of the total polymorphic loci (557) detected. The fragment sizes and F_{st} values of these loci are presented in Table 1. The remaining 31 loci which did not show consistent outlier behavior during the ten runs might represent false positives. Such loci might exhibit a variation pattern in the tail of the neutral distribution simply by chance (type I error), due to statistical bias or due to some degree of departure between the simulated and the empirical data. However, such outlier loci due to chance are not expected to exhibit parallel trends in several comparisons while true outliers would stand out in each comparison. It is further possible that demographic events like bottlenecks to produce apparent selection signatures. The relatively high heterozygosity estimates which ranged from 0.24 to 0.26 within the tested 16 village populations make genetic bottlenecks an unlikely event. Hence, the 18 loci identified as true outliers might represent loci under directional selection or those that are linked to genomic areas that are under directional selection. Although the actual cause of this outlier behavior cannot be inferred without perusing these loci further, it is possible at least some of these loci could represent insecticide resistant genes. Other sources of diversifying selection which might have caused the outlier behavior might include odor reception, parasitic pressures and sexual selection.

Conclusions and Recommendations

This is the first report on detection of selection signatures by genome scan in *A. albopictus*. As observed 3.2 % of candidate loci are under diversifying selection, indicating that *A. albopictus* is indeed undergoing local adaptation. This might in part reflect the pressure put on the mosquito genome as a result of continues mosquito control campaigns. In order to discern the exact cause of selection pressure and to identify the genes under adaptive evolution, AFLP bands need to be sequenced and matched to their corresponding genomic locations which would be a future extension of this study.

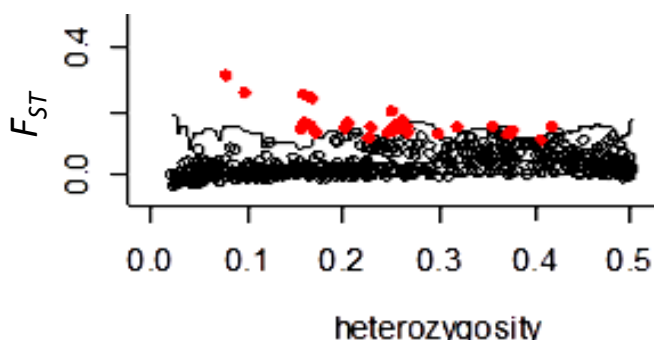


Figure 1. Example plot corresponding to a single run in Dfdist drawn between F_{ST} and heterozygosity showing outlier loci: Each dot indicates an AFLP locus while outliers are presented as red dots. The line immediately below the outlier loci represents the 99 % confidence interval.

Table 1. AFLP Loci under diversifying selection

Marker	Fragment length (bp)	F_{ST}
EcoR1_ ATC & Mse1_ CTG	136	0.2487
EcoR1_ ATC & Mse1_ CTG	225	0.1351
EcoR1_ ATC & Mse1_ CAT	190	0.1485
EcoR1_ ATC & Mse1_ CAT	281	0.1476
EcoR1_ ATC & Mse1_ CAT	292	0.1477
EcoR1_ TCT & Mse1_ CGA	192	0.1151
EcoR1_ TCT & Mse1_ CGA	300	0.1385
EcoR1_ TCT & Mse1_ CGA	422	0.3111
EcoR1_ TCT & Mse1_ CTC	105	0.1366
EcoR1_ TCT & Mse1_ CTC	131	0.1326
EcoR1_ TCT & Mse1_ CTC	147	0.1476
EcoR1_ TCT & Mse1_ CTC	182	0.1474
EcoR1_ TCT & Mse1_ CTC	193	0.2381
EcoR1_ TCT & Mse1_ CTC	228	0.2547
EcoR1_ TCT & Mse1_ CTC	232	0.1694
EcoR1_ TCT & Mse1_ CTC	316	0.1364
EcoR1_ TCT & Mse1_ CTC	382	0.1573
EcoR1_ TCT & Mse1_ CAC	269	0.1970

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RECYCLING OF SPENT GRIT INTO ASPHALT

N. R. G. Gamage^{1*}

¹*Holcim Lanka Ltd, Palavi, Puttalam, Sri Lanka*

Introduction

In the cleaning of the metallic surfaces in the ship building industry, there is used grit made of cast iron or steel. After usage in the sandblasting operations, the material loses the abrasive properties, entering to the waste category. In Sri Lanka, ship cleaning and equipment maintenance generates 4,614 tons of spent sandblasting grit per year. The spent grit, consisting of silica sand plus a small amount of slag-derived grit, has the physical characteristics of coarse-grained beach sand and also contained fragments of coatings. The spent grit has the potential for exhibiting hazardous characteristics since the coatings included lead-based primers, copper and butyltin-containing antifouling topcoats. Table 1 shows the heavy metal content in spent sand blasting grit.

Table 1. Heavy metal content in spent sand blasting grit (SGS Lab test report, 2013)

Heavy Metal	Result (mg kg⁻¹)
Arsenic (as As)	724
Cadmium (as Cd)	21
Chromium (as Cr)	271
Lead (as Pb)	499
Mercury (as Hg)	Not detected
Phosphorus content as P ₂ O ₅	1347

The spent grit was difficult to treat by conventional cement based solidification/stabilization methods. The lead and copper contaminants were contained in the organic portion of the paint chips, thereby limiting the ability of inorganic binders to stabilize the lead and copper. This study presents the feasibility of reusing the waste grit as a substitute for fine aggregate in the road construction industry. This is of both economic and environmental importance because by recycling them, the grit waste dumps disappear not polluting the environment.

* nadik.gamage@hotmail.com

Materials and Methods

Spent grit (Fig. 1) sample was sieved and particle size distribution is shown in Fig. 2.

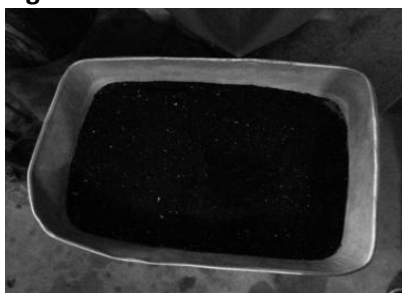


Figure 1. Spent Grit sample

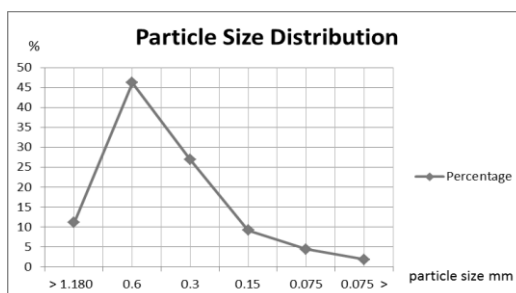


Figure 2. Particle size distribution of used sand blasting grit

Three asphalt samples were made by replacing 5 % of fine aggregate with grit. The quantity of Grit and rock was used as shown in Table 2. 4.76 % bitumen content was selected as it was the optimum bitumen percentage for that type of rock.

Table 2. Quantity of grit, rock and bitumen in asphalt

Bitumen		Aggreg ate type	Total aggregate weight	Sieve Size (mm)									
% weight				20	10	5	2.4	1.2	0.6	0.3	0.15	0.075	Pan
4.76	57.1	Rock	1082 (90%)	11	201	199	205	81	80	71	89	84	60
		Grit	60 (5%)						32	28			
Total Aggregate weight			1143										
Total Asphalt			1200										

Marshall Stability and flow of asphalt samples (Fig. 3) was measured and calculated.

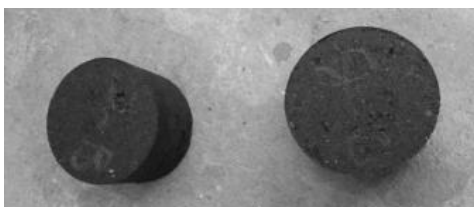


Figure 3. Asphalt sample

Toxicity Characteristic Leaching Procedure (TCLP) was performed for asphalt sample. The aim of this work has been to investigate leaching of asphalt that

may have an influence on the release of trace elements such as chromium, zinc and manganese, magnesium that are regarded as pollutants.

Results and Discussion

Based on the results in Table 3, it was observed that 5 % (weight percentage of total mixture) of grit can be fed into asphalt by maintaining required parameters of asphalt.

Table 3. Marshall Stability and Flow of Asphalt sample

Specification No : ASTM - D - 1559 -76											
No. of Blows : 75, each face											
Sample No:	Bitumen % by wt of Mix	Specimen height (mm)	Specimen wt.			Bulk volume (cm ³)	Bulk Specific Gravity (gcm ⁻³)	Marshall Stability			Flow (0.25 mm)
			in air (g)	Saturated Surface Dry (g)	in water (g)			Measured (kN)	Correc-tion Factor	Corrected (kN)	
A	4.76	60.0	1192	1192.4	707.2	485.2	2.46	14.45	1.09	15.75	12
B	4.76	59.0	1187	1187.8	703.9	483.9	2.45	14.91	1.09	16.25	12
C	4.76	60.0	1187	1188.0	705.1	482.9	2.46	12.48	1.09	13.60	6
Average							2.457			15.20	10
Marshall Stability / kN = Min 8; Marshall Flow/0.25mm = Min 8 Max 16 (As per RDA specifications)											

There are four bins in asphalt plant having less than 4 mm, 7.5 mm, 16 mm and 24 mm particle size respectively. Spent Grit can be fed into bin 1 (less than 4 mm particle) using front end loader by 13 % of total weight of bin 1. Balance percentage would be normal 87 % fine aggregate as rock.

Average Asphalt production per day (average plant) = 500 MT
 Grit consumption per day = 25 MT
 Number of plant working days per year = 200
 Grit consumption per year (single average plant) = 5,000 MT

Based on the above results, total ship building industry in Sri Lanka spent grit would be consumed by one average plant itself. There are five layers of road construction; 200 mm thickness of Gravel layer is laid on the soil first and then 150 mm of Aggregate Base Coat (ABC) is spread on it. After that there is a prime coat which is a bitumen layer. Then Tack coat is applied and after that asphalt is put on it.

Bitumen is a binding agent and it is insoluble in water. So Tendency of leaching of asphalt is very low. Furthermore, bitumen is highly impermeable to the passage of water. As asphalt layer would be above the bitumen layer (prime coat) it is confirmed that there is no tendency of any leachate coming out of asphalt.

Table 4. TCLP Test Results (ITI Lab test report, 2013)

Test/unit	Test method	Test values	Limit of
Lead (as Pb)	AAS/Flame	Not	0.1
Cadmium (as		Not	0.02
Chromium (as		Not	0.05
Copper (as Cu)		Not	0.05
Barium (as Ba)		Not	0.5
Antimony (as		Not	0.5
Zinc (as Zn)	AAS/VGA	0.27	
Arsenic (as As)		Not	0.001
Selenium (as Se)		Not	0.001
Mercury (as Hg)		Not	0.001

Note: AAS - Atomic Absorption Spectrometry VGA - Vapor Generation Accessory

As per results in Table 4, only Zinc was detected as leachable metal tested by the Toxicity Characteristic Leaching Procedure (TCLP) but it is well below the regulatory limit. Regulatory limit for same is 5 mg L⁻¹ maximum. So the leachable metals tested by the TCLP test did not exceed regulatory limits.

Conclusions and Recommendations

Based on the results of Marshall Stability test and TCLP test, it was concluded that 5 % of spent grit can be fed into asphalt by maintaining required physical performance characteristics of asphalt as well as disposal the spent grit environmental friendly manner.

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EFFECT OF NITROGEN ON GROWTH AND OIL CONTENT OF *Chlorella vulgaris* FOR BIODIESEL PRODUCTION

A.M. Gammanpila^{1*}, C.P. Rupasinghe¹ and S. Subasinghe²

¹Department of Agriculture Engineering, Faculty of Agriculture, University of Ruhuna, Mapalana, Kamburupitiya, Sri Lanka

²Department of Crop Science, Faculty of Agriculture, University of Ruhuna, Mapalana, Kamburupitiya, Sri Lanka

Introduction

Petro-diesel fuel has been used as a source of energy since many years. However, it is understood that the petroleum reserves are to be depleted in less than 50 years at the rate of present consumption. Further, the use of mineral oil has several negative environmental effects, such as emission of greenhouse gases.

In recent years biodiesel has received considerable attention as a biodegradable and renewable source of energy. It can be produced from renewable biomass by transesterification of triglycerides. Production of biodiesel from microalgae is a newly emerging field because of their high oil content and rapid biomass production.

Microalgae are photosynthetic microorganisms which convert sunlight, water and CO₂ to sugars, from which macromolecules such as lipids and Triacylglycerols (TAGs) are synthesized. Some of the species of microalgae already possess high oil concentration and they can be manipulated to produce more oil. A number of factors have been shown to influence the oil concentration of algae, such as nitrogen deficiency, phosphate limitation and temperature fluctuation. Nutrient deficiency, on the other hand, is a major limiting factor in the cultivation of microalgae as it directly impacts on the cellular metabolic activities and biochemical constituents, thereby creating a physiological imbalance which ultimately results in lower photosynthetic activity and slower growth.

The present study was conducted to investigate the growth response and oil content of a green alga, *Chlorella vulgaris*, by varying the concentrations of nitrate (NaNO₃) in the growth medium with an aim to obtain optimum NaNO₃ concentration for biomass cultivation for biodiesel production.

Materials and Methods

The pure *Chlorella vulgaris* cultures were grown in 25 ppt (parts per thousand) saline water in an outdoor culture system under protected house for 32 days growth period. The medium used for cultivation was Guillard and Ryther's modified F/2 media. The original nitrogen source concentration in

* shanigam@yahoo.com

the Guillard and Ryther's modified F/2 media was 0.075 gL^{-1} of NaNO_3 . The experiment was performed with three treatments, such as; half concentration of original, original concentration and double of the original concentration. The cultures were aerated with mechanical aerators. All the glassware and media were sterilized prior to inoculation. All the experiments were carried out in triplicates.

Cell growth analysis: Optical density (O.D) measurement at 665 nm was used to monitor cell growth by UV/visible spectrophotometer (Shimatzu, Japan) three times per week. All cultures were initiated with an O.D. of about 0.18. Based on the constructed calibration curve ($R^2 = 0.9428$), the numbers of cells were determined with relevant to the measured O.D.

Oil extraction: The cells were harvested at the stationary phase by chemical flocculation using 0.1 g of $\text{FeCl}_3 \text{ L}^{-1}$. The cells were washed once with distilled water and dried in an oven for 6 hours at 105°C for dry weight estimation. The oil content of the dried algae samples were then determined using Soxhlet apparatus.

Statistical analysis: The statistical analysis was done by using Minitab 14 version. One-way Analysis of Variance (one-way ANOVA) was performed at 95 % level of probability in order to test the significance differences of oil content and growth performances under different NaNO_3 concentrations. When the test resulted $p < 0.05$ a Turkey post-hoc test was performed for pair wise comparisons.

Results and Discussions

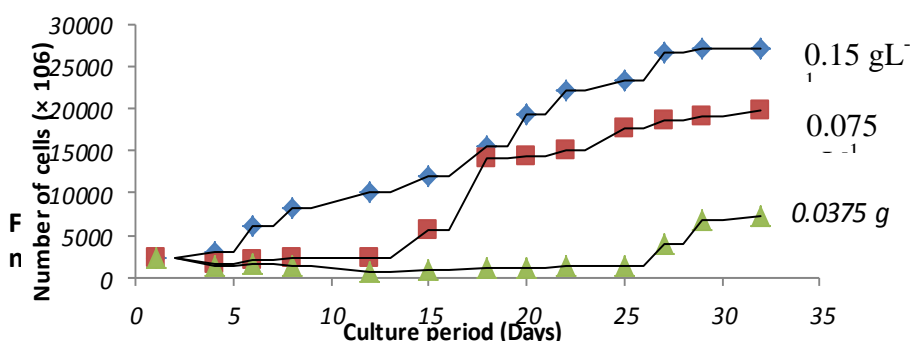


Fig. 1 shows the growth curve of *Chlorella vulgaris* grown in Guillard and Ryther's modified F/2 medium with different concentrations of NaNO_3 in the medium.

As evident from the graph, by increasing the NaNO_3 concentration, growth of *Chlorella vulgaris* also increased. At day 32, maximum number of cells (27286×10^6) was recorded in the culture with double the concentration of nitrate (0.15 g L^{-1}). Number of cells in the culture medium with half of the original nitrate concentration (0.0375 g L^{-1}) decreased initially and enters to lag phase after 25 days. Table 1 shows the dry biomass and lipid extraction results of the *Chlorella vulgaris* cultures grown in different nitrate concentrations.

Table 1. Dry biomass, oil content and oil yield of the *Chlorella vulgaris* cultures grown in different NaNO_3 concentrations

NaNO_3 Concentration (g L^{-1})	Dry biomass (g L^{-1})	Oil content (%)	Oil yield (g L^{-1})
0.0375	1.14 ± 0.07^a	21.01 ± 1.09^a	0.24 ± 0.01^a
0.075	1.66 ± 0.14^b	18.30 ± 0.26^b	0.30 ± 0.02^b
0.15	2.32 ± 0.08^c	13.83 ± 0.98^c	0.32 ± 0.02^b

Note: ^{a, b, c} indicate the values which were significantly different at $p < 0.05$.

Dry biomass and the percentage oil contents in the cultures with different NaNO_3 concentrations were significantly different ($p < 0.05$). Cultures with highest nitrate concentration (0.15 g L^{-1}) resulted significantly a higher dry biomass but significantly lower percentage oil content than the other cultures. A significantly highest oil accumulation and lowest dry biomass was recorded when the cultures were grown at half the original nitrate concentration (0.0375 g L^{-1}). However, the oil yield at half the original nitrate concentration was significantly lower than other treatments.

According to the results, *Chlorella vulgaris* growth is directly proportional to the concentration of NaNO_3 in the medium. As nitrate concentration increased in the medium, enhancement in biomass concentration was recorded and an increasing trend is observed in oil content as the nitrate concentration decreased. This result is in accordance with that of Nigam et al. (2011). They reported a loss of biomass and higher oil accumulation when green alga, *Chlorella pyrenoidosa* was exposed to nitrogen deficient conditions. Further, it was reported that, transferring cells to nitrogen-free medium, with no nitrogen source also led to an increase in oil content of 4% against control. It has earlier been found that under nitrogen starvation conditions, nitrogen containing macromolecules and carbon reserve compounds like carbohydrates and fats are accumulated.

According to the results, it can be concluded that, nitrogen starvation in the culture medium can enhance the oil content of *Chlorella vulgaris* but lower the growth rate and hence decrease the final oil yield.

Acknowledgement

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**MOLECULAR CYTOGENETIC CHARACTERIZATION OF THE FIRST
REPORTED SRI LANKAN CHILD WITH A *DE NOVO* 9P INVERTED
DUPLICATION (p13.3p23)**

I.U.H. Godapitiya¹, I. Kariyawasam¹, V. Udalamaththa¹, U.N.D. Sirisena¹ and
V.H.W. Dissanayake^{1*}

¹*Human Genetics Unit, Faculty of Medicine, University of Colombo*

Introduction

Compared to other rare chromosomal disorders, duplication of the short arm of chromosome 9 (partial trisomy 9p) is not uncommon. With the first reported case in 1970 more than 150 patients have been reported so far. After abnormalities in chromosomes 21, 18, and 13, trisomy 9p is the fourth most common chromosome anomaly in a live-born. In most reported patients, the trisomic segment of 9p duplications was transmitted from a parent carrying a reciprocal balanced translocation and only a few were due to *de novo* duplications. Patients with partial trisomy of the short arm of chromosome 9 often present a wide spectrum of phenotypic features including developmental delay, craniofacial abnormalities such as bulbous nose, hypertelorism and limb abnormalities with small nails and fifth-finger clinodactyly. The objective of this study was to perform cytogenetic molecular characterization of a child having dysmorphic features, intellectual disability, delayed bone age, development and speech.

Materials and Methods

A four year old Sri Lankan male child who presented to the Human Genetics Unit with delayed bone age, development and speech with intellectual disability and dysmorphic features was clinically evaluated. Cytogenetic studies were performed after obtaining ethical approval from the ethics review committee, Faculty of Medicine, University of Colombo. Peripheral venous blood samples were obtained from the child and his parents after obtaining a written informed consent. Metaphase chromosome spreads preparation from peripheral blood lymphocyte cultures and GTL-banding was performed according to standard methods. Thirty metaphase spreads of each parent were captured and analyzed along with twenty metaphase spreads of the child. The maximum band resolution achieved in the child's spreads was 550 according to the International System of Cytogenetic Nomenclature 2009. Fluorescent *in situ* Hybridization (FISH) technique was performed on the metaphase chromosome spreads of the child using gene/region specific FISH probes (Empire Genomics) according to the manufacturer's protocol. The

* vajirahwd@hotmail.com

initial cell culture was harvested using standard cytogenetic protocol. The fixative (Carnoy's 3:1 methanol:acetic acid) in sample tube was changed until supernatant was colourless, then refixed in fresh fixative prior to slide preparation. Three drops of suspension were added to the slide in a vertical angle. The slides were gently rotated to make a thin cell suspension and were kept parallel until a grainy appearance was observed to drain the excess suspension. The BAC clones used for the specific genes were RP11-627M21 for *DMRT 1* gene located at 9p24.3 and RP11-145E14 for *RMRP* gene located at 9p13.3. The probes were fluorescently labeled from the time of manufacture with spectrum Green and spectrum Red respectively. From each of the probe mixtures, 10 µL were added on to the slides (2 µL probe + 8 µL hybridization buffer) separately. A clean 22 × 22 mm cover slip was applied on to each slide and the edges were sealed using rubber cement. The probes were hybridized with metaphase chromosomes in a ThermoBrite Hybridizer; denaturation at 73 °C for 2 minutes, followed by hybridizing at 37 °C for 16 hours. After the hybridization, the slides were taken and the cover slips were removed and placed in a pre-warmed WS1 (0.4xSSC/0.3 % NP-40) at 73 °C solution and left to stand in WS1 (agitating ~10sec) for exactly 2 minutes. Then the slides were transferred to WS2 (2xSSC/0.1 % NP-40) at room temperature for 1 minute. The slides were dried in the dark, and counterstained with 10 µL of 4', 6-diamidino-2-phenylindole (DAPI) and covered with 22 × 22 mm cover slips. After 15-30 minutes the slides were visualized and the images were captured using the epifluorescence microscope.

Results and Discussion

The following dysmorphic features were noted: strabismus, low set anteverted large ears (bat ears), broad nasal root, short philtrum, a mouth with downturned corners with a prominent lower lip, short and broad hands with short fingers with 5th finger clinodactyly, bilateral simian creases and micropenis with bilateral testes (Fig. 1). Cytogenetic analysis detected a karyotype of 46, XY, add (9pter) (Fig. 2).

All thirty metaphase spreads of each parent showed a normal karyotype. Since it was only thirty spreads were analyzed from each parent, we cannot completely exclude mosaicism but can infer that the child's karyotype may probably be due to a *de novo* rearrangement as the parents did not show any phenotypic features. In the FISH analysis, the probe targeting the *RMRP* gene at 9p13.3 confirmed an inverted duplication 9p and the probe targeting the *DMRT1* gene at 9p24.3 terminal verified no terminal deletion/duplication (Fig. 3).

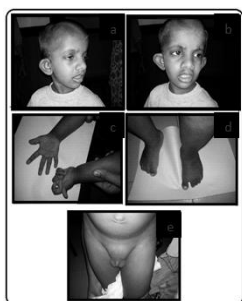


Figure 1. Clinical phenotype of the patient: (A) Side view showing frontal bossing (B) front view showing low set anteverted large ears (bat ears), broad nasal root, short philtres, a mouth with downturned corners with a prominent lower lip, (C) short and broad hands with short fingers with 5th finger clinodactyly and bilateral simian creases (D) short leg showing sandal gap (E) micropenis with bilateral testis.

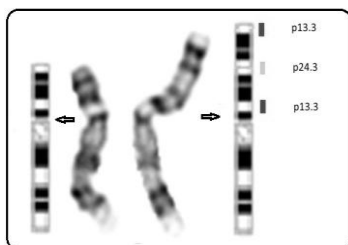


Figure 2. Identification of the 9p inverted duplication by conventional karyotyping: Ideogram of chromosome 9 showing the normal and the inverted duplicated region (red, green, red markers) and cut-out of the abnormal and normal chromosome 9 in G-banding at a resolution of 550 bands.

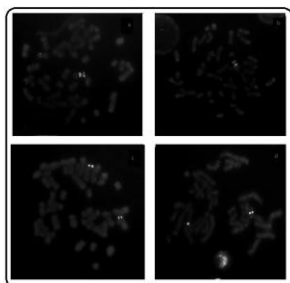


Figure 3. Characterization of the 9p inverted duplication by FISH analysis: (A) and (B) FISH analysis using spectrum red for RMRP gene at 9p13.3, (C) and (D) FISH analysis using spectrum green for DMRT 1 gene at 9p24.3. Arrow indicates the inverted duplication of 9p13.3 region.

Hence the final karyotype of the patient after molecular-cytogenetic characterization was 46, XY, ish dup(9)(p13.3p23). So far, *de novo* duplications of this chromosomal region have previously been described in approximately 15 patients worldwide. To our knowledge this is the first reported case of a child with *de novo* 9p inverted duplication in Sri Lanka. The 9p22.1; p23 region had been proposed to be the critical region for the 9p duplication syndrome phenotype. Duplications of 9p11.2 to 9p13.1 are believed to be natural chromosome variants with no reported abnormal phenotype. Published reports show that haplo insufficiency of *DMRT* 1, 2, and 3 results in gonadal abnormalities in the male. Even though in our patient the expected FISH signals for *DMRT* 1 gene were seen, we can suggest that due to the inversion of the duplicated segment which has involved the 9p terminal region may have disrupted one of the above mentioned genes in order to give the micropenis phenotype.

Conclusions and Recommendations

We can imply that the phenotypic features observed in this child are in concordance with the spectrum of clinical features seen in children with duplication of the 9p22.3-p23 critical region. Further genetic evaluation with advanced molecular cytogenetic techniques such as microarray along with FISH for break point analysis would be useful to confirm the diagnosis and to identify the affected genes.

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DO DIFFERENT COOKING METHODS AFFECT GLYCAEMIC INDEX OF RICE?

M. D. T. L. Gunathilaka¹, and S. Ekanayake^{1*}

*¹Department of Biochemistry, Faculty of Medical Sciences,
University of Sri Jayewardenepura, Nugegoda, Sri Lanka*

Introduction

Glycaemic index (GI) reflects the blood glucose response after a starchy meal. Many factors affect the GI including degree of starch gelatinization which depends on the amount of water added and the method of cooking. During cooking, heat and water soften the hard compact starch granules causing these to imbibe water, swell and eventually disrupt. As a result of that, individual amylose and amylopectin molecules are released leading to gelatinization. This makes starch more bioavailable for the action of enzymes causing increased glucose absorption. Rice is the staple food in Sri Lanka contributing to glycaemic carbohydrates and thus to the glycaemic response. Several studies had been conducted in the world, as well as in Sri Lanka to determine the GI of different rice varieties. Though it is believed that some basmati rice varieties imported to Sri Lanka have low GI, adequate research data on GI of these varieties is not available. Thus, the present study was conducted to evaluate the GI and the effect of two different cooking methods on GI of two imported basmati varieties (Pakistan basmati rice:-PBR; Indian basmati rice:-IBR) commonly purchased by Sri Lankans.

Materials and Methods

After following an unstructured interview type market survey, two basmati rice varieties were selected for the study IBR and PBR. GI of the two rice varieties was determined using the standard method. The GI study was designed as a randomized cross over study. Volunteers ($n=10$) were advised to undergo 8-10 hour fasting the day before each test day. A dietary recall of the previous night was taken from each volunteer on each test day due to secondary meal effect and the palatability and adequacy of portion size of two basmati rice varieties were recorded. About 200 μ L of fasting blood sample was taken from the participants by finger prick using an Accu Check meter. Finger tips were wiped thoroughly with surgical spirit and disposable lancets were used to prick fingers.

Two basmati rice varieties were cooked separately in a rice cooker (Panasonic) and microwave (LG) by adding water (1 cup of rice (110 g): 1 cup of water (150 mL).

* sagarikae@hotmail.com

The test meal (IBR and PBR) containing an available 50 g carbohydrate portion (IBR:-147 g and PBR:-151 g) to be consumed within 10-15 minutes with 250 mL of water was provided. Further finger prick blood samples (as above) at 30, 45, 60, 90 and 120 minutes were obtained. Eppendorf tubes containing dried residues of NaF (10 µL) were used for blood collection. Blood glucose was analyzed using a glucose kit (GOD-PAP/BIOLABO–France). Likewise, test meal was given on two different days to the same participants by keeping a three day wash out periods.

Above procedure was followed with the standard. Glucolin (gsk Glaxco Wellcome Ceylon Ltd, Sri Lanka) that contained dextrose monohydrate was used as the standard food and 55 g corresponding to 50 g available carbohydrate was dissolved in 250 mL and given on two different days to the same participants by keeping a three day wash out period. The following equation was used to calculate the Glycaemic Load (GL) of each individual and finally GL of a food was calculated as an average of ten values.

$$GL = \left[GI \times \frac{\text{weight of carbohydrate in one normal serving portion}}{100} \right]$$

The data was analyzed using the Statistical Package for Social Science (SPSS) Software (19th version) and Microsoft office Excel 2007. Chemical compositions are presented as mean \pm SD and the GI values as mean \pm SD. The results were analyzed using independent sample t-test by using SPSS (19th version) with 95 % taken as confidence interval.

Results and Discussion

The GI values of IBR cooked in the rice cooker (GI=54 \pm 8) or microwave (GI=43 \pm 6) belonged to low GI category (Table 1). These values are compatible with the values obtained from another study conducted by Srinivasa et al. (2013) who reported that the GI of an Indian basmati rice variety cooked in an electric cooker that belonged to low GI (54.93) category. The GI values of PBR cooked in rice cooker (GI=64 \pm 12) or microwave (GI=56 \pm 14) belonged to medium GI category (Table 1). Henry et al. (2005) studied commercially available four basmati in the UK and observed that the GI of all four basmati rice varieties belonged to medium GI (GI values- 52, 57, 67 and 69) category. Thus, present study results are compatible with the reported data for basmati rice varieties.

The GI values of the two rice varieties cooked using the rice cooker or microwave were not significantly different ($p < 0.05$). However, there was a percentage reduction in GI values in PBR (12.5 %) and IBR (20.4 %) when cooked in a microwave oven compared to rice cooker method. GL is the product of the GI and the amount of available carbohydrate of the actual

portion size. Irrespective of the rice variety and the cooking method, all the rice portions provided a high GL.

Table 1. Glycaemic indices of PBR and IBR cooked in a rice cooker and a microwave ($n=10$)

	PBR cooked in a rice cooker	PBR cooked in a microwave	IBR cooked in a rice cooker	IBR cooked in a microwave
Mean GI (\pm SD)	64 \pm 12	56 \pm 14	54 \pm 8	43 \pm 6
Peaking time	45 minutes	30 minutes	30 minutes	30 minutes
% peak reduction (against glucose)	19%	20.4%	23%	24.3%
% reduction in GI compared to rice cooker	-	12.5%	-	20.4%
Glycaemic load	32	28	27	22

Conclusions and Recommendations

GI of PBR and IBR cooked in a rice cooker and microwave belonged to medium and low GI categories respectively. Microwave cooking reduces the GI.

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EFFECT OF HERMETIC CONDITIONS OF HERMETICALLY SEALED LARGE CAPACITY COCOONS FOR STORAGE OF SOYBEAN (*Glycine max*)

D.M.C.C. Gunathilake^{1*}, W.M.C.B. Wasala¹ and B.M.K.S. Thilakarathne¹

¹ *Research and Development Center, Institute of Post-Harvest Technology,
Jayanthi Mawatha, Anuradhapura, Sri Lanka*

Introduction

Studies conducted in Sri Lanka have shown that nearly 14 % to 18 % of soybean is lost during postproduction operations due to improper post-harvest techniques. These studies also found, major component of these losses occurred during storage practices. On the other hand, sometimes soybean price drops even less than cost of production at the harvesting season due to high supply. Hence, it needs to promote farmers to store their product till the off-season in order to overcome this problem. Hermetically sealed cocoon storage is one of the suitable storage methods for grains and pulses (durable crops) because it can overcome most of storage problems. However, they should be dried up to 12 % moisture content before hermetic storage (Donahaye et al. 1991). Many research studies carried out abroad and few studies conducted in Sri Lanka revealed that growth of fungus, insects and pests can be inhibited under the hermetic storage conditions; and also metabolic rate of pulse will be decreased due to lack of oxygen (Donahaye et al. 1991; Villers et al. 2009). No research studies have been reported in Sri Lanka for evaluation its impacts in altering of physical properties and quality parameters of soybean under commercial scale hermetic (modified atmospheric) storage in local climatic conditions. Therefore, this research study was conducted by storing soybean in commercial scale hermetic storage conditions for 7 months and compared the occurrence of quality changers in soybean seeds such as moisture content, germination percentage, nutritional properties (protein) and physical properties namely bulk density, thousand seed mass, kernel hardness and colour in comparison with soybean stored in conventional warehouse.

Materials and Methods

Soybean variety PB-01 was used for this study. Seed was dried up to 12 % moisture content. The 1220 kg of soybean seeds bagged in poly-sack was stored inside the cocoon and sealed. Cocoon was placed in warehouse and the same amount of soybean seeds bagged in poly-sack bags place warehouse as a conventional storage method as treatments. The 20 soybean samples as methods outlined by Bal et al. (1978) were used initially and after the storage trail to measure moisture content, germination percentage, nutritional

* champathi@hotmail.com

properties (protein) and physical properties namely bulk density, thousand seed mass, kernel hardness and colour as responses. American oil chemists' society (AOCS) method (AC 2-41 (14)) was adapted to analysis the moisture content of soybean. Standard germination test, 50 soybean seeds placed in germination paper wrapped with adequate water and covered with polyethylene sheet and kept room temperature for 7 days was performed. Oxygen level inside the sealed cocoons was measured using oxygen meter manufactured by Grain Pro Inc, USA once a two week till the end of the experiment. Inside temperature of hermetically sealed cocoon and warehouse were measured using thermo couples. Soybean seed color was measured by using Mini-scan XE plus Hunter Lab Colorimeter (L, a, and b values were measured). Compression test (yield stress) by seed hardness tester was carried out to measure seed hardness. Force at rupture was considered as the hardness. Five replicates were considered for compression test and the measured force were averaged. 1000 seed mass obtained by measuring mass of the 10 seeds for ten times at 3 replicates averaged and multiplying average weight by 100. Bulk density apparatus was used to measure bulk density of soy bean. Each treatment was replicated two times. Analysis of Variance (ANOVA) on Complete Randomized Design (CRD) was performed and treatment means were separated by the DNMT at $\alpha = 0.05$ level of significance.

Results and Discussion

Change of oxygen level and temperature inside the hermetically sealed cocoon: Fig. 1 shows the reduction of oxygen percentage inside the hermetic cocoon. Oxygen percentage of hermetic cocoon reduced from 20.5 % (ambient environment O₂ percentage) to 9.4 % during 7 months storage. The temperature fluctuation was also low inside the cocoon in comparison to warehouse.

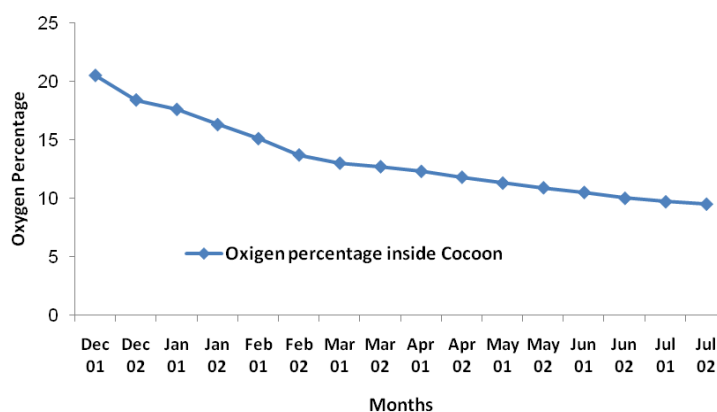


Figure 1. Change of oxygen% inside of the sealed cocoon during the experiment

Comparison of moisture content, germination percentage, nutritional properties and physical properties:

Table 1 shows the mean moisture content (MC) of samples. It was clear from the results that soybean seed moisture percentage was significantly changed in conventional warehouse samples in comparison to its initial moisture percentage and moisture percentage of hermetically sealed stored samples for 7 month. During the experimental period, dry climate (low relative humidity) was observed therefore seed moisture was evaporated. But seed moisture was not evaporated under sealed hermetic condition. Germination percentage (GM) of the cocoon sample after 7 months storage was shown a significant reduction compared to initial values and warehouse samples. Warehouse samples have shown higher germination percentage than cocoon samples. This result proves previous results obtain by Narayan et al. (1988), that seed viability can be lost under low oxygen storage conditions. Protein percentage (PRO) of the soybean seed has been protected under hermetic storage than conventional warehouse storage, because metabolic function of seed is reduced under low oxygen storage. However, protein content was reduced with time at both storage methods. It was not significant. Thousand seed mass (TSM) significantly reduced in conventional warehouse storage compared to hermetically sealed cocoon. Hermetically sealed cocoon preserved soybean seed mass due to slow metabolic rate of seeds and also weight difference of soy bean i.e. total initial weight to total final weight for 7 months also low in cocoon in comparison to warehouse storage. Growth of fungus, insect and pests were inhibiting under the hermetic storage conditions due to lack of oxygen and the metabolic rate of seed was also be decreased due to lack of oxygen (Donahaye et al. 1991; Villers et al. 2009).

Table 1. Change of seed parameters (responses) stored under different storage conditions

Storage method	MC %	PRO %	GM %	TSM	BD	Hd	L	SC a	b
Initial	11.2 ^a	36.31 ^a	96.12 ^a	146.5 ^a	775.2 ^b	7.04 ^a	54.15 ^a	3.70 ^a	15.34 ^a
Mean values									
Cocoon	11.1 ^a	36.04 ^a	44.45 ^c	141.7 ^b	790.9 ^a	5.85 ^c	53.34 ^a	3.25 ^a	15.36 ^a
Mean values									
Warehouse	9.92 ^b	35.12 ^a	70.32 ^b	130.4 ^c	789.2 ^a	6.35 ^b	51.98 ^a	4.14 ^a	15.87 ^a
Mean values									

Note: *Columns having same letter are not significantly difference at $\alpha = 0.05$ by DMRT

MC-moisture percentage, PRO-protein percentage, GM-germination, TSM-thousand seed mass, BD-bulk density, Hd-seed hardness, and SC-seed colour

Bulk density (BD) was increased in both cocoon and warehouse samples against its initial value. However, non-significant change of bulk density was observed between cocoon and warehouse samples for 7 months storage. Therefore, soybean seed mass increased compared to volume increased. Seed hardness (HD) values were reduced compared to its initial values in both storage methods and the hardness of the cocoon samples also reported low value than warehouse samples for 7 months storage. Soybean seed colour (SC) was reduced slightly from its initial value in both storage methods. However seed colour was also slightly lower in warehouse stored soybean in comparison to cocoon stored soybean for 7 months of storage.

Conclusions and Recommendations

Oxygen percentage of soybean stored cocoon reduced from 20.5 % (ambient environment O₂ percentage) to 9.4 % for 7 months. Temperature fluctuation inside the hermetic cocoon was very low. Change of moisture percentage during storage was not significant among storage methods. Germination was significantly low in cocoon sample. Protein percentage was slightly reduced in both storage methods. Seed mass was preserved in hermetic storage and seed hardness. Bulk density of soybean increased during storage and it was highly increased in hermetically sealed cocoon. Soybean seed colour was preserved slightly by sealed cocoon storage. Hermetically sealed cocoon storage has some advantages in comparison to conventional warehouse storage for large scale commercial soybean/pulse storage. However, it is recommended to maintain hermetically sealed condition throughout storage period to gain those advantages.

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EVENT CLASSIFICATION AND ABNORMAL EVENT DETECTION IN VIDEOS

**H.M.S.P.B. Herath^{1*}, P.H. Perera¹, W.S.K. Fernando¹, M.P.B. Ekanayake¹,
G.M.R.I. Godaliyadda¹ and J.V. Wijayakulasooriya¹**

*Department of Electrical and Electronic Engineering, Faculty of Engineering,
University of Peradeniya, Sri Lanka*

Introduction

Series of actions performed by an object forms the structure of a video event. For an example a man walking down a street could be treated as a video event and the event itself is defined by the characteristics of his motion throughout the video. Hence an event can be represented using a feature space which is a collection of features associated to an objects' motion such as object location, speed, direction etc. Event classification is the process of assigning an event identity to a newly observed event. It is used excessively in applications such as security monitoring, traffic surveillance, emergency detection systems and abnormal event detection systems etc. The paper proposes a supervised classification approach to classify such events using the principle of Dynamic Time Warping (DTW) and in the process to identify abnormal events using a conformity score.

Materials and Methods

In supervised classification (Zhang and Schuller 2012) set of event classes are predefined and representative templates of each event class is maintained. A distance measure is evaluated between an observed event and every event template to determine the class identity of the event.

Event Distance Measures:

Since events are defined using associated feature trajectories, event similarity can be evaluated in terms of temporal features which describe them. Temporal feature trajectories of tracked object location, speed or any informative feature could be extracted from a foreground estimation and tracking methodology similar to the one proposed in (Fernando 2014).

Use of conventional signal comparison processes like correlation, direct mean square error, and trajectory statistics comparison are inappropriate in such feature trajectory comparison since the length of compared trajectories would be different. This brings up the necessity of using more specific trajectory comparison methodologies in event detection applications. The method considered should be able to compare multi-dimensional feature trajectories of different lengths and generate. Moreover, it should reflect a clear scalar distance between the compared events. Since the comparison is

* spbh@gmail.com

carried out between two trajectories, it is important that the comparison mechanism captures trajectory shape and placement information.

Dynamic Time Warping (DTW):

DTW method is a method that focuses on the trajectory shape and placement during the comparison process removing certain non-linear variations in the time domain of the sequences. As explained by Muller (2007), DTW is an existing method that could be used in comparing temporal trajectories (signals) of different lengths. This method is used in finding an optimal temporal alignment between the compared two trajectories.

DTW for event comparison:

Employment of DTW for event comparison requires a cost function to be defined such that higher dimensional features could be incorporated. In handling this cost matrix is proposed using squared norm of the vector differences according to,

$$C(N - i, j) = \left| \overline{\text{Trajectory2}(i)} - \overline{\text{Trajectory1}(j)} \right|^2 \quad (1)$$

where N is the length of *Trajectory2*. For the explanation of the method consider the speed trajectories of two different events illustrated in Figure 1. The corresponding cost matrix calculated using the equation (1) is shown in Figure 2 where brightly shaded areas denote higher costs.

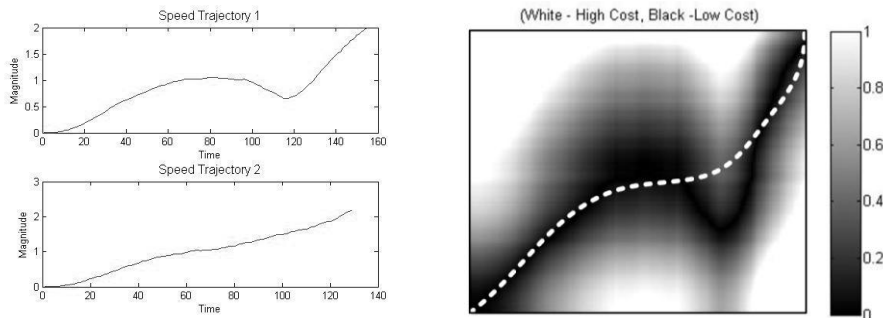


Figure 1. Two Speed Trajectories to be Compared using DTW

Figure 2. Alignment Cost Matrix used in DTW

Conformity score of two events is calculated based on a quantity, total alignment cost. In DTW the total alignment cost between the trajectories is defined to be the sum of the costs when moving from the bottom-left corner to the top-right corner of the cost matrix in a specified path. The cost associated with minimal cost path that is available is defined to be the minimum cost of alignment. As an additional constraint for the optimal path selection, the motion of optimization is restrained such that the path always moves forward in time (Muller 2007). A possible alignment path is illustrated in Figure 2 on the cost matrix. Conformity score is obtained by dividing the total alignment cost by an approximated length of the aligning path.

Event classification:

At periodic intervals an event is compared against the set of predefined event templates using the DTW method and corresponding conformity scores for each template is obtained. Each event is attributed to the event template that gives the minimum conformity score. Moreover, obtained conformity scores are compared against a predetermined threshold T , and if the obtained conformity score is greater than T , the corresponding event is classified as an abnormal event as in,

$$I = \begin{cases} \text{abnormal} & ; d_k > T \text{ for } \forall k \\ \text{argmin}_k(d_k) & ; \text{otherwise} \end{cases} \quad (2)$$

where d_k is the conformity score between the observed event and the k^{th} event template. I is the identified event. To determine the threshold T an analysis on a training data set is carried out. In the analysis, conformity score of arbitrarily selected events with training set population are calculated and conformity score histogram is constructed. The resulting histogram has two separable regions that correspond to “conformity scores with same event class” and “conformity score with different event classes”. Parameter T is selected to be equal to a conformity value that separates these two regions. Figure 3 illustrates how this process was carried out for a sample data set. The obtained value for T in this application was 11.

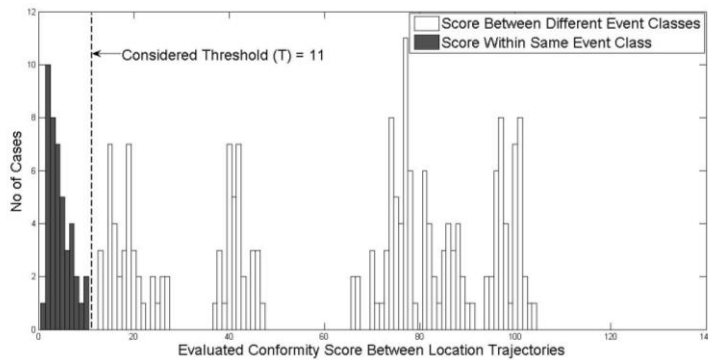


Figure 3. Process of Obtaining the Threshold (T)

Results and Discussion

The overall event classification was tested in an outdoor environment focusing on human motions. Figure 4(a) shows an image of the environment and Figure 4(b) shows two of the template trajectories for event classes that had multiple occurrences throughout the experiment. The experiment was conducted over 50 object location trajectories and multiple occurring event trajectories were imposed as the reference events. Figure 5 (a) & (b) illustrates frame sequences where each of the events is identified and indicated with number corresponding to the event. The representative numbers for events are in accord with the ones marked in Figure 4(b). Each of

the events (a) & (b) in Figure 5 undergoes an unclassifiable time period initially where the conformity score denotes them as unclassifiable according to equation (2) and they are denoted by "0".

Proper identification of the event class has been observed after this observation time period. Events that do not match any of the templates are classified as abnormal events according to equation (2) and they continue to be marked as "0" following the observation time.

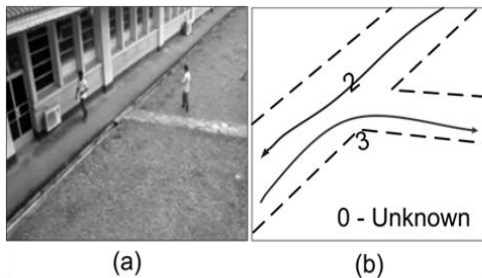


Figure 4. Two of the events templates

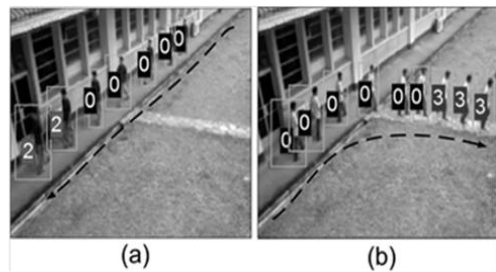


Figure 5. Frame sequences where the denoted event is classified correctly

Conclusions and Recommendations

The adoption of DTW into event classification problem enables to compare feature trajectories based on the shape of the trajectory accurately. This adoption gives better comparison results compared to standard algorithms due to the inherent ability of removing nonlinear temporal variations. The results obtained for a set of real life videos demonstrates that DTW accurately distinguishes events based on feature trajectories. Abnormal activities that differ from standard event templates can be identified using the conformity score threshold.

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VIRAL BURDEN AND DIVERSITY IN ACUTE RESPIRATORY TRACT INFECTIONS IN CHILDREN IN SELECTED AREAS OF SRI LANKA

**J.A.A.S. Jayaweera^{1,2*}, F. Noordeen², A.J. Morel³, F.N.N. Pitchai²,
H.S. Kothalawela², A.M.S.B. Abeykoon² and J.S.M. Peiris⁴**

¹*Department of Microbiology, Faculty of Medicine and Allied Sciences,
Rajarata University of Sri Lanka*

²*Department of Microbiology, Faculty of Medicine, University of Peradeniya*

³*Teaching Hospital, Gampola*

⁴*School of Public Health, University of Hong Kong, Hong Kong*

Introduction

Acute respiratory tract infection (ARTI) represents the most common acute illness evaluated in childhood. ARTIs range from the common cold, a mild, self-limited catarrhal syndrome to life threatening lower respiratory tract infections. Viruses account for most ARTIs (Meneghetti 2011). Several viruses are associated with ARTI and associated respiratory diseases. The most frequently reported viruses in newborns and children under 5 years are respiratory syncytial virus, parainfluenza types 1, 2, 3, adenovirus, influenza virus types A, B, corona virus, coxsackievirus, enterovirus, human boca virus and human metapneumovirus (Muller-Pebody et al. 2002).

Viruses cause majority of ARTI but only small percentage of these infections results in severe or fatal disease. Viral ARTI lead to secondary bacterial infections in healthy children by lowering immunity in the respiratory tract and that allows the invasion of bacteria (Graham 1990). There is evidence for marked seasonal variation in viral incidence, which is higher during the colder months in countries with temperate climates (Musher 2003). In countries with tropical climates the seasonality is variable according to the temperature-dependent local weather pattern such as humidity or rainfall (Muller-Pebody et al. 2002). Differences in the prevalence of viral infections can also be observed between community groups. Epidemiological studies on ARTIs of viral aetiology in the Sri Lankan paediatric population have not been performed. However, there is a recent study mainly related to influenza virus in patients attended for treatment for ARTI to North Colombo Teaching Hospital in the Western Province of Sri Lanka showing different seasonal patterns (Perera et al. 2010). By identifying seasonality pattern and the prevalence of viral causes in childhood ARTI, we will be able to take early preventive actions to assist to reduce childhood ARTI as well as associated morbidity and mortality. At the same time by identifying the viral aetiology, the number of consultations and irrational use of antibiotics can be reduced.

* jaas820703@yahoo.com

Materials and Methods

The ethical approval for the study was obtained from ethical and higher degree committees of the Faculty of Medicine, University of Peradeniya. This is a epidemiological study of inward paediatric patients (1 month to 5 years) with clinically diagnosed ARTI. Nasopharyngeal aspirates (NPA) of patients with ARTI were obtained at Teaching Hospital Gampola (THG) and the Professorial Unit, Anuradhapura (PUA) from March 2013 to July 2014.

Indirect immunofluorescence (IF) test by DAKO IMAGEN™ (UK) respiratory screening reagents were used to screen for 8 respiratory viruses. Polymerase chain reaction (PCR) was performed using virus specific primers on the IF negative NPA sediments for hMPV, hBoV and corona viruses. Basic descriptive analyses were done by means of central tendency.

Results and Discussion

Out of 430 from children with ARTI from THG, 262 (61 %) were males. Patients with bronchiolitis were 309 (72 %); bronchopneumonia was 73 (17 %), laryngotracheobronchitis was 34 (8 %) and pharyngitis was 9 (2 %). Mean age of inward patients with ARTI was 1.22 years (SD±1.08). Out of 385 children with ARTI from PUA, 268 (69.5 %) were males. Mean age of patients in the PUA cohort was 1.04 years (SD±0.93). In the PUA cohort, bronchiolitis was noted in 271 (70.5 %), laryngotracheobronchitis was noted in 29 (7.6 %), bronchopneumonia was noted in 26 (6.8 %), acute infective rhinitis was noted in 24 (6.2 %) and pharyngitis was noted in 12 (3.2 %) children. A few patients (n=55) had cyanotic congenital cardiac anomalies, 34 from PUA and 21 from THG. Sum of 172 (40 %) patients in THG were exposed to passive tobacco smoking and 26 (6 %) were exposed to dust following industrial air pollution. Whereas in PUA cohort, 85 (22 %) were exposed to passive smoking and 169 (44 %) were exposed to dust of road construction.

Out of 430 and 385 NPAs from THG and PUA, 156 (36.3 %) and 144 (37.3 %) were positive for viral screening using IFA. Following typing, RSV was found in 96 and 84 samples from TAG and PUA, respectively. Para influenza 1, 2 and 3 viruses were positive in 4, 21 and 3 from THG and 3, 15 and 5 in PUA, respectively. Adenovirus was positive in 20 and 22; influenza A in 3 and 2 and influenza B in 9 and 14 in THG and PUA, respectively. RT-PCR detected 11 hMPV. Six hMPV, 8 parainfluenza 1 and 18 parainfluenza 2 were positive from RSV positive samples in THG. hBoV and corona viruses were not detected in our study cohort either as primary pathogen or as co-infection with RSV.

Out of 300 respiratory virus positive cases 88 (61 %) and 118 (75.6 %) patients were on antibiotics in PUA and THG cohorts, respectively. There was a significant different noted in ARTI between males and females (P=0.01; α ,

0.05). Peak viral detection was found in both centres in May-July in 2013; May-June in 2014 in THG. A peak in April 2013 and in December-January 2014 was noted in PUA (Figs. 1 and 2)

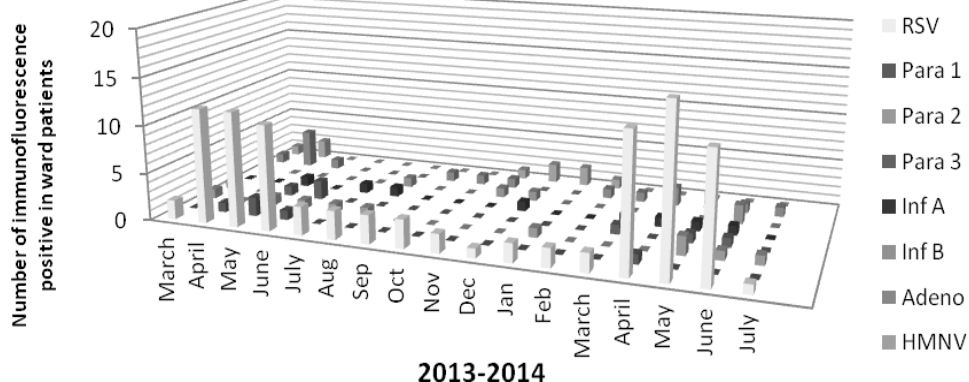


Figure 1. Number of IFA and PCR positive NPA with viral aetiology in inward patients with ARTI in THG from March 2013-July 2014

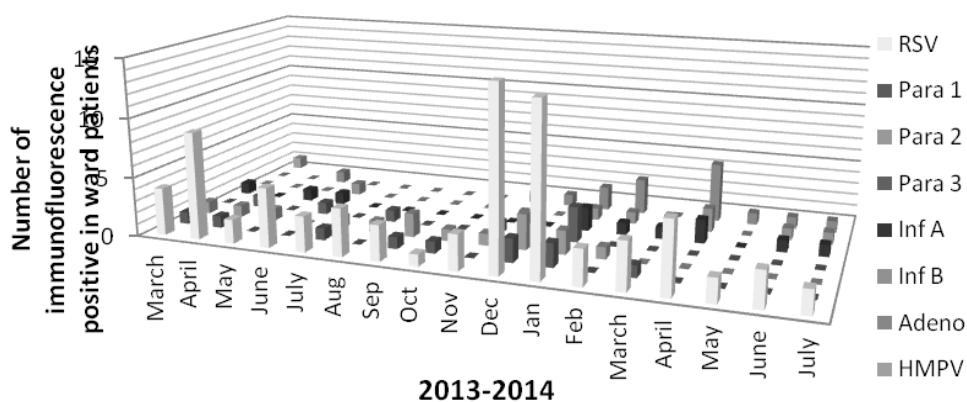


Figure 2. Number of IFA and PCR positive NPA with viral aetiology in inward patients with ARTI in PUA from March 2013-July 2014

Conclusions and Recommendations

Depends on IFA viral aetiology accounted for 36.9 % of the ARTI cases in THG and PUA. RSV is the most common virus responsible for childhood ARTI. Peak viral detection was found in both centres in May-July in 2013; May-June in 2014 in THG. A peak in April 2013 and in December-January 2014 was noted in PUA.

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**USE OF INFORMATION ON RAINFALL BEHAVIOUR TO ASSIST TIMELY
MANAGEMENT OF AGRONOMIC OPERATIONS IN RUBBER
PLANTATIONS IN THE AGRO ECOLOGICAL REGION – WL1a**

P. W. Jeewanthi¹, W. Wijesuriya^{1*} and J. K. S. Sankalpa²

¹*Biometry Section, Rubber Research Institute of Sri Lanka, Dartonfield, Agalawatta*

²*Agricultural Economics Unit, Rubber Research Institute of Sri Lanka, Dartonfield,
Agalawatta*

Introduction

Agronomic operations in rubber plantations from planting through felling are closely linked with the amount and pattern of rainfall. Hence, rainfall can be regarded as the major climatic factor which directly affects the productivity of rubber plantations. Observational evidence indicates that changes of climatic factors such as rainfall, have significantly affected diverse set of natural and human systems and activities in many countries (Hansen 2002). This necessitates detail analyses of rainfall for effective planning of agro-management practices in rubber plantations. Understanding the rainfall patterns provide information on adverse conditions with respect to both dry and wet conditions in a particular area.

Yogarathnam (2001) reported that the ideal annual rainfall for rubber falls within the range of 1650 mm to 3000 mm. The tree performance is severely affected when rainfall over a six months period recorded less than 500 mm especially when rainfall is not uniformly distributed over the year. It seems that rainfall is not a limiting factor in the rubber growing areas of the Low Country Wet Zone of Sri Lanka. However, no detail studies were found in literature in the recent years giving emphasis on comparing the crop calendar with the existing rainfall patterns to suggest appropriate dates for different agro-management practices for rubber cultivation. Hence, studying the rainfall pattern, probability of occurring a wet day, number of wet days and length of wet and dry spells within a particular period is also of immense importance as rainfall amount in planning agricultural operations in Rubber cultivation. This study was conducted as an initial step to identify the rainfall behavior in three major rubber growing areas employing Markov chain models, a classical model to express a stochastic process (Gabriel and Neumann 1962).

Materials and Methods

Data: Daily rainfall values of three meteorological stations in the Agro Ecological Region, WL-1a namely; Agalawatta, Avissawella and Ratnapura

* wasanawij@hotmail.com

were used covering the period from 1980 to 2010. A wet day (a dry day) is defined as one with ≥ 3 mm (<3 mm) of rainfall per day according to the water requirement of Rubber.

Markov Chain Model and Estimation of Parameters: The process of occurrence of wet(W) and dry(D) days can be described by a 2-state Markov chain with wet and dry days as the two states. The transition probability matrix for a given standard week P describing the 2-state Markov chain model is;

$$P = \begin{pmatrix} P_{00} & P_{01} \\ P_{10} & P_{11} \end{pmatrix}$$

with $P_{00} + P_{01} = 1$ and $P_{10} + P_{11} = 1$, where P_{00}, P_{01}, P_{10} and P_{11} are the transition probabilities; P_{00} = probability of occurring a dry day given that the preceding day was a dry day [$P(D/D)$], $P_{01} = P(W/D)$, $P_{10} = P(D/W)$ and $P_{11} = P(W/W)$. As the next step, each day in a year was grouped into the categories; D/D, W/D, D/W and W/W. This procedure was adopted subsequently for all the years considered for the analysis. Then the maximum likelihood estimates of probabilities; P_{00}, P_{01}, P_{10} and P_{11} were obtained for the 52 standard weeks.

According to Cox and Miller (1965), after a sufficiently long period of time, it is expected to settle down to a condition of statistical equilibrium with steady state or equilibrium probabilities which are independent of the initial conditions. So for a given standard week (SW), the limiting probabilities are given by $\pi_0 = \frac{1-P_{11}}{1+P_{01}-P_{11}}$ and $\pi_1 = \frac{P_{01}}{1+P_{01}-P_{11}}$ corresponding to dry and wet days, respectively (Ross, 1996). According to the above proof, probability of day in a given SW being a wet day is π_1 . Let, X= number of wet days within a given SW, $X \sim \text{binomial}(n, \pi_1)$ with $n=7$ or 8 (8 in only two occasions). Therefore, the expected number of wet days ($E(X)$) for a given SW is defined as, $E(X) = n \cdot \pi_1$. The dry spell (or run) lengths are very important statistical descriptors of dry periods in a rubber growing area. Let, D is the length of dry spell in a given SW, $D \sim \text{Geometric}(\pi_1)$. Therefore, the expected lengths of the dry spells ($E(D)$) for a given SW is defined as, $E(D) = 1/\pi_1$.

Results and Discussion

As depicted in Fig. 1, the probability of a day in a given SW being a wet day in Agalawatta and Ratnapura is very much similar over the year. However, a marked deviation was found for Avissawella, where the mid-year dry period is more pronounced. The estimated number of wet days per SW also showed the same sequence (Fig. 2). In Avissawella, the number of wet days equals 3 in 22 weeks out of 52. In Agalawatta and Ratnapura, the number of wet days equals 4 in 24 and 25 weeks out of 52, respectively. None of the weeks had 5 wet days in Avissawella (Table 1). The Expected dry run if the beginning of a SW is given a dry day is depicted in Fig. 3. Accordingly, during the initial part of the year, in Avissawella a relatively dry period is observed compared to the

other two areas. Although less in magnitude, the similar sequence is observed during the mid-year dry period.

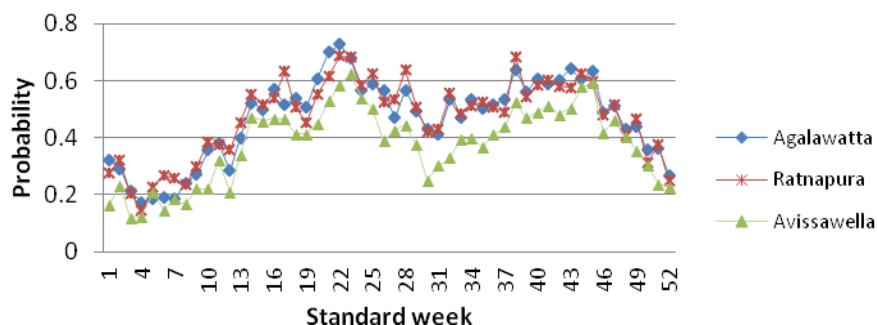


Figure 1. Probability of a day in a given standard week being a wet day

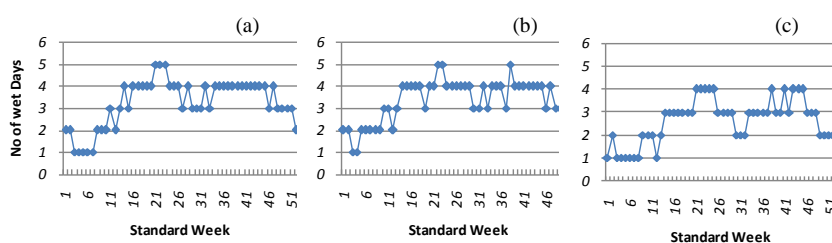


Figure 2. Expected number of wet days per standard week in (a) Agalawatta, (b) Ratnapura and (c) Avissawella

Table 1. No of wet days per standard week in Agalawatta, Ratnapura and Avissawella

No of wet days per week	No of weeks out of 52 standard weeks		
	Agalawatta	Ratnapura	Avissawella
1	5	2	8
2	7	10	12
3	13	12	22
4	24	25	10
5	3	3	0

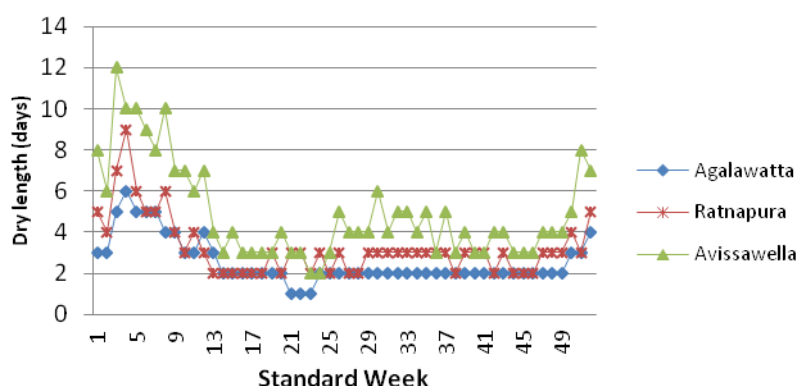


Figure 3. Expected dry run if the beginning of a standard week is given a dry day

Conclusions and Recommendations

A marked difference is observed in Avissawella for probability of a day in a given standard week being a wet day, expected number of wet days and expected dry run. Planting season which coincides with 16th to 22nd week should be strictly followed in Avissawella area. A slight delay in planting which frequently happen due planting material demand will have a relatively low risk in Agalawatta and Ratnapura. Further, more emphasis should be paid on moisture conservation measures during the initial part of the year and the mid-year in Avissawella compared to the other two areas. Since it records more number of expected wet days in Ratnapura and Agalawatta during the year, more attention should be paid on arresting rain interference for harvesting using rain guards.

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ALTERED APPEARANCES: SOCIAL MEANINGS OF DRESS DURING THE KOTTE PERIOD

P.V.M.Karunaratne^{1*}

¹*Department of Textile and Clothing Technology, Faculty of Engineering,
University of Moratuwa, Moratuwa*

Introduction

Fashion of dress gives social meanings. The social meanings of dress are embedded in various social contexts or various circumstances. These meanings are constantly altered by the appearances in which they are found in social/ historical context which frame space and time. The objective of the study is to find out how the meanings are embedded and altered by the appearances in which they are found in social contexts framing space and time. The historical context of the Kotte era (1411-1597 AD) was thoroughly explored through its significant dresses and analyzed theoretically in order to understand social meanings of new styles that were reconfigured elements of clothing created by the royalty of that period after they embraced Christianity.

Materials and Methods

Trouser (*pantaloons*), coat (*cabaya*), jacket (*hettaya*) short jacket with collar (*manthe hettaya*) and articles of clothing (items) such as attached collar (*tippet*), were observed and analyzed. The systematic pictorial analysis reveals many new meanings that were created, which influenced transmission of their “context”. The conceptual framework of the study is based on the qualitative research method. Sequence of in-depth observational studies was carried out with selected temple murals, cloth paintings, wood and stone carvings, sculptures and ivory carvings at the Munich Treasury in Germany. The literature review employed original documents, manuscripts, chronicles, records of foreign travellers, published research and inscriptions. The validity of the data was confirmed by cross checking with literary sources.

Results and Discussion

The king was the culminating point of the pyramid in society and represented the state and the center of political life. The concept of king is distinguished by the term *Maharaja* (great king) or *Rajadhiraja* (ruler of the king) *Mahipathi* (lord of the earth) and *Deva* (god). The title corresponds to ‘Our Majesty’. Culture built certain stereotypes for the king as a living god that affected the cognition of people in society. The representation of the god in sculpture or in

* virajinik@uom.lk

paintings therefore could be hypothesized as the representation of a king. Because the artists of the period had no imagination of how such a god should be represented, they saw the live model of the king who was considered the lord of the earth. The king used 64 royal ornaments which signified royalty had to wear at his consecration and at important state occasions. He used fine white silk cloth for the lower dress and shawls for head covering. It was because of the culturally built social code regarding silk and fine cotton that people identified royalty or luxurious sophistication. Color became the signified purity according to tradition. The pictorial references given below show that the king dressed differently according to the day and the place where he represented himself.



Figure 1



Figure 2



Figure 3



Figure 4

Figure 5

Fig. 1 shows king Buwanekabahu V11 (1521-1555) at the royal platform *Chitakuta Mandape* at the palace of Kotte. He wore 64 royal ornaments, a long wrap cloth and is wearing the multi tiered crown, holding a lotus flower, one of the five emblems of the Lord Buddha. Fig. 2 shows the king receiving a visitor in the king's audience hall by wearing a head-dress instead of a crown with limited upper body jewelry. Fig. 3 shows the king participating in a ritual ceremony in front of the god Shiva. The king wore a different type - dress. Fig. 4 shows prince Dharmapala at the coronation ceremony in Portugal wearing a Sri Lankan king's dress exclusive to the emperor viz the headband, the single necklace of pearls called *ekavali* and the divine dagger called *acchijja – cchurika*. Fig. 5 shows prince Dharmapala is wearing a long coat over a long piece of cloth after he was crowned as a Portuguese king. His (*cabaya*+ long wrap cloth) new dress resembles and obeys the Portuguese crown.

Altered appearances found in new social contexts



Figure 6 Figure 7 Figure 8 Figure 9 Figure 10 Figure 11 Figure 12

Figure 13

It is noticed that most of the early kings of the Kandyan era were educated under Portuguese Franciscans and baptized by them. King Wimaladharmasooriya 1 (1591-1604) (Fig. 6) was known as Konappu Bandara who was grown and educated under Franciscans by the name of *Dom Joao de Austria* Mudaliyar. He embraced Buddhism and became the king of the Kandyan kingdom, as it was an important qualification for the kingship. The son of king Senarath (1604-1635) (Fig. 7) king Rajasinghe 11's (1635-1687) (Fig. 8) dress habits also show how he changed his mind with the impact of the environment he had grown up. Knox says, 'he was not wont to keep to one fashion, but changes as his fancy leads him'. King Wimaladhramasooriya 1 introduced the *Kameesa hettaya* or *Juan hettaya* with a *tippet* (collar) (Figs. 11, 12, 13) while prince Dharmapla first introduced the *cabaya*. King Wimaladhramasooriya 1 wore the *thuppotti* for the lower dress. King Rajasinghe 11 wore a long sleeved jacket with long breeches.

The study revealed that an article of clothing (*tippet*) has many possible meanings and so there is likely to be a certain degree of ambiguity. The articles of clothing that are worn together may have numerous possible meanings and they also interact with one another to produce additional ambiguities. It is also observed that the jacket of king Wimaladhramasooriya 1 had an attached collar known as the *tippet*. This dress article was also utilized by kings Keerthi Sri Rajasinghe and Sri Wrickrama Rajasinghe. However the size of the article is seen to be bigger than the *tippet* of Wimaladhramasooriya 1. It is known that these kings practiced oiling their hair as etiquette and they separately attached the *tippet* to the neck line of the jacket as a protective dress item which they supposed would not allow oil to run into their jacket. In this way the signification of *tippet* would be fashion attitudes, or daily habits of the royalty. However, king Wimaladhramasooriya 1's *tippet* being attached to the jacket would signify a different signification from Keerthi Sri Rajasinghe and Sri Wrickrama Rajasinghe's *tippet*. Furthermore, the study also revealed that there is a high degree of social variability in the link between signifier (dress) and signified (concept) when it comes to appearance. The royal dress with long sleeved coat (*cabaya*) and a lower piece of cloth of the Kotte period developed to a special costume called *thuppotti* during the Kandyan era. The *thuppotti* means the whole garment which was worn by the elite of Kandy. The *thuppotti* dress has now become a fashionable garment at wedding ceremonies. Many people love to wear *thuppotti* on their wedding day. Social order is established regarding dress of each category. Social codes were more stable than in the present days to interpret dress messages.

Conclusions and Recommendations

The study revealed that dresses or clothes are semantically unstable, because the meaning is directly related to the context, when clothes are considered as texts; it was typical in hierarchical societies for clothes to function as 'closed' texts with a relatively stable and fixed meaning. This is also proved by the dress of prince Dharmapala; king Wimaladhramasooriya 1 and king Rajasinghe 11 These dresses were embedded in the Portuguese influenced social, cultural and historical context of the sixteenth and seventeenth centuries. These four royal personages were brought up in the Portuguese social context from their childhood. The kings Keerthi Sri Rajasinghe and Sri Wrickrama Rajasinghe's dresses were also embedded in Dutch and British influenced social contexts. Within that western influenced social context all the seven dresses of the above said kings prevailed at a certain time period, remaining as fixed dresses of the royalty. These examples also show that dress signs rely heavily on social contexts in order for a message to be comprehended. Therefore, the applicability of fashion meanings of the Kotte period to the present as well as to the future is possible.

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PHENOLIC CONTENT AND ANTIOXIDANT ACTIVITIES OF LEGUMES AS AFFECTED BY DIFFERENT PROCESSING CONDITIONS

K.D.D. Kumari^{1*}, A.L.F. Sifna¹ and G.A.P. Chandrasekara¹

¹*Department of Applied Nutrition, Faculty of Livestock, Fisheries and Nutrition,
Wayamba University of Sri Lanka*

Introduction

Legumes are plants belonging to the family *Fabaceae*. They are edible seeds, borne in pods that often open along with two seams. Legumes have two types according to the way of storing energy. The leguminous oilseeds such as soya bean and groundnut store energy as lipids. The pulses such as peas, beans, and lentils store energy as starch. Legumes are rich in proteins, and a good source of essential amino acid lysine, but are deficient in sulphur containing amino acids such as, methionine and cysteine. Therefore legumes are valuable complementary food to cereals due to differences of individual amino acid compositions. Phenolic compounds such as phenolic acids, flavanoids, and condensed tannins and their antioxidant activities are identified and characterized in food legumes. Phenolic compounds can act as antioxidants via different mechanisms and provide beneficial health effects to humans. Processed legumes play a vital role in the diet of Sri Lankans. Therefore a study was conducted to determine the phenolic content and antioxidant activities of legumes commonly consumed in Sri Lanka as affected by different processing methods.

Materials and Methods

Six different types of commonly consumed legumes, namely *Phaseolus aureus* (Green gram; cultivar: Ari), *Phaseolus mungo* (Black gram; cultivar: Anurada), *Cicer arietinum* (Chickpea; Red, Yellow), *Vigna unguiculata* (Cowpea; cultivar: Varuni), and *Glycine max* (Soybean; cultivar: PB1) were used in this study. Samples were processed by soaking (12 hour, room temperature in water 1:5 (w/v)), peeling (manually), boiling (boiled in distilled water 1:1 (w/v) for 30 minutes) and germination. The raw and processed legume samples were freeze dried and defatted using hexane (1:5 w/v). Soluble phenolic compounds were extracted using 50 % acetone (v/v), shaking in a water bath, for 8 h at 10±5 °C. The extracts were tested for their total phenolic content (TPC) by folin-ciocalteu reagent, and total flavonoid content (TFC) by Aluminium (III). Antioxidant activities were measured by reducing power (RP) capability of donating electrons, ferrous ion chelating activity (FICA), and

* disnakumare@yahoo.com

trolox equivalent antioxidant capacity (TEAC) by ability to reduce the 2, 2'-azinobis (3-ethylbenzothiazoline-6-sulfonate) radical.

All experiments were carried out in triplicates and data are presented as mean \pm standard deviation. The significance of differences of mean values among each processing methods were determined using one-way analysis of variance (ANOVA). Statistical analyses were performed using SPSS version 16 software.

Results and Discussion

Table 1. TPC of legume samples with different processing methods

Sample	TPC μ moles of gallic acid equivalents per g of defatted meal (dm)				
	Raw	Soaked with peel	Soaked and peeled	Soaked, peeled and boiled	Germinated
Chick pea	4.1 \pm 0.2 ^a	4.3 \pm 0.2 ^a	3.5 \pm 0.1 ^a	0.8 \pm 0.0 ^a	2.8 \pm 0.0 ^a
Cow pea	27.6 \pm 0.5 ^b	20.6 \pm 0.2 ^b	6.3 \pm 0.1 ^b	5.8 \pm 0.1 ^b	7.1 \pm 0.5 ^b
Soya bean	9.7 \pm 0.1 ^c	8.5 \pm 0.4 ^c	10.4 \pm 0.0 ^c	9.7 \pm 0.5 ^c	5.1 \pm 0.1 ^c
Black gram	17.5 \pm 0.3 ^d	5.6 \pm 0.6 ^d	2.8 \pm 0.2 ^d	6.3 \pm 0.0 ^d	9.7 \pm 0.2 ^d
Green gram	14.6 \pm 0.1 ^e	13.5 \pm 0.1 ^e	7.4 \pm 0.2 ^b	3.4 \pm 0.2 ^e	13.5 \pm 0.7 ^e

Note: Values in each column with same letters are not significantly different ($p>0.05$)

The TPC content of raw legumes ranged from 2.25 to 27.55 μ moles of gallic acid equivalents per g of dm (Table 1). Soaking and boiling processes reduced the TPC of legumes, compared with raw counterparts.

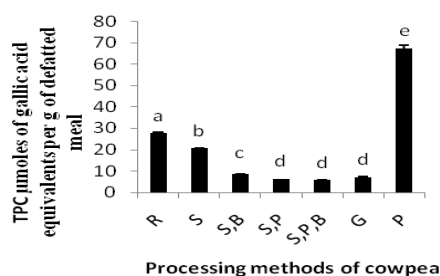


Figure 1. TPC of cowpea with different processing methods

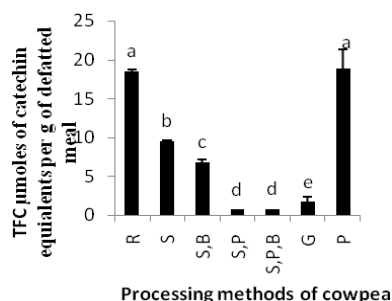


Figure 2. TFC of cowpea different processing methods

Note: (R-Raw cowpea with peel ; S- Soaked; SB-Soaked, Boiled; SP-Soaked, Peeled; SPB-Soaked, Peeled, Boiled; P-Peel)

Letters in columns of each category indicates the significantly different values $P < 0.05$

The Figs. 1 and 2 show the reduction of phenolic contents of cowpea, due to different processing methods. Peel contained the highest TPC and TFC of cowpea sample. Several studies showed that the outer part of the plant foods, peel or the outer cover of the seeds contain more bioactive compounds, especially to protect the plant foods from pest attacks. The dark colour seed coats contain high antioxidant properties due to their high content of polyphenolic compounds.

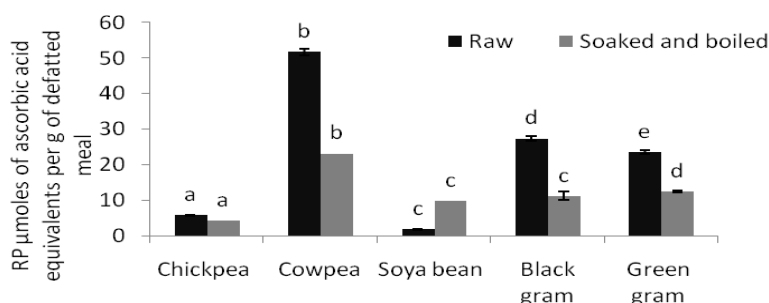


Figure 3. RP of raw samples, and soaked, boiled legume samples

Note: letters in columns of each category indicates the significantly different values $P < 0.05$

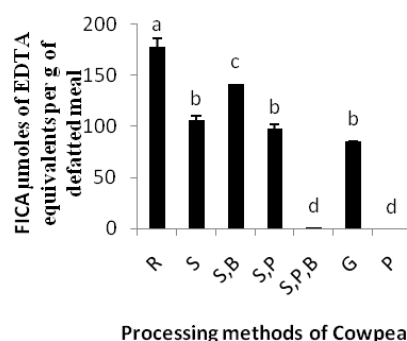


Figure 4.
FICA of cowpea with different processing methods

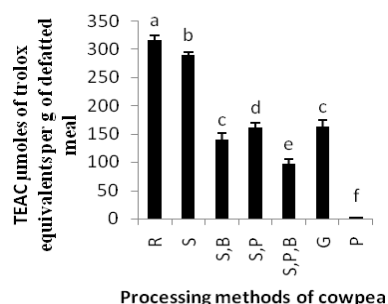


Figure 5. TEAC with different processing methods

Note: letters in columns of each category indicates the significantly different values $P < 0.05$

Processing methods changed the antioxidant activities of raw legumes (Figs. 3, 4 and 5). The reduction of TPC content due to soaking and boiling is reason of having the low RP values of the legume samples. Soaking will softens the cell walls of legumes and soluble phenolics can leach into the soaked water. According to the Mohamed et al. (2011), soaking of legume seeds reduced phenolic contents and more reduction observed with the increasing of period of soaking.

Thermal processes have a large influence in phenolics availability in foods which depends on their magnitude and duration. Xu and Chang (2008) reported that the soaking, boiling and steaming significantly reduced the phenolic contents and antioxidant activities of cool season food legumes including green pea, yellow pea, chick pea and lentil.

Conclusions and Recommendations

Legumes are rich sources of phenolic compounds and they possess antioxidant activities. Antioxidant activities are significantly different among the legumes. Cowpea contained more phenolic compounds per g of defatted meal among other legumes examined. The peel of the legumes contains the highest content of phenolic compounds except soya bean. The processing changes the phenolic content and their antioxidant properties of legumes. Legumes are potential source of antioxidants which can be utilized as functional foods in the management of non communicable diseases. Further studies are warranted to identify best food processing methods to retain more phenolic contents and their antioxidant activities of legumes.

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THERMAL CHARACTERIZATION OF GRAPHITE, GRAPHITE-OXIDE (GO) AND GRAPHENE (GN) BY THERMOGRAVIMETRIC ANALYSIS (TGA/DTA), AND OPTIMIZATION OF TEMPERATURE OF GRAPHITE-OXIDE TO GRAPHENE BY HTA/XRD

C.H.Manoratne^{1*}, I.R.M.Kottegoda¹ and S.R.D.Rosa²

¹*Materials Technology Section, Industrial Technology Institute, Colombo 7, Sri Lanka*

²*Department of Physics, Faculty of Science, University of Colombo, Sri Lanka*

Introduction

Exports of local minerals are discouraged without a value addition in the present strategies of government policies. Adding value to the local minerals is advised to increase the foreign revenue; as a result graphite is identified as a potential mineral that can be value added. Graphene (Gn) and graphite-oxide (GO) are such value added products which have become wonder materials in the present research and development in every part of the world. Since the separation of single-layer graphene (SLG) in 2004 has made scientists to explore the suitable applications of Gn as it shows two dimensionality and unique physical properties (Janowska et al. 2010). The intermediate product of GO has also been attracted by the scientists recently as a potential carbon-based nanomaterial for producing graphene (Su et al. 2009). Chemical composition (purity), particle size, surface area, surface morphology, and the thermal behavior etc. are some of important properties need to be analyzed as they highly govern the final properties in any application (Wissler 2006). To name few, the powder size influences the ultimate properties of graphite oxide and graphene (Jiang et al. 2000). Therefore, in this partial study an attempt was devoted to study the relative fraction of edge sites and the oxidation behavior of graphite and its derivatives and to optimize the conversion temperature of GO to Gn.

Materials and Methods

Commercially available local graphite powder as received with the average particle size (d₅₀) ~ 110 µm was reduced in particle size ((d₅₀) ~ 37 µm) by using ball-milling. A planetary ball mill was used by maintaining the ball-to-powder ratio of 5:4. Total pebbles volume is taken as 50 % (V) of the mill volume. Pebbles in three sizes, i.e. large in 20 % by V volume, medium size pebbles in 40 % by V, and small size pebbles in 40 % by V. Water is added sufficiently enough to wet the powder about 8 % by V. The effect of ball milling time was optimized by collecting milled powders at an interval of 1 h up to 24 of milling and measuring particle size distribution.

* manoratne@yahoo.com

The resulting powder was dried and used for synthesizing of graphite oxide. Modified hummer method (Kottegoda et al. 2001) was adopted for the synthesis of GO and Gn. Materials were characterized by using SDT Q600 Simultaneous Thermogravimetric Analyzer under N₂ flow. The weight change and true differential heat flow on the same sample from ambient to 1500 °C was measured. The reduction or the conversion of graphite-oxide to graphene was systematically studied with the High Temperature Attachment (HTA) attached to the XRD (Rigaku Ultima IV/HTA). The morphology of the products was studied by the SEM (LEO 1420VP).

Results and Discussion

Thermal characterization of graphite, graphite-oxide and graphene:

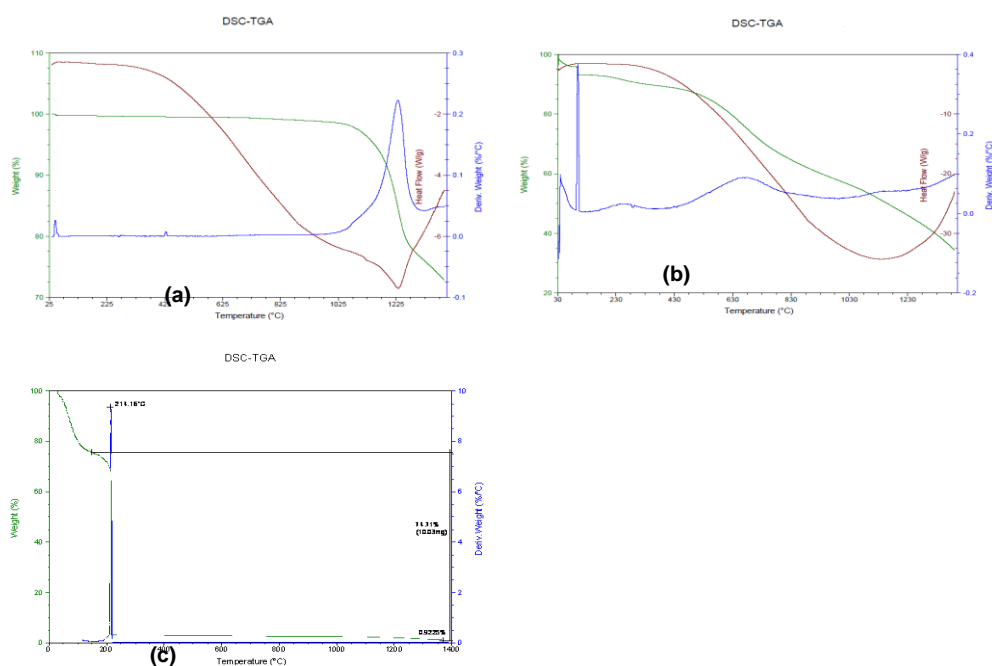


Figure 1. TGA/DSC plots for (a) Graphite, (b) GO, and (c) Gn

The Fig. 1 shows the simultaneous measurement of weight change and true differential heat flow (TGA/DTA) on the sample of graphite (a), graphite-oxide (b) and graphene (Gn) from ambient to 1500 °C. As seen in the Fig. 1(a) the graphite was highly stable up to 1000 °C, and revealed pyrolysis of oxygen-containing functional groups available in the graphite layers as the derivative dW/dT shows three peaks as a function of temperature appearing at around 50 °C, 430 °C and 1230 °C, respectively. The GO shows slight mass decrease from room temperature to 150 °C and significant decrease from 150 °C to 200

°C, and slowly decreased up to 600 °C (Fig. 1b). Fig. 1(c), the corresponding thermal characteristics of Gn on the other hand showed an enhanced thermal stability up to ≈ 214 °C.

Optimization of conversion temperature of GO to Gn by HTA/XRD:

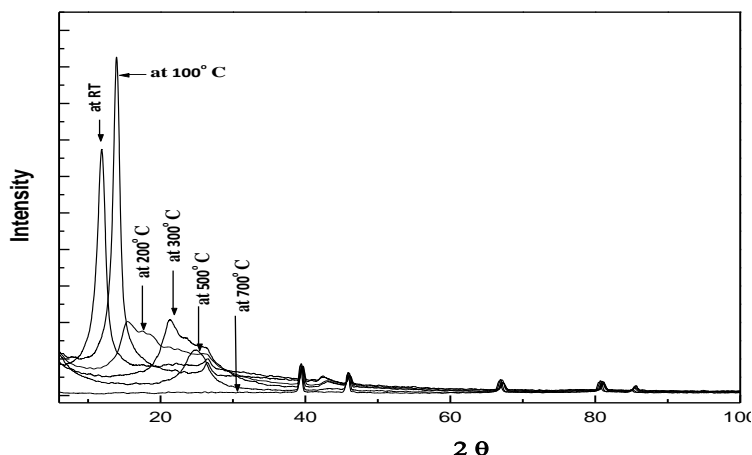


Figure 2. XRD spectra at different temperature of 100 °C, 200 °C, 300 °C, 500 °C, 700 °C

The GO was subjected to different temperatures of 100 °C, 200 °C, 300 °C, 500 °C and 700 °C. It was observed that there is an optimum low temperature range (300 °C- 500 °C) in which the GO can be converted to Gn (few layers) as confirmed with the change of characteristic peak of GO at $2\theta \sim 11^\circ$ to Gn at $2\theta \sim 24-26^\circ$ (Fig. 2). The BET surface area measures is about 290-308 $\text{m}^2 \text{g}^{-1}$ which depends on the graphene layers.

Microstructure Analysis:

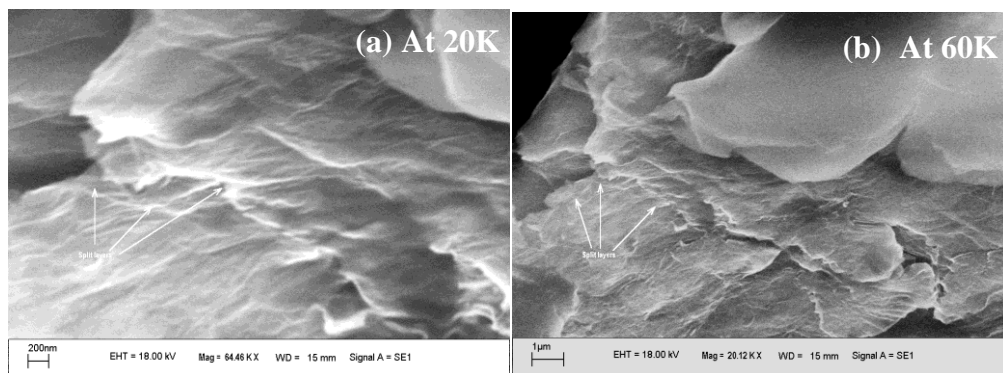


Figure 3. SEM images of Gn at the magnification of (a) x 20, 000, and (b) x 60, 000

The morphology of Gn was observed at the magnifications of (a) x 20,000 and (b) x 60,000. As marked by the arrows it can be observed that the graphite structure has been split during the synthesis process (oxidation/reduction) and the few layers of graphene are conspicuous.

Conclusions and Recommendations

Thermogravimetric analysis reveals pyrolysis of oxygen-containing functional groups available in the graphite layers, GO, and to some extent in Gn layers. The XRD spectra obtained at different temperature also confirmed this phenomenon. The temperature (range) for conversion of GO to Gn was optimized. The findings will be effectively used in optimum practical synthesis of GO, Gn, and graphene composites for the applications in Li-ion batteries in the future.

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IN VITRO ANTIBACTERIAL ACTIVITY OF CATECHINS FROM REFUSE GREEN TEA AGAINST DRUG-RESISTANT *Pseudomonas aeruginosa*

S.S. Mediwake^{1,2}, I.S.K. Vithanage^{1,2}, V. Thevanesam², E.W.M.A. Ekanayake²
and B.M.R. Bandara^{1*}

¹Department of Chemistry, Faculty of Science, University of Peradeniya

²Department of Microbiology, Faculty of Medicine, University of Peradeniya

Introduction

Pseudomonas aeruginosa is a Gram-negative bacterium causing many nosocomial infections such as urinary tract infections, pneumonia and bacteremia. Emergence of multi-drug resistance in *P. aeruginosa* makes treatment of these infections difficult. New strategies to combat drug-resistance are urgently required. However, new antibiotics are hardly introduced by pharmaceutical companies due to the high cost of producing new drugs and the eventual development of resistance to new antibiotics in organisms (Okeke et al. 2005).

Some natural products show antimicrobial properties. Catechins extracted from *Camelia sinensis* L. (tea) have displayed antibacterial activity against Gram-positive organisms like *Staphylococcus aureus* and Gram-negatives like *Escherichia coli* (Taguri et al. 2004). Recent studies in our laboratory reveal that tea catechins have moderate antibacterial activity against drug-sensitive *P. aeruginosa* (Karunathilake et al. 2007). The objective in the present study is to isolate catechins from refuse green tea and to evaluate the crude catechin fraction (CCF) for antibacterial activity against drug-resistant *P. aeruginosa*. Over 10-12 % from the total tea harvest is removed as refuse tea - the tea left-over from veins and stems of the tea leaf – during manufacture. New uses of refuse tea will enhance its value and help boost the tea industry in Sri Lanka.

Materials and Methods

Refuse green tea samples were obtained from a local factory and CCF was extracted with 70 % aqueous methanol followed by washing with dichloromethane and then with ethyl acetate. Ethyl acetate fraction was concentrated and freeze dried to obtain CCF as a brown colored solid.

Thirty *P. aeruginosa* strains were collected from Teaching Hospital Peradeniya. The identification of *P. aeruginosa* strains was done using Gram stain, motility, oxidase, catalase, DNase and oxidative-fermentative tests. The antibiotic sensitivity was determined using agar disc diffusion method (CLSI method). The Minimum inhibitory concentration (MIC) of ciprofloxacin

* bmrbandara@gmail.com

against each strain was determined using agar dilution method. *P. aeruginosa* NCTC 10662, ATCC 27853 were used as the control strains. The antibacterial activity screening of 5 mg mL⁻¹ CCF against all test strains was screened using agar well diffusion assay while MIC of CCF against *P. aeruginosa* was determined using agar dilution method on Mueller Hinton Agar (MHA).

Results and Discussion

Twenty three *P. aeruginosa* strains that had MIC values of ciprofloxacin ranging 8×10^{-3} - $>0.128 \times 10^{-3}$ mg mL⁻¹ were considered as drug-resistant strains. The CCF extracted from refuse tea inhibited six strains of drug-resistant *P. aeruginosa* in well diffusion assay, and the diameter of the inhibition zones ranged 18 - 21 mm (Fig. 1). The MIC value of the CCF corresponding to each of the 30 drug-resistant *P. aeruginosa* strains was determined using the agar dilution assay (Andrews 2001), and the results are given in Table 1. MIC values of CCF were in the range 0.3 - >1 mg mL⁻¹.

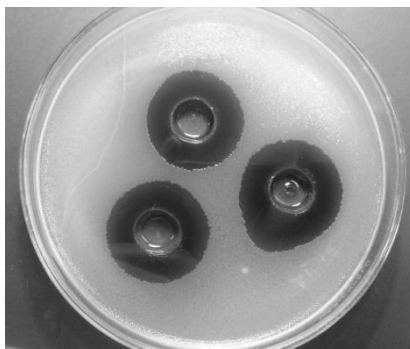


Figure 1. Zones of inhibition around wells in well diffusion assay for catechins against *P. aeruginosa* (well diameter, 12 mm).

CCF had higher MIC values when compared with ciprofloxacin. Irrespective of the sensitivity of ciprofloxacin for some strains, catechins displayed higher MIC values. In a previous study, the aqueous extract of green tea has exhibited MIC 0.8 mg mL⁻¹ against drug-resistant *P. aeruginosa* (Radji et al. 2013) while the CCF extracted from fresh tea flush has an MIC value of 0.5 mg mL⁻¹ for eleven *P. aeruginosa* strains (Karunathilake et al. 2007). Our results together with those from the previous studies suggest that refuse tea catechins are as potent as those available from green tea and fresh tea flush.

Table 1. MIC values of ciprofloxacin and CCF against control, drug-resistant and drug-sensitive strains of *P. aeruginosa* determined using agar dilution assay

<i>P. aeruginosa</i> (PA) strain	MIC ($\times 10^{-3}$ mg mL ⁻¹) of Ciprofloxacin	MIC (mg mL ⁻¹) of CCF
PA NCTC 10662	0.25	0.6
PA ATCC 27853	0.25	>1
PA 1	0.25	0.7
PA 2	<0.12	>1
PA 3*	64	>1
PA 4*	128	0.3
PA 5*	64	>1
PA 6	<0.12	>1
PA 7	<0.12	>1
PA 8*	32	>1
PA 9*	32	>1
PA 10*	>128	>1
PA 11*	>128	0.3
PA 12*	>128	0.3
PA 13*	32	>1
PA 14	0.5	>1
PA 15*	16	>1
PA 16*	64	0.3
PA 17*	32	0.6
PA 18*	64	>1
PA 19*	32	>1
PA 20*	8	1
PA 21	2	0.3
PA 22*	>128	>1
PA 23	0.25	>1
PA 24*	>128	>1
PA 25*	64	0.6
PA 26*	32	0.6
PA 27*	64	0.3
PA 28*	8	>1
PA 29*	>128	>1
PA 30*	64	>1

Note: PA*- Ciprofloxacin resistant *P. aeruginosa* strains

Conclusions and Recommendations

CCF isolated from refuse green tea displayed moderate antibacterial activity against some of the strains of drug-resistant *P. aeruginosa*. Further studies on the CCF, to enhance its potency, and its mechanism of action on the bacterial strains are recommended.

Acknowledgement

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IN-VITRO AND FIELD EFFICACY OF KASUGAMYCIN ON *Xanthomonas campestris* pv. *campestris*, THE BLACK ROT PATHOGEN IN CABBAGE

L.D.C. Mendis^{1*}, K.P. Somachandra² and M.L.M.C. Dissanayake¹

¹*Faculty of Agricultural Sciences, Sabaragamuwa University of Sri Lanka*

²*Regional Agricultural Research and Development Centre, Bandarawela*

Introduction

Cabbage black rot (*Xanthomonas campestris* pv. *campestris* (Xcc)) is considered as a very destructive disease of Crucifers worldwide (Sikora 2004). Bacterial diseases in plants are usually controlled by cultural practices and application of antibiotics, up to certain extent. However several limitations and constraints are associated with such practices. Even though copper is not recommended by Department of Agriculture (DOA) as it is a heavy metal, some farmers apply copper-based fungicides to check the disease.

Kasugamycin is an antibiotic which belong to active toxicity profile of category IV and it only used in agriculture; there are no known medical or veterinary uses (EAP 2005). It possesses weak or almost no antibacterial activity against common pathogenic bacteria in human or animals (Tamamura et al. 1998). Therefore, use of Kasugamycin in agriculture may not create an adverse effect on environment or human health. Although with all these black rot control methods, still the disease causes heavy losses in cabbage cultivation in the upcountry. Therefore, there is a pressing need to find new feasible control method for black rot disease. Recent studies carried out at RARDC, Bandarawela revealed that Kasugamycin have inhibitory effect on Xcc. Thus, this study was undertaken to investigate the applicability of using Kasugamycin 2 % (Kasumin®) as a feasible control method for black rot disease in cabbage (*Brassica oleracea* var. *capitata*) variety “Green Coronet”.

Materials and Methods

In-vitro experiment was conducted to identify the suitable concentration of Kasugamycin to suppress the growth of Xcc. The selected concentrations were tested in open field under natural disease pressure. Infected leaves with black rot symptoms were collected from field and confirmed the presence of bacteria by oozing out of the bacteria cells under the light microscope. Hundred µl of bacterial suspension was poured onto nutrient agar (NA) plates. Plates were kept in incubator in invert position at 28 °C temperature. Bacterial colonies developed in NA plates after 72 hours were picked up and streaked in yeast dextrose chalk (YDC) for purification; a pathogenicity test was carried out for verification.

* dinu.chamari@gmail.com

Three concentrations (100, 200 and 300 ppm) of Kasugamycin solutions were prepared using the commercial formulation 2 % Kasumin^R. They were tested against Xcc using diffuse disk test (Kirby and Bauer 1961). Copper fungicide (1000 and 2000 ppm) and sterilized distilled water were used as an alternative chemical control and negative control respectively. Diameter of the zone of inhibition was calculated using a metric ruler after 1-2 days after inoculation and data were statistically analyzed to see the differences between treatments.

100 ppm and 200 ppm of Kasugamycin and 2000 ppm copper (DOA recommended concentration) were used for field study. Chemicals were applied as foliar sprays at the time of symptom appearance and continued weekly; four sprays were done during the three months cropping period. Black rot disease severity was calculated using randomly selected eight plants in each plot using 1-4 scale described by Williams et al. (1972) while number of infected leaves plant⁻¹ was counted at the end of the trial and recorded as disease incidence. Data were analyzed using analysis of variance (ANOVA) by SAS and treatment means were separated using the Tukey's Studentized Range (HSD) test at P=0.05.

Results and Discussion

Isolation and identification of Xanthomonas campestris pv. Campestris:

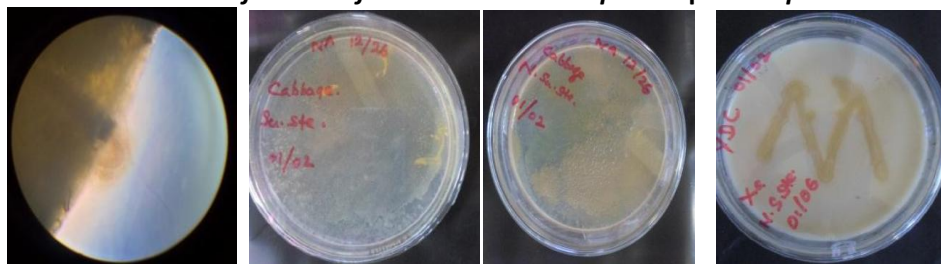


Plate 3.1.1:
Bacterial ooze
under 100
magnifications

Plate 3.1.2: Cultured Xcc suspension
on Nutrient Agar media

Plate 3.1.3 Xcc streak
on Yeast Dextrose
Chalk media

Presences of bacteria were confirmed by oozing out of bacterial ooze under 100 × magnification of light microscope (Plate 3.1.1). Bacteria were identified by light gold color colonies on NA medium (Plate 3.1.2) and yellow color culture characteristic on Yeast Dextrose Chalk medium (Plate 3.1.3). Black rot symptoms were observed on leaves of cabbage plants injected with bacterial suspension after 10 to 14 days and no symptoms were developed in leaves inoculated with sterilized distilled water.

In vitro test: According to results, Kasugamycin at 200 and 300 ppm concentrations showed the significant inhibition (1.5 cm) of bacterial growth and Copper fungicide showed minimum inhibition zone 0.6 cm at both concentrations used (Fig. 3.2.1).

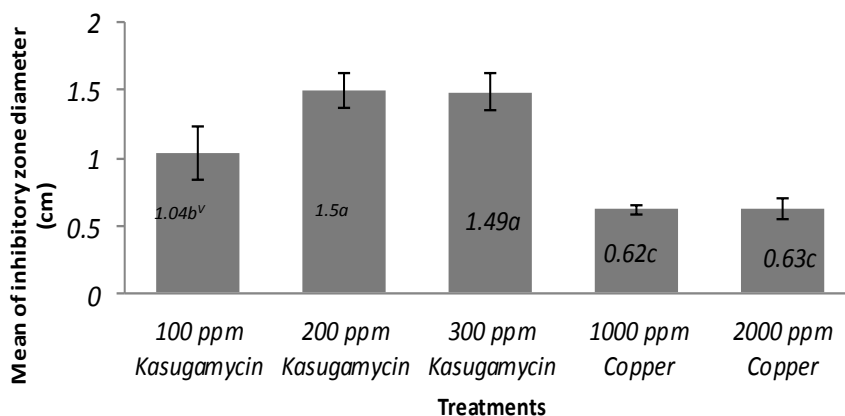


Figure 3.2.1: Mean of inhibition zone diameters of Xcc growth under in-vitro condition

Table 1. Treatments of the field experiment

Treatments	Disease incidence	Disease severity*
	Number of infected leaves	Means of Black rot disease rating
100 ppm Kasugamycin	8.31 ^b	3.09 ^a
200 ppm Kasugamycin	8.28 ^b	2.24 ^b
2000 ppm Copper	9.68 ^a	2.71 ^{ab}
Control	8.74 ^{ab}	2.42 ^{ab}
CV %	5.06	6.47

Note: Means followed by the same letter are not significantly different according to Tukey's Studentized Range (HSD) test at $P \leq 0.05$.

**Based on 1-4 scale of Williams (1972) where 1 represent lowest severity and 4 represent highest*

Kasugamycin in the rate of 100 ppm and 200 ppm showed significant ($p=0.05$) reduction of disease incidence (8.28) with compare to copper fungicide (Table 3.3.1). However high concentration of Kasugamycin 200 ppm showed lowest diseases severity with compared to lower concentration of Kasugamycin and copper. It revealed 200 ppm of Kasugamycin is best rate to reduced disease severity and disease Incidences. Though, 100 ppm Kasugamycin showed remarkable suppression of Xcc in in-vitro tests, the effect was not observed in the field. Cabbage plants treated with 100 ppm Kasugamycin showed severe black rot symptoms. It may be due to the lack of retention of the chemical in plant tissues or the dosage is insufficient under field conditions. Higher

Kasugamycin dose reduced disease incidence however, both doses tested were not suppressed the disease severity. Results revealed that though Kasugamycin showed superior performances in *in-vitro* tests, but it is less effective to reduced disease severity in field, once cabbage cultivation is infected with pathogen. It may due to effect of various factors at natural environmental conditions which influence the chemical management of plant diseases in the field.

Jones and Jones (1985) explained that chemical control methods were relatively ineffective when disease pressure is high. Therefore this phenomenon cannot be explained with the data generated from present study. Further studies were in need to give a firm conclusion on the behavior of Kasugamycin in field level under Sri Lankan context.

Conclusions and Recommendations

Application of Kasugamycin (200 ppm) is suitable to reduce black rot incidence of cabbage in field level. The conclusion is drawn with the *Yala* season a result, therefore further studies in *Maha* season is in need to go for a recommendation.

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EFFECT OF WATER QUALITY PARAMETERS ON BENTHIC MACROINVERTEBRATES

D.S.N. Munasinghe^{1*} and M.M.M. Najim¹

¹*Faculty of Science, University of Kelaniya, Kelaniya, Sri Lanka*

Introduction

Small hydropower projects offer one of the most promising energy resources for long term sustainable development. Small hydropower can, however, exert multiple impacts on local environment, and the impacts which are perceived to be of critical importance are ecological, centered on aquatic flora and fauna. The underlying cause could be attributed to the non-maintenance of sufficient environmental flow downstream, below the weir. The alteration of flow regimes is often claimed to be the most serious and continuing threat to the ecological sustainability of rivers because the duration and seasonal timing of associated low flow conditions can strongly influence organisms directly and via changes to habitat.

Benthic macroinvertebrates are an important component of the river biota and are characteristic as stream health indicators. There is a wealth of literature suggesting that macroinvertebrate community composition is tightly linked to in stream hydraulic conditions (i.e flow variation) inadvertently suggesting that the major factor governing benthic macroinvertebrate composition is in stream flow conditions. This research focused on presenting a comparative account of the water quality and the benthic macroinvertebrate species composition and diversity upstream and downstream of the Stream impoundment caused by Mini Hydro Diversion in the Hungampola South/Morontota village section of the Gurugoda Oya.

Materials and Methods

In the present study six sampling sites (sites A-F) were rationally established in the Hungampola South and Morontota village section in the Gurugoda Oya to capture the effects of different flow regimes on benthic macroinvertebrates. The study was carried out from May to November 2013 with monthly intervals between sampling occasions.

At one given sampling site, a transect line perpendicular to the stream was divided into three equal parts and sampling was carried out in the center of these three sections separately. Macroinvertebrates were collected using a standard D-framed dip net consisting of a D-shaped metal frame (0.3 m width and 0.3 m height) holding a conical net (mesh aperture 400 μ m). The collected

* dinukemunasinghe@kln.ac.lk

samples were preserved in a solution of Rose Bengal containing 5 % formaldehyde. Later in the laboratory, the material retained was wet sieved through a mesh (0.5 mm aperture) and identified to the nearest possible taxonomic category using standard identification keys. After initial identification, macroinvertebrates were enumerated in their different taxonomic groups.

At each sampling site, each time a biological sample was taken, the physico-chemical parameters of the overlying water immediately above the bottom were measured using standard sampling and analytical procedures. The parameters thus measured were temperature, pH, conductivity, water flow velocity, Total Dissolved Solids (TDS), Dissolved Oxygen (DO) and five day Biological Oxygen Demand (BOD₅).

Results and Discussion

During the sampling period (May - November 2013), sixteen (16) different macrobenthic invertebrate taxa belonging to three phyla, viz; Annelida, Mollusca and Arthropoda, were found in the six study sites. In these sites, the number of taxa recorded upstream, above the weir (Sites C, D, E), was higher than that of the downstream sites, below the weir (Sites A, B, C). The total species richness of all the study sites ranged from 7 to 13 during the sampling period and further, the species richness in these six sites increased during months categorized by high rainfall (June and October). *Pila* sp. dominated all the six study sites where its mean percentage contribution to the total macrobenthic data matrix was 74.49 %.

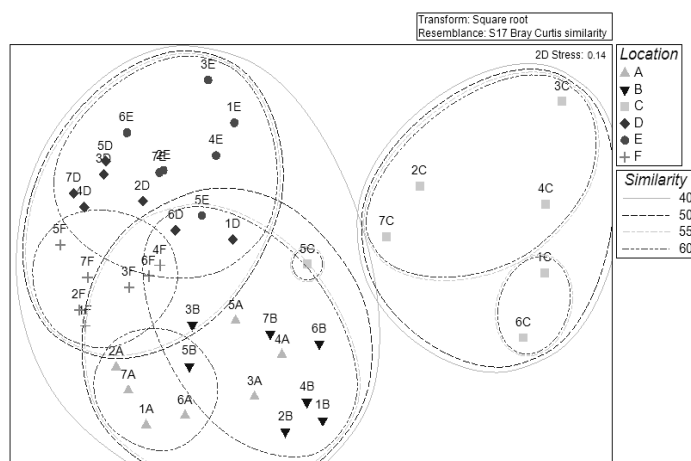


Figure 1. Non-metric Multidimensional Scaling (MDS) plot depicting the temporal and spatial variation of the macrobenthic community assemblages during the study period from May- November 2013.

The Non – Metric Multidimensional Scaling (NMDS) plot created for the macrobenthic invertebrate community structure in the present study (Fig. 1) explicitly shows how different Site C is in macrobenthic community composition compared to the other sites and also the high level of intra-site variation during sampling events during the study period.

A high cumulative percentage of 71.6 %, of the total variation of the physico-chemical parameters could be explained by both PC₁ and PC₂ axis of the Principal Component Analysis (PCA) (Fig. 2). The radiating axes from the center of the physico-chemical parameter matrix incorporated into the PCA plot shows that relatively higher levels of DO and temperature influence the macrobenthic community structure at site C in comparison to sites A and B and also that in comparison to site E, site F is more dominated by the depth of the water column, flow velocity and volume flow rate (Fig. 2).

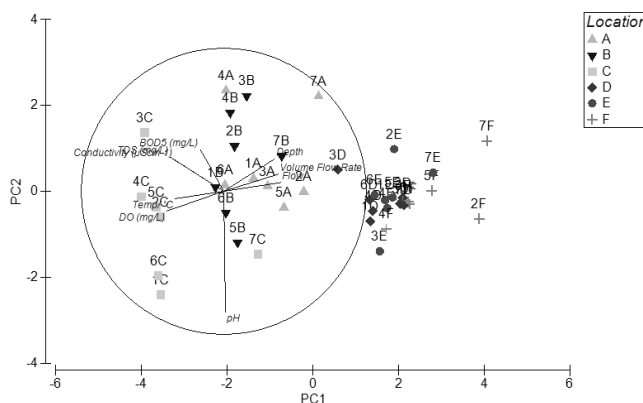


Figure 2. Ordination of the sampling occasions based on PC₁ and PC₂ scores of Principal Component Analysis (PCA) between physico-chemical parameters among the six sites. The radiating axes from the center of the physico-chemical parameter matrix are incorporated into the PCA plot.

The results of the present study demonstrate a clear relationship between flow regime changes and benthic macroinvertebrate community structure up - and downstream of the weir. The species richness of sites D, E and F, relatively pristine areas with the normal diversity of environmental conditions, were comparatively higher than the regulated sites A, B and C throughout the sampling period from May – November 2013.

Ordination of macroinvertebrate communities based on average abundance for each sequence demonstrated three distinct clusters (Fig. 2) at around 50 %

similarity in the macrobenthic invertebrate community structure over the period of record (May – November 2013) (Fig. 2; NMDS 2D stress 0.14). ANOSIM confirmed that assemblage composition differed statistically between these three clusters ($P < 0.05$, Global R 0.751). The most noticeable fact was that site C clustered separately and had the most intra-site variations of taxa between sampling occasions. It was interesting to note that Site C was mostly dominated by nymph varieties throughout the sampling period. Varieties representing orders Ephemeroptera, Plecoptera and Tricoptera (EPT) (For study purposes, the presence of mayfly nymph, stonefly larva and caddisfly larva are taken as the presence of EPT taxa and as EPT taxa richness) were prominent. Rader and Belish (1999) suggested that an increase in water temperature in the dewatered zone in one of their study streams created more favourable conditions for Mayflies, in turn causing increased densities. Also the presence of in-stream rock substrata increases mayfly abundances. High DO levels recorded at site C could also be an important factor in EPT taxa abundance, especially Order Plecoptera which has a high requirement for oxygen. Analysis of physico-chemical properties at this location also show that extreme values were recorded here.

Conclusions and Recommendations

The study revealed how the species richness and species diversity of the macrobenthic fauna up and downstream of the Stream impoundment caused due to Mini Hydro diversion, as well as the prevailing physico - chemical parameters among the study sites in the Hungampola South/Morontota village section of the Gurugoda Oya varied significantly. It can be also inferred that the flow regime changes play an important role in structuring the benthic macroinvertebrate communities in the region.

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DEGREE OF MATERNAL TRANSFER OF ANTIBODIES AGAINST MEASLES, MUMPS AND RUBELLA TO NEWBORNS OF PRE- AND POST-VACCINATION ERA MOTHERS

**M. Nadhikala^{1,2}, P.P.S.L Pathirana^{1*}, S. Peiris⁴, S.M. Handunnetti³
and G.C.S. Galagoda²**

¹*Department of Parasitology, Faculty of Medicine, University of Colombo*

²*Department of Virology, Medical Research Institute, Sri Lanka*

³*Institute of Biochemistry, Molecular Biology and Biotechnology, University of Colombo*

⁴*Epidemiology Unit, Ministry of Health, Sri Lanka*

Introduction

Primary protection against measles, mumps and rubella at birth is provided mainly by maternal antibodies and it gradually decreases with time. These antibodies could hamper the humoral antibody response of infant to vaccination. According to Kacica (1995), the antibody levels were found to be less in mothers born after the beginning of vaccination era when compared with women born before the vaccination era who had natural measles. In Sri Lanka measles vaccination era can be considered as after 1987 since coverage reached 60 % of the population in the national immunization programme that year. The mothers who are below 27 years of age have had measles vaccination and thus could be expected to have anti-measles antibodies which could protect their infants during early infancy. Mothers, who are older than 27 years who were born in pre vaccination era, may not be immune to measles unless they had a natural infection.

The prevalence of maternal antibody is based on 3 principal determinants: starting level of maternal antibody, placental transfer of maternal antibody and the rate of decay of maternal antibody after birth. The starting level of antibody depends on the mothers' vaccination and infection history. Gestational age defines placental transfer therefore preterm neonates receives significantly fewer antibodies (i.e., longer gestation results in a higher ratio). Nicoara (1999) states active placental transfer of Immunoglobulin G (IgG) begins during the third trimester and increases sharply thereafter. At the end of gestation, IgG concentrations in fetal serum exceed maternal levels. Leuridan (2010) suggests HIV- type 1 infection and placental malaria negatively affect placental transfer resulting in less antibodies in infants. The rate of decay is related to maternal age and maternal antibody titer. Early antibody loss in infants is also related to lower maternal age. Decay of maternal antibodies has led to the necessity of

* sisira_pathirana@yahoo.com

vaccination to develop active immunity. Data on antibody prevalence in infants or their mothers was lacking in Sri Lanka to decide the proper timing of the measles, mumps and rubella combined (MMR) vaccination for it to be effective to achieve maximum benefit from vaccination. Therefore studying the level of maternal antibody and the degree of maternal transfer will help for vaccination programme. The objectives of the study were to compare antibody titres in pre- and post-vaccination era mothers (born before and after the year 1987, respectively) and to determine the degree of maternal antibody transfer to their newborns.

Materials and Methods

This was a cross sectional descriptive study. Two age groups of mothers; below and above 27 years of age who are in the last trimester of pregnancy, were recruited to the study from De Soysa Hospital for Women, Colombo. Blood samples were collected from 100 mothers (venous blood) and their newborns (using cord blood) after getting their written consent. Data on previous history of measles, mumps and rubella infections, vaccination history of the mother, gestational age of baby and weight at delivery of baby were collected. Pre-term and low birth weight babies were excluded. Mothers with any known immunological impairments and who were undergoing administration of blood products or immunoglobulins were also excluded.

Mothers and their newborns were categorized into two groups based on the age of the mother; pre- and post-vaccination era mothers and their newborns. There were 57 mothers of pre-vaccination era and 43 mothers of post-vaccination era. 65 % of them had history of prior infections of measles, mumps or both. Ninety eight of the mothers had been vaccinated against rubella and 44 against measles. Antibody levels were measured using commercial ELISA kits (Serion Immundiagnostica GmbH) following manufacturers' instructions. The cut-off level of sero-positivity for measles, mumps and rubella were considered as 200 mIU mL⁻¹, 100 U mL⁻¹ and 20 IU mL⁻¹, respectively. The data was analyzed using SPSS software. Mean antibody levels were compared using Student's *t*-test. The ethical clearance for this study was obtained from Ethics review committee of Medical Research Institute, Colombo.

Results and Discussion

ELISA results showed 91 % and 93 % of pre- and post- vaccination era mothers respectively were sero-positive for measles. Similarly, 98 % and 95 % of newborns delivered by pre- and post- vaccination era mothers respectively were sero-positive for measles. The mean IgG concentrations of measles antibodies in newborns exceeded that of maternal levels in a ratio of 2.2:1 in

pre-vaccination era mother-infant pairs and in 1.88:1 ratio in post-vaccination era mother-infant pairs. The level of measles antibodies in newborns was significantly correlated with that of their mothers ($r=0.818$, $p<0.001$ in pre-vaccination era group; $r=0.827$, $p<0.001$ in post-vaccination era group). Pre-vaccination era mothers and their newborns were seropositive for mumps (86 % and 89 %) whereas 93 % and 88 % of sero-positivity was seen in post-vaccination era mothers and their newborns, respectively. All the mothers who had anti-mumps antibodies had transferred antibodies to their newborn. The ratios of IgG concentrations of pre- and post-vaccination era mother-infant pairs were 0.67:1 and 0.74:1, respectively. The level of mumps antibodies in newborns was significantly correlated with that of their mothers ($r=0.751$, $p<0.001$ in pre-vaccination era group; $r=0.841$, $p<0.001$ in post-vaccination era group).

The prevalence of sero-positivity for rubella in pre-vaccination era mother-infant pairs were 94 % and 100 %, respectively and in post-vaccination era it was 95 % and 97 %, respectively. A significant positive correlation was observed between the level of rubella antibodies in mothers and their newborns ($r=0.764$, $p<0.001$ in pre-vaccination era group; $r=0.865$, $p<0.001$ in post-vaccination era group). The IgG concentration of newborns was higher than that of their mothers in both groups (1.26:1 in pre-vaccination era mother-infant pairs and 1.34:1 in post-vaccination era mother-infant pairs).

The results showed that pre-vaccination era mothers had significantly higher levels of antibodies compared to post-vaccination era mothers because they are most likely to have acquired immunity through infection whereas post-vaccination era mothers had less chance for natural boosting due to high vaccination coverage. Although the pre-vaccination era mothers were older than the post-vaccination era mothers, they had significantly higher levels of antibodies against all three diseases, compared to post-vaccination era mothers. This may imply that immunity acquired through infection is long lasting than immunity acquired through vaccination. The degree of maternal antibody transfer to the newborn infants appears to depend on the antibody levels of the mother. Thus the newborns of pre-vaccination era mothers had higher levels of antibodies transferred to them than the newborns of post-vaccination era mothers. An efficient transfer of maternal antibodies has occurred and a majority of the infants were sero-positive (97 % for measles, 89 % for mumps and 99 % for rubella).

Conclusions and Recommendations

Mothers who were born in the pre- vaccination era and their newborns had higher levels of IgG compared to post-vaccination era mothers and their newborns. There is a high degree of maternal antibody transfer to the

newborns against all three diseases giving rise to protective levels of maternal IgG levels in newborns. It is recommended to expand the study consisting larger sample size with assessing functional antibody levels.

Acknowledgement

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IDENTIFICATION OF DIFFERENTIALLY EXPRESSED GENES IN *Brassica juncea* AND *Camelina sativa* UPON *Alternaria brassicae* CHALLENGE

N.M.C. Nayanakantha^{1*}, S. Rawat² and A. Grover²

¹*Rubber Research Institute of Sri Lanka, Dartonfield, Agalawatta, Sri Lanka*

²*National Research Centre on Plant Biotechnology,
Indian Agricultural Research Institute, New Delhi, 110012, India*

Introduction

The necrotrophic pathogen *Alternaria brassicae* causes the globally devastating alternaria black spot disease of Indian mustard (*Brassica juncea*), an important oilseed crop. No satisfactory source of resistance against *A. brassicae* has been identified among crop *Brassica* species. However, resistance to *A. brassicae* has been found in distantly related species of the family Brassicaceae such as *Camelina sativa*. A large number of defense-related genes are up- or down-regulated during plant-pathogen interactions. Therefore, it is important to understand the network of genes that govern the resistance to *A. brassicae* in *C. sativa* as compared to *B. juncea*. In this context, suppression subtractive hybridization (SSH) and macroarray (reverse Northern blot) techniques were employed to identify putative defense-related genes expressed during the compatible *A. brassicae* - *B. juncea* and incompatible *A. brassicae* - *C. sativa* interactions.

Materials and Methods

Forty five days old plants of *C. sativa* and *B. juncea* were inoculated with *A. brassicae*. Leaf samples were collected at different time intervals viz., 0, 2, 4, 8, 12, 24, 48, 72 and 96 h post inoculation. Control samples were collected from sterile distilled water treated plants. Total RNA was isolated from the *A. brassicae* inoculated (tester) and water-inoculated (control/driver) leaf tissues of *B. juncea* and *C. sativa* using TRIzolTM Reagent (Invitrogen, USA). For each plant species, Poly(A)⁺ RNA was purified separately from pooled RNA from infected samples using an Oligotex mRNA Midi Kit (Qiagen, US). SSH was performed using the PCR-Select cDNA Subtraction Kit following the guidelines of the manufacturer (Clontech, USA). Subtracted cDNA were inserted directly into pGEM-T Easy T/A cloning Vector (Promega, USA) and then were transformed into *Escherichia coli* DH5 α competent cells (New England Bioscience, UK) in Luria Agar plates with ampicillin (100 mg L⁻¹), X-gal (20 mg mL⁻¹) and IPTG (200 mg mL⁻¹).

The cDNA inserts of the recovered positive clones, based on blue/white color selection were amplified by colony PCR in a Thermal Cycler (MJ Research,

* nayanakanthachamil7@gmail.com

USA) using M13 forward and M13 reverse primers. All PCR products were analyzed by electrophoresis on 1.0 % agarose-gel. The PCR products of the recovered clones of ≥ 300 bp showing a single band were spotted onto Hybond-N+ nylon membranes (Amersham International, UK) in 96-well format in duplicate. Single stranded cDNAs from all four RNA populations were synthesized by using a first strand cDNA synthesis kit (Thermo Scientific, USA). The radio-labeled double stranded cDNAs were synthesized using a decalabelling kit (Thermo Scientific, USA) and $\alpha^{32}\text{P}$ -dCTP (BRIT, India). Nylon membranes were hybridized with denatured control and experimental cDNA probes. The membranes were washed and then exposed to Kodak X-ray films and stored at -80°C for 2–3 for autoradiography. Membranes were scanned and the images were quantified by Image quant software (Imagequant Inc., USA). Those showing >2 fold differential expression were selected for sequencing in an ABI3700 capillary sequencer. Raw nucleotide sequences were manually processed for base calling, vector masking, and ambiguous nucleotides. The Lasergene 7-1 software was used to assemble the good quality ESTs. ESTs were compared with database nucleotide collection (nr/nt) using blastx and blastn (<http://www.ncbi.nih.gov>). Computational annotation of the EST datasets was performed using the Blast2GO software V1.3.3 (<http://www.blast2go.org>.) with E value thresholds of $e-10$.

Results and Discussion

Suppression subtractive hybridization is a relatively simple and efficient tool that can be used to generate cDNAs enriched for differentially expressed genes of both greater and lesser abundance. Subtracted cDNA libraries identified 56 unigenes from *B. juncea* and 75 unigenes from *C. sativa* as shown in Table 1. Representative reverse Northern blots and an agarose gel photograph for colony PCR are shown in Plate 1. The number of transcripts coding for several proteins involved in plant defense (including pathogenesis related proteins) or biotic stress were significantly higher in the resistant *C. sativa* than in the susceptible *B. juncea*. Pathogenesis-related proteins viz., *Defensin*, *thionin* and *PR14* were identified in *C. sativa* while only *thionin* were identified in *B. juncea* (Tables 2 and 3). In most SSH studies with pathogen challenge in other plants, a large number of defense-related genes have been identified apart from major PR genes (Yu et al. 2010). In the present study also, defense-related proteins such as WRKY 70 transcription factor, MLP-like protein, serine/threonine kinase and metallothionin were identified in *C. sativa* but not in *B. juncea*. These proteins might play an important role in *C. sativa* for its resistance to *A. brassicae*. This is the first report on analysis of differentially expressed genes in *C. sativa* and *B. juncea* against *A. brassicae* by SSH technique.

Table 1. Complete sequence analysis of the subtractive cDNA libraries of *B. juncea* and *C. sativa*.

Species Name	No. of clones used for macroarray	No. of clones sequenced	Quality ESTs	No. of contigs	No. of singlets	Total No. of unigenes
<i>B. juncea</i>	672	195	130	29	27	56
<i>C. sativa</i>	768	205	146	46	29	75

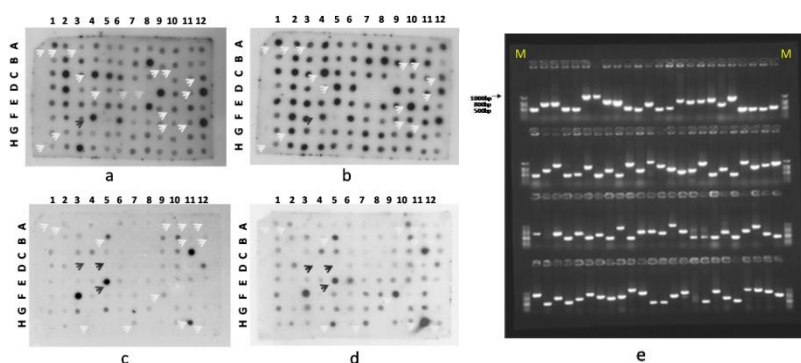


Plate 1. Representative differential screening results of macroarrays of the SSH libraries of *B. juncea* and *C. sativa* (a-d). (a): control cDNAs of *C. sativa*, (b): *A. brassicae* challenged cDNAs of *C. sativa*, (c): control cDNAs of *B. juncea*, (d): *A. brassicae* challenged cDNAs of *B. juncea*. A representative agarose gel photograph of colony PCR showing different insert lengths ranging from 400bp - >1kb (e).

Conclusions and Recommendations

In this study, a screening was performed for genes preferentially expressed in resistant *C. sativa* compared to that in susceptible *B. juncea* challenge inoculated by *A. brassicae*. Some of those genes were found up-regulated in both plant species and may play an important role during *A. brassicae* infection. Further characterization and functional analysis of these putative genes will enhance our understanding of the molecular mechanisms occurring in resistant *C. sativa* and susceptible *B. juncea* during *A. brassicae* infection. It will provide novel target genes for the development of *A. brassicae* resistant transgenic *B. juncea* through genetic engineering. This study will undoubtedly

Table 2. Defense/ biotic stress-related genes identified from the SSH library of *B. juncea* using *Blas2GO* analysis

Contig No.	Accession No. of matching sequence	Matching sequence	Matching sequence from data base	Sequence identity (%)
7	ACH88385.1	<i>B. oleracea</i> (var gemmifera)	vegetative storage protein vsp2	81.6
41	ABO32545.1	<i>B. oleracea</i>	lipoxygenase (LOX)	98
44	ACJ68108.1	<i>B. napus</i>	thionin	84.2

shed light on defense signaling pathways in *C. sativa* conferring its resistance

Table 3. Defense/biotic stress-related genes identified from the SSH library of *C. sativa* using *Blas2GO* analysis

Contig No.	Accession No. of matching sequence	Matching sequence	Matching sequence from data base	Sequence identity (%)
4	NM_110996.2	<i>A. thaliana</i>	serine/threonine kinase	86
5	NP_001077806.1	<i>A. thaliana</i>	mlp-like protein 28 (miraculin)	85.8
40	NM_102161.3	<i>A. thaliana</i>	polyketide cyclase dehydrase and lipid transport-like protein (PR14)	79.2
44	NP_181879.1	<i>A. thaliana</i>	defensin like proteins	79.4
45	NM_111773.3	<i>A. thaliana</i>	metallothionein	74
52	NP_191199.1	<i>A. thaliana</i>	WRKY transcription factor 70	76
67	ACJ68108.1	<i>A. thaliana</i>	thionin	80.2

to *A. brassicae*.

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RELATIONSHIP BETWEEN RENAL FUNCTION PARAMETERS CYSTATIN C, SERUM CREATININE AND BLOOD UREA WITH BODY FAT CONTENT MEASURED BY BIO- IMPEDANCE ANALYSIS OF APPARENTLY HEALTHY FEMALES LIVING IN A SELECTED URBAN AREA OF COLOMBO DISTRICT

M. S. N. F. Nihla^{1*} and P. P. R. Perera¹

¹*Department of Biochemistry, University of Sri Jayewardenapura*

Introduction

The term 'Renal function' indicates the state of kidney and its efficiency to filter blood. Loss in renal function leads to renal failure which is mainly caused by diabetes or hypertension. Obesity is a leading cause for diabetes and hypertension so as for renal diseases. The prevalence of renal disease is increasing dramatically in the world. Between 8-10 % of the adult population have some form of kidney damage, and every year millions die prematurely of complications related to chronic renal diseases (Dupuis 2013).

Body fat content can be accurately determined by bio impedance analysis (BIA) because it has the ability to distinguish fat from muscle. BIA measures percentage body fat, visceral fat content and percentage fat distribution in different compartments of the body by sending an electrical pulse throughout the body. Therefore, BIA can define obesity and central obesity more accurately than anthropometric measures. Renal function could be screened through simple routine tests. Cystatin C, serum creatinine and blood urea are used as biomarkers of kidney dysfunction. Detecting kidney dysfunction at an earlier stage will prevent or slow down the incidence of chronic renal disease and mortality rate. No research has been recorded in Sri Lanka regarding the association of kidney function and BIA. The main objective of this study was to determine the relationship of renal function parameters with body fat percentage and visceral fat content of apparently healthy females living in a selected urban area of Colombo district.

Materials and Methods

This study was carried out as a descriptive cross sectional study in a community by collecting data from 132 apparently healthy females of Colombo district with no diagnosed long term diseases and who were between 18 to 65 years of age. Sample size was calculated using an online sample size calculator (www.raosoft.com/samplesize.html). Study instruments used were an interviewer administered questionnaire, non-stretchable commercial measuring tape and a Bio electrical impedance

* nihal.nilufer@ymail.com

analyze machine (Omron Karada Scan Body Fat Analyzer HBF -358, Omron, Japan)

An information sheet and a consent form were given to the participants and the procedure was explained to them. Written consent from participants was obtained and data collected using an interviewer administered questionnaire. Bioelectrical impedance analyzer was a portable machine which measures the total body fat percentage, visceral fat content and fat distribution percentages in different body compartments by sending an electrical impulse throughout the body. The machine was kept on a flat surface and the participant was requested to stand on the machine bare footed, holding machine at arm's length for about 1 min. Height of the participant was fed into the machine and readings obtained. Blood samples were collected by a trained person under aseptic conditions. Blood was drawn from the arm in a sitting position using a sterile syringe after cleaning the area with surgical spirit. Then serum was separated and stored at -20 °C until analysis. Blood urea, serum creatinine and serum cystatin C were analyzed using a fully automated Konelab[®] analyzer.

Data were analyzed using the statistical package for social sciences (SPSS) version 16 and Microsoft excel software. Descriptive analysis was performed to identify the prevalence of different variables (cystatin c, serum creatinine, blood urea, body fat percentage and visceral fat content). One way analysis of variance was used to compare the means of continuous variables. Pearson's *Chi*-squared test was used to compare the frequency of categorical variables between groups. Pearson/Spearman correlation was used to analyze the correlation between variables where appropriate. A *p*- value < 0.05 was considered statistically significant. Ethical clearance for the study was obtained from the Ethics Review Committee, Faculty of Medical Sciences, University of Sri Jayewardenepura (Approval No. HB-04-13).

Results and Discussion

Among 132 participant's 40 persons belonged to Sinhala ethnic group and 92 persons to Muslim ethnic group. Table 1 gives the average values of study variables. The study population is at an increased health risk to obesity related diseases due to their high amount of fat deposition in the body. According to the frequency distribution of body fat percentage classification of the study, about 28.0 % of the study population had a 'high' body fat percentage and 59.8 % of the population had a 'very high' body fat percentage. Only 10.6 % had 'normal' body fat percentage and 1.5 % had a 'low' body fat percentage. This increased body fat percentage of the population reflects the increase in obesity of this population.

Table1. Average values of bio impedance analysis variables and renal function parameters

Variable	Mean	Reference Value
Body Fat Percentage (%)	35.6	Very High Low: 5.0- 19.9 Normal: 20.0- 29.9 High: 30.0- 34.9 Very High: 35.0- 50.0
Visceral Fat	9.1	Normal Normal: 1- 9.9 High: 10- 14.9 Very High: 15- 30
Cystatin C (mg L ⁻¹)	0.95	Normal 1-50 years: 0.55- 1.15 >50 years: 0.63- 1.44
Serum Creatinine (mg dL ⁻¹)	0.74	Normal 0.6- 1.1
Blood Urea (mmol L ⁻¹)	3.7	Normal 2.1- 7.1

Colombo district is a highly urban area and residents of Colombo district have a life style which leads to obesity. Consumption of fast food and lack of exercise leads to obesity. Females have a higher prevalence of obesity than males and this may be another reason for the higher proportion of obesity in this study population.

Table 2. Correlations between renal function parameters and BIA variables

	Correlation coefficient (r)	Significance
Cystatin C with Body fat %	0.195	0.025
Cystatin C with Visceral fat content	0.039	0.660
Serum Creatinine with Body fat %	-0.067	0.444
Serum Creatinine with Visceral fat content	-0.139	0.112
Blood Urea with Body fat %	-0.011	0.901
Blood Urea with Visceral fat content	-0.121	0.167

Visceral fat content of the individual reflects their abdominal obesity. According to the frequency distribution of visceral fat content of the study, it shows that 60.6 % of the study population had a 'normal' visceral fat content, 21.2 % of the population had a 'high' visceral fat content and 18.2 % had a 'very high' visceral fat content. Another study done in Sri Lankan females in

2008 using DXA scan indicate an average body fat mass of 20.3 kg and 7.7 kg of visceral fat mass (Weeraratna et al. 2008).

Research studies have shown that renal function decreases with increasing body fat percentage (Oh et al. 2014). The present study showed a significant correlation ($p=0.025$) between participants' cystatin C concentration with their body fat percentage. Neither Serum creatinine or blood urea had a significant relationship with body fat percentage when analyzed with Pearson's correlation and Pearson's chi-square tests.

This study does not show any association for visceral fat content of the participants' and their cystatin C, serum creatinine and blood urea concentrations. This contradicts a study done to examine the association of visceral fat with renal function which revealed that renal function has a significant association with and visceral fat while using cystatin C and not with serum creatinine (Young et al. 2008).

Conclusions and Recommendations

Renal function parameter cystatin C significantly correlates with the individual's bio impedance variable body fat percentage. Renal function parameter serum creatinine and blood urea does not relate with participants' body fat percentage. Renal function parameters cystatin c, serum creatinine and blood urea do not shows a relationship with visceral fat content of the participants.

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COMBINED APPLICATION OF 1-METHYLCYCLOPROPENE (1-MCP) AND ASCORBIC ACID BEST IMPROVES THE POSTHARVEST LONGEVITY AND QUALITY OF *Alstroemeria* SPP. CUT FLOWERS

**B.S. Obadamudalige^{1*}, C.K. Beneragama², G. Wijesuriya²
and S.M.M.R. Mawalagedera¹**

¹*Department of Plant Sciences, Faculty of Agriculture, Rajarata University of Sri Lanka, Puliyankulama, Anuradhapura, Sri Lanka*

²*Department of Crop Science, Faculty of Agriculture, University of Peradeniya, Peradeniya, Sri Lanka*

Introduction

Alstroemeria spp., commonly known as Peruvian Lily (Choon 2012), is a highly demanded cut flower due to availability of numerous and vibrant colors. *Alstroemeria* is an ethylene sensitive genus in which the ethylene induced postharvest degradation symptoms include; petal wilting, transparency of petals, petal drop, premature yellowing of leaves and short life of leaves (Choon 2012). These changes reduce the economic value of *Alstroemeria* spp. cut flowers. The ethylene induced postharvest degradation of cut flowers can be mitigated by applying chemicals, which inhibit ethylene biosynthesis and/or block ethylene receptors. 1-Methylcyclopropene (1-MCP), a blocker of ethylene receptors, is structurally similar to ethylene that has a ten-fold more affinity to ethylene receptors than ethylene. Ascorbic acid (vitamin C) is an antioxidant, which donates electrons to neutralize free radicals. The objective of the study was to improve the vase life and to maintain postharvest quality of *Alstroemeria* spp. with 1-methylcyclopropene and ascorbic acid application alone and in combination.

Materials and Methods

Inflorescences of red *Alstroemeria* spp., with at least three florets were completely red were harvested from fields of commercial growers in Meepilimana, Sri Lanka. The inflorescences were treated either with 200 mL of 57 mM ascorbic acid or with 1-MCP (0.25 ppm) or with both together for six hours. The tested concentrations and exposure duration were optimized in preliminary studies (Obadamudalige et al. 2014). Vase life was counted as the number of days taken for the shedding of first petal in florets. The Percentage fresh weight loss (FWL) for inflorescences was calculated. Total chlorophyll content of leaf extracts and, anthocyanin and glucose contents were measured in petal extracts. The analysis of variance (ANOVA) procedure was performed for the vase life, percentage fresh weight loss and concentrations

* buddhikas143@gmail.com

of chlorophyll, anthocyanin and glucose. All analyses were performed using MINITAB 16.0. Significances were defined at $p < 0.05$.

Results and Discussion

Combined application of 1- MCP and ascorbic acid, gave the longest vase life of 10 days compared to all other treatments. All treatments showed longer vase life than that achieved in the control ($p=0.001$; Table 1). Ethylene induced senescence is delayed by 1-MCP since its ability to bind permanently to ethylene receptors while ascorbic acid can decrease ethylene synthesis, reduce respiration and transpiration rates in cut flowers (Serek and Sisler 2001). The extension of vase life by 1-MCP and ascorbic acid applied alone was seen in numerous other cut flower species; *Alstroemeria* spp., *Diathus caryophyllus* and *Curcuma alismatifolia* (Serek and Sisler 2001).

Table 1. Effect of treatments for 6 hrs on the vase life of cut *Alstroemeria* spp. inflorescences

Treatment	Vase life (days)*
Control	3.5±0.2 ^c
1-methylcyclopropene (0.25 ppm)	8.25±0.4 ^{ab}
Ascorbic acid (57 mM)	7.5±0.6 ^b
1-methylcyclopropene + ascorbic acid	10.25±0.8 ^a

Note: *Mean±SE (n=3), different letters indicate significant differences between treatments ($p<0.05$).

Percentage fresh weight loss (FWL %) was not significantly different among treatments ($p=0.12$; Fig. 1a). The best postharvest concentrations of petal anthocyanins were maintained in 1-MCP treatment and in the combined treatment, which was one to two-fold higher than other treatments ($p=0.001$; Fig. 1b). Present study reflects the application of ascorbic acid alone had no effect on anthocyanin concentrations compared to the control (Fig. 1b). The reason behind that was, the combination treatment maintain high anthocyanin concentration throughout the period compared to the inflorescences treated with ascorbic acid alone.

Glucose concentrations were highest in petals from inflorescences treated with 1-MCP alone compared to the control (Fig. 1c). This may be due to 1-MCP decelerates or delay the ethylene-induced respiration climacteric thus maintaining more glucose in tissues (Chutichudet et al. 2010). However, petals from inflorescences treated with ascorbic and 1-MCP+ascorbic acid had glucose levels similar to those of the control (Fig. 1c).

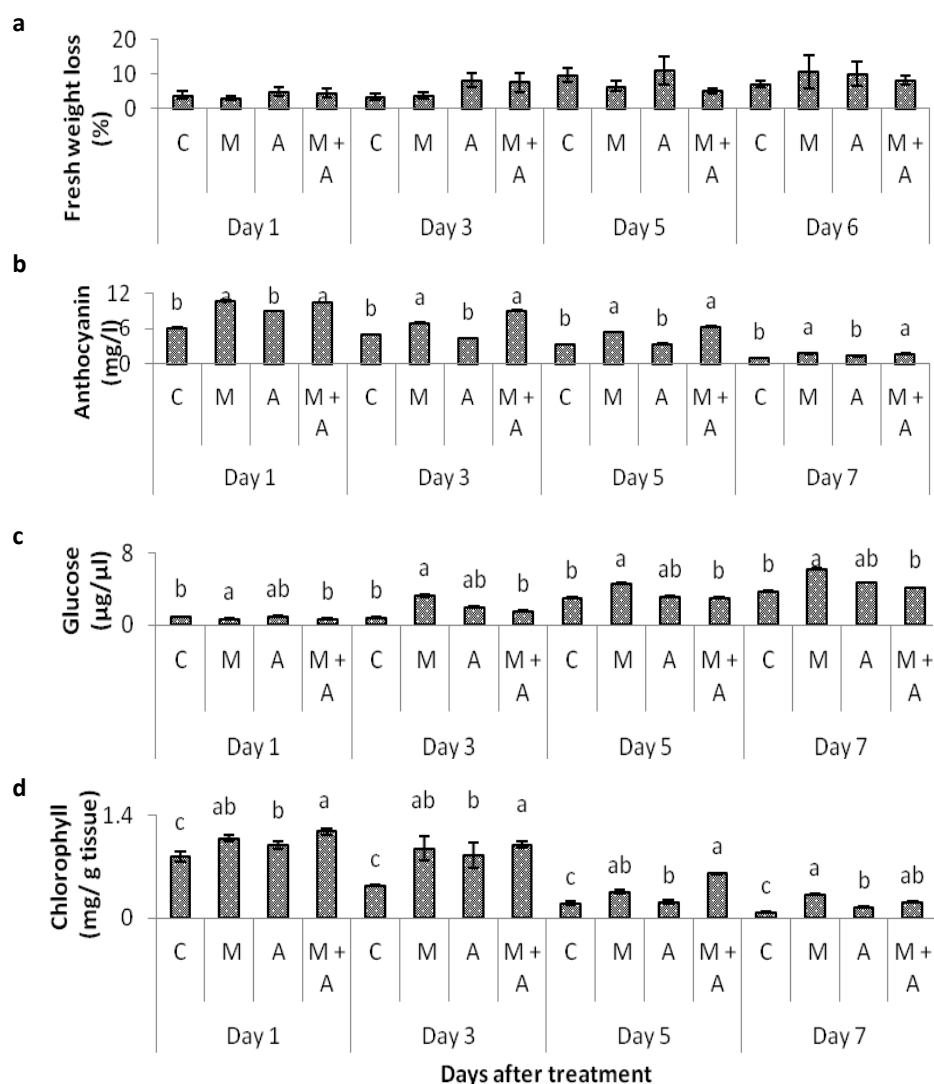


Figure 1. Effect of treatments on (a) percentage fresh weight loss, concentrations of (b) anthocyanin, (c) glucose and (d) chlorophyll in cut *Alstroemeria* spp. inflorescences measured one, three, five and seven days after treatments with (C) control, (M) 1-methylcyclopropene 0.25 ppm, (A) ascorbic acid 57 mM and the (M+A) combination of the two for 6 hrs. Mean \pm SE ($n=3$) Bars with different letters indicate significant differences across treatments within the day ($p<0.05$).

The combined treatment and 1-MCP alone, best maintained the chlorophyll concentration in leaves, which was five and three times higher compared to those in the control at the seventh day ($p<0.05$; Fig. 1d). The retention of green color in the leaves is an important postharvest quality parameter in

Alstroemeria spp. in which ethylene induces rapid degradation of chlorophyll. Blocking of ethylene receptors by 1-MCP may have inhibited the ethylene induced chlorophyll degradation in leaves of *Alstroemeria* spp.

Conclusions and Recommendations

Of the tested postharvest treatments of *Alstroemeria* spp., combined application of 1-MCP 0.25 ppm and ascorbic acid 57 mM was the most effective treatment for extending vase life and maintaining postharvest quality. Experiment can be repeated under different storage temperatures before, after and during 1-MCP and ascorbic acid treatments to improve vase life and postharvest quality of *Alstroemeria* spp. cut flowers.

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THE FACTORS HAMPERING THE INNOVATION IN SMALL AND MEDIUM SCALE FIRMS IN THE FOOD AND BEVERAGE INDUSTRY IN SRI LANKA

H.M.L. Peiris¹, G.H.I. Anjalee^{1*} and A.M.T.P. Athauda¹

¹*Department of Agribusiness Management, Faculty of Agriculture and Plantation Management, Wayamba University of Sri Lanka, Makandura, Gonawila*

Introduction

Food and beverages industry is one of the fastest growing sectors in the Sri Lankan economy and has the potentials to grow further in short and medium term. The product range of the sector consists of processed vegetables, fruits and juices, confectionary and bakery products, rice and cereals, beverages and animal fodder. In year 2011, the contribution to the Gross Domestic Product (GDP) by the food and beverages industry was recorded as 7.9 %. In Sri Lanka, there are 3340 registered small and medium scale food and beverage firms are located island wide and 55.2 % of them are established in the Western and North Western provinces of the country.

Innovation is considered as the backbone of entrepreneurship and widely recognized as a key strategy to gain competitive advantages against nations and firms. In entrepreneurial literature it is widely evident that, small firms are much more innovative than large scale firms. The ability to introduce innovation often depends on the characteristics of the small firm. Less bureaucracy, owner expertise, and intimacy between owners and customers can facilitate implementation of innovation. However, small firms are particularly restricted for introducing innovation by their limited physical resource base.

In this light, the study was focused on identifying barriers for innovation among small and medium scale firms in the food and beverages industry in Sri Lanka and in turn, to investigate the degree to which those barriers have an effect on the number of product, process and management innovations of food and beverages processing firms.

Materials and Methods

The data were collected from 155 managers or owners of small and medium scale firms in the food and beverage industry via face-to-face interviews conducted in the Western and North Western provinces covering five districts namely; Colombo, Gampaha, Kalutara, Kurunegala and Puttalam. The sample included six firm types namely; Tea and Non-Alcoholic Beverages (TNAB),

* ishara.anjalee@gmail.com

Coconut Products (CP), Dairy Products (DP), Fruits, Vegetables and Processed Foods (FVP), Bakery Products (BP) and Other firms (OF). The sample was selected through random sampling technique based on a list obtained from the Department of Census and Statistics, Sri Lanka. A questionnaire was structured and used both Sinhala and English versions and each innovation barrier was scaled using five point Likert Scale ranging from “strongly agree” to “strongly disagree” giving 5 to 1 value range to identify the magnitude of the each barrier to the entrepreneur.

Kruskal-Wallis test was conducted among the both internal and external barriers for innovation separately. An index was developed to measure the internal and external barriers. Ordinal Logistic Regression Model was used to evaluate the relationship between number of firm innovations and internal barriers, external barriers for innovation, firm age, average annual turnover, industry type, and number of employees.

Results and Discussion

The data on innovations were collected under three categories namely; product, process and management innovations. Regardless the firm types, majority of the firms were involved in introducing product innovations compared to other two categories (Fig. 1). Management related innovations were the least common type of innovations among the firms.

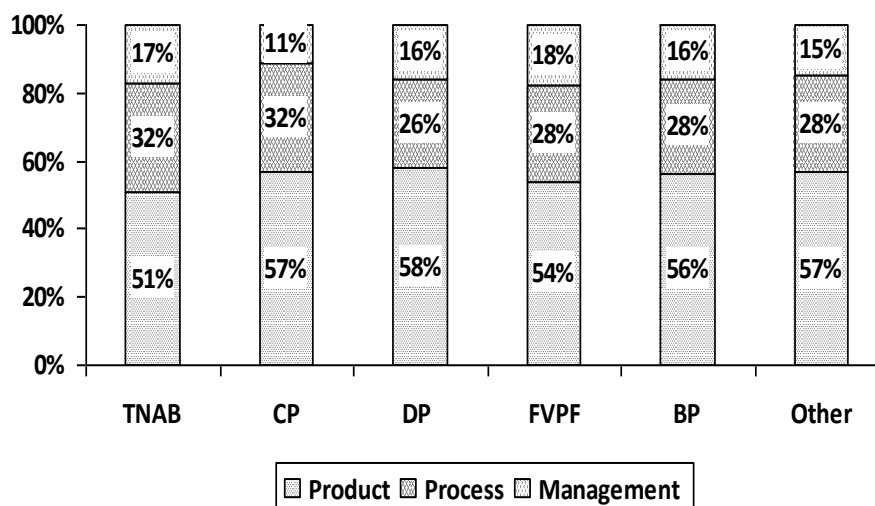


Figure 1. Product, process and management innovations in each firm type

Note: Non-Alcoholic Beverages (TNAB), Coconut Products (CP), Dairy Products (DP), Fruits, Vegetables and Processed Foods (FVPF), Bakery Products (BP)

The median values of barriers to innovation were examined by Kruskal-Wallis Test separately for two categories. The probability value ($P = 0.000$) for the

both analysis of internal and external barriers depicted that at least one internal/ external barrier is different from other internal/external barriers. Four out of eight internal barriers were having a median value of three or higher while, four out of six external barriers also had a median value of three or higher (Table 1).

Table 1. Results of Kruskal-Wallis test

Internal Barriers			External Barriers		
Barrier	MV	AR	Barrier	MV	AR
High cost	4.00	1101.9	Insufficient government support	4.00	602.1
Lack of qualified personals	4.00	878.7	Lack of information about technology	4.00	507.6
Difficulty of keeping qualified employees	3.00	1039.4	Economic turbulence	3.00	529.5
Manager resistance to change	3.00	636.3	Lack of market information	3.00	215.7
Poor accessibility to financial resources	2.00	830.4	Lack of regional infrastructure	1.00	715.1
Excessive risk	1.00	299.9	Lack of external partner opportunities	1.00	223.0
Lack of internal employee training	1.00	353.6			
Employee resistance to change	1.00	845.9			

Note: MV=Median Value; AR = Average Ranking

Table 2. Results of ordinal logistic regression

Variable	Coefficient	Probability
Internal barriers	-0.0352	0.605
External barriers	-0.0679	0.089*
Firm age	-0.3510	0.103
Firm type: TNAB	1.6623	0.072*
CP	0.8999	0.130
DP	0.9846	0.061*
FVPF	-0.7692	0.128
BP	-0.6021	0.584
Average Number of Employees: <5	0.7352	0.606
5 – 10	-0.1670	0.899
11 – 15	-0.6954	0.557
16 - 20	-1.4529	0.227
Average Annual Turnover : < Rs. 1 million	2.3479	0.056*
Rs. 1 – 2 million	0.4251	0.701
Rs. 2 – 3 million	-0.5050	0.632
Rs. 3 – 4 million	-0.3327	0.723

The ordinal logistic regression examined the relationship between number of innovations and number of internal barriers, external barriers, firm age, and industry type, average number of employees and average annual turnover of firm (Table 2) and the results indicate that the internal barriers are insignificant to innovation hence have a negative relationship with number of innovations. External barriers were significant at $\alpha=0.1$ and the negative coefficient suggests that there is a negative relationship with number of innovations. Among firm types, TNAB and DP were significant and have a positive relationship to number of innovations. Out of average annual turnover, only less than one million rupees was significant and has a positive relationship to innovations.

Conclusions and Recommendations

For a small and medium scale firm in food and beverage industry, the major internal barriers for innovation are high cost, lack of qualified personals, difficulty of keeping qualified employees and manager's resistance to change. The critical external barriers were insufficient government support, economic turbulence, lack of market information and lack of information about technologies. Only external barriers were found to be significantly associated with number of innovations. Relationship of both internal and external barriers showed a negative impact on the number of innovations introduced by the firm. Among the significant external barriers, the highest ranked barriers were insufficient government support and lack of information about technology. The findings of this study open avenues for relevant authorities and policy makers to re-think on the growth and development of food and beverage sector. Even though, food and beverage sector has been identified as one of the fastest growing industries in the Sri Lankan economy, these barriers to introduce new products and processes would one day affect on hampering that progress. Therefore, it is recommended that the government should make the policy decisions regarding the growth and development of small and medium scale firms in the food and beverage industry in Sri Lanka in order to harvest the optimum benefits from the sector.

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PROVISION OF LOW COST GELLING AGENTS FOR *IN VITRO* SEED GERMINATION OF SUNFLOWER (*Helianthus annuus* L.)

P.C.D. Perera¹ and N. Dahanayake^{1*}

¹*Department of Agricultural Biology, Faculty of Agriculture, University of Ruhuna, Mapalana, Kamburupitiya, Sri Lanka*

Introduction

In-vitro cultivation of plant tissues is generally carried out in a solid or semi-solid nutrient medium, with gelling agents. Unreasonably price of agar and fear of over-exploitation of its resources necessitated the search for low costs materials as alternative to agar. During last two decades, there has been increase in the efforts to look for suitable substitutes for agar (Singh et al. 2013). Gelling agent such as agar which is usually added to increase media viscosity contributes 70 % of the media costs (Mohomad et al. 2009). Various brands and grades of agar, agarose, phytigel and gelrite were used for *in vitro* propagation. Agar, the conventional gelling agent, has a number of drawbacks that negatively affect culture growth and differentiation in many cases (Mohomad et al. 2009). Cheaper agar alternatives include various types of starch and gums, which have been investigated in commercial micro-propagation. For example gelrite can be replaced with starch gelrite mixture. Other options include white flour, laundry starch, semolina, potato starch, rice powder and sago (Mohomad et al. 2009) and corn starch, tapioca, isubgol, guar gum, xanthan gum and karaya gum (Singh et al. 2013).

Sunflower (*Helianthus annuus*, Asteraceae) is an important oil seed crop in the world after soybean and groundnut. Wheat flour (*Triticum aestivum*), corn flour (*Zea mays*), cassava flour (*Manihot esculenta*), rice flour (*Oryza sativa*), potato (*Solanum tuberosum*), barley flour (*Hordeum vulgure*), gelatin (edible) and stabilizer (uses in ice-cream preparation) is easily available in the market of Sri Lanka and have a possibility to be used as potential solidifying agents. Therefore, present study was carried out to evaluate the feasibility of various commercial products; agar, wheat flour, corn flour, cassava flour, rice flour, barley flour, gelatin and stabilizer as solidification agents to minimize the cost of gelling agents.

Materials and Methods

Plant materials: Seeds of Sunflower were purchased from the Seed and Planting Material Division, Department of Agriculture, Sri Lanka. First, seeds rinsed with distilled water and then washed using 70 % (v/v) alcohol for 3 min. Thereafter, the seeds were soaked in 20 % (v/v) Chlorox solution

* daha_27@yahoo.com

(Sodium hypochlorite) for 20 min. Those seeds were rinsed in sterile distilled water for 4-5 times. They were dried onto sterile filter papers. Seed coat was removed with sterile scalpels and pliers. Seeds were introduced to glass tubes (1 seed per tube) containing 3ml of hormone free Murashige and Skoog (MS) basal medium, which was solidified by different gelling agents as stated earlier. The culture medium used for all the experiments was based on MS medium with 30 gL⁻¹ sucrose (Dahanayake et al. 2012) and addition of 100 gL⁻¹ different gelling agents (agar - control, wheat flour, corn flour, cassava flour, potato flour, rice flour, barley flour, stabilizer and gelatin) without plant growth regulators. Medium was autoclaved for 21 minutes at 121 °C after adjusting the pH to 5.8.

Experimental design: The lengths of the aseptically raised plant seedlings and number of seeds germinated were recorded after four weeks period. Experiment was repeated three times. All experiments reported here were repeated at least five times with three replicates. Statistical analysis was performed with Duncan's Multiple Range Test using SAS software (version 9.1.3). The cost analysis was performed using the following formulae;

$$\text{Cost deduction compared with agar (\%)} = 100 - \left[\frac{\text{cost of alternate gelling agent / kg}}{\text{cost of agar / kg}} \right] \times 100$$

Results and Discussion

Seedling height as influenced by different gelling agents: The highest seedling height of 6.83 cm observed at the end of the fourth week in MS medium solidified with agar was not significantly different from corn flour (6.43 cm). The lowest seedling height was observed in medium solidified with gelatin (0.12 cm). In addition, treatments rice flour (1.6 cm) and wheat flour (1.73 cm) were not significantly different from each other, while cassava flour (0.77 cm), stabilizer (0.29 cm), grind potato (0.18 cm) and gelatin (0.12 cm) were also on par with each other (Table 1).

Seed germination percentage: Highest numbers of seed germination percentage were observed in medium solidified with agar (85 %). Second highest number of seed germination percentage was corn flour (80 %) while third was observed by barley (70 %). Agar, corn flour and barley flour were not significantly different. Number of germinated seeds percentage in wheat flour (40 %) and rice flour (40 %) were not significantly different from each other; besides, cassava flour (10 %), stabilizer (10 %), gelatin (5 %) and grind potato (5%) also not significantly different with each other (Table 1).

Corn flour has been found to be the best alternative to agar among used gelling agents. Similar results were observed by Henderson and Kinnersley (1988) and Daud et al. (2011). Growth and differentiation of plant cell cultures was increased when media were gelled with corn flour instead of agar.

Table 1. Effect of different gelling agents on seed germination and seedling growth of sunflower

Treatment (Gelling agent)	Mean height of the seedlings (cm)	Seed germination (%)
Agar	6.83 ^a	85 ^a
Corn Flour	6.43 ^a	80 ^a
Barley Flour	3.77 ^b	70 ^b
Wheat Flour	1.73 ^c	40 ^c
Rice Flour	1.60 ^c	40 ^c
Cassava Flour	0.77 ^d	10 ^d
Stabilizer	0.29 ^d	10 ^d
Potato Flour	0.18 ^d	05 ^d
Gelatin	0.12 ^d	05 ^d

Note: Means followed by the same letter in each column are not significantly different at 5% level with Duncan's Multiple Range Test.

Higher yield of anthocyanin and dry weight of embryos were found in wild carrot cultures grown on media gelled with corn starch. The starch-mediated increase in growth and differentiation of wild carrot cells was accompanied by an increase in density of the cultures shown by higher dry weight/fresh weight ratios (Henderson and Kinnersley 1988). Furthermore the present study revealed that corn flour was better than agar when considering mean weight of plantlets and number of seeds germinated. However, Daud et al. (2011) observed that wheat flour and rice powder were better than corn starch. In the present study corn flour was the best gelling agent than barley, rice flour and wheat flour.

Table 2. Comparison of cost different gelling agents

Treatment (Gelling agent)	Price of 1 kg gelling agent Rs.	Concentration of gelling agent gL ⁻¹	Cost deduction compared with agar %
Agar	12,000	8	0.00
Corn Flour	650	100	94.58
Barley Flour	522	100	95.65
Wheat Flour	110	100	99.08
Rice Flour	68	100	99.43
Cassava Flour	50	100	99.58
Stabilizer	735	100	93.88
Potato Flour	160	100	98.67
Gelatin	1,800	100	85.00

Cost analysis of agar and corn flour: Economics of using alternative gelling agents was calculated by comparing the standard price of agar (Table 2). The cost of 1 kg of agar was about Rs. 12,000.00 Sri Lankan currency and it is the most expensive gelling agent compared to the other gelling agents tested in this study. Use of corn flour, as an alternative solidifying agent, reduced 94.6 % of expenditure made towards purchase of gelling agent i.e. agar. Comparatively other gelling agents also reduced around 85 %-99 % of expenditure than agar.

Conclusions and Recommendations

Plant height and seed germination percentage between corn flour and agar were higher than the other gelling agents and were not significantly different. Corn flour (100 gL⁻¹) has shown a greater potential as a cheaper alternative gelling agent for agar based medium for *in-vitro* seed germination of Sun flower.

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IMPACT OF CARTOON MOVIES ON BEHAVIOUR OF SCHOOL CHILDREN: A CASE STUDY OF MIHINTHALE DS DIVISION

D. A. M. Perera^{1*}, H.A.B.W. Hettiarachchi² and R.G.S.K Udayangani²

¹*Department of Accountancy, Wayamba University of Sri Lanka*

²*Department of Social Sciences, Rajarata University of Sri Lanka*

Introduction

Children spend most of their leisure time in front of television watching cartoon programs. Since it is the most frequent and easily accessible source of entertainment some parents use cartoon movies to keep the children occupied while they are busy with their work. Therefore, time which was previously spent by children in outdoor activities is now replaced with watching cartoons. As a result, most of children in Sri Lanka begin watching cartoons at an early age and by the age of three or four children become enthusiastic viewers. This has become a problem in Sri Lanka, because many children who are becoming addictive to some aggressive cartoons on television have become violent. Many children are attracted through contents of the cartoon programs and they inculcate some positive and negative habits in them (Ali and Muhammad 2013). Among the others, violence is a vital part of most of the cartoon programs, and therefore, children might be induced and attracted by violent content in cartoon movies. Hence, some scholars argue that cartoons have only negative impacts on children (Kellogg 1992; Anderson 2003; Saturnine 2004). Many parents are also in the view that some cartoon movies persuade children's behaving antagonistic and disreputable. Alternatively, some scholars argue that cartoons have a tremendous impact in improving the cognitive aspects among school children (Ginmann 2003). These contradictory versions show the research gap in this field. Considering the importance of exploring the effects of cartoon on the children's psychological development, and their behavior patterns, the main objective of this study was to trace the impact of television cartoon programs on the behavior of the school children.

Materials and Methods

Fifty seven school going children of the age group 6 -13 years were randomly selected from two government schools, situated in Mihinthale DS division of Anuradhapura district of Sri Lanka, as the sample of this study. Considering the sensitiveness of the respondents, this study was designed on survey method and most of the data were obtained through observations. Few focus group discussions also conducted with the parents and the teachers to gather the necessary information. Before starting the data collection, researchers have visited these schools several times and had few focus group discussions

* methsila240@yahoo.com

with the teachers in the primary section and with some parents. The necessary measures were also taken to minimize the influence on children's behavior during the data collection stage. The selected children were given few blank papers and crayons and then, they were requested to draw their protagonists. Further, they were given a chance to sign a song in front of other students. No instructions or guidance were provided for the students; they were given the maximum freedom to paint their heroes as they like. While they were painting their pictures, researchers also checked the children's costumes and accessories, such as school bags, pencil cases, lunch boxes, water bottles, cover pages of books. The main purpose of checking their costumes and accessories were to explore the pictures (i.e. different cartoon characters) printed on them. The gathered data were analyzed by using non-parametric test in Statistical Package for Social Sciences (SPSS) software version 17.

Results and Discussion

This study has explored that the violence presented in cartoons programs have impacted on children behavior. Most of school children observed during the data collection have imitated their favorite cartoon characters in their leisureliness. Since they were requested to entertain the peers and the audience, 27 students (47 percent) caroled different songs as they like. Interestingly, the majority (59 percent, $n=16$) of them have used songs in cartoon movies to entertain their peers, and most of those presentations were comparatively aggressive. They have imitated the movements of different cartoon characters. Moreover, it was revealed that they force their parents to buy the same costumes or accessories as displayed by different cartoon characters. It was observed that 81 percent of the children are attracted by the dresses of cartoon character and most of the accessories of these students contained different cartoon characters were printed on.

There was no significance difference ($p<0.05$) among the groups on measure of selecting cartoon movies. Yet, boys preferences exhibited significantly more antagonism than girls ($p<0.05$). Violence is also integral part of the even those cartoons which are comedic in their genre (Ali and Muhammad 2013). Many boys (54 percent) indicated that they like to fight with their classmates and other children after watching these comedic fights in cartoon movies. Most of the students (68 percent) spend daily 1 to 2 hours in watching cartoons on television in their leisure time. Further it revealed that the cartoon movies dubbed in Sinhala is the most favorite cartoons of 91 percent children. Based on these figures, it can be understood that cartoon watching is the most favorite hobby of the school children (6-13 years of age) in Mihinthale DS division.

It also revealed from the study that '*Ben Ten*' is the most popular cartoon among the boys with the average of 78.6 percent. The most favorite cartoon character of the school going boys is boy in Ben-10. The girls have indicated that they prefer to watch '*Heda Weda Kello*' with the average of 55.2 percent. Their most favorite cartoon character is '*Meena*' with other two girls in '*Heda Weda Kello*' cartoon with the average of 51.7 percent. Majority of the children (63 %) like to watch comedy and a very low number like to watch love and action in cartoon programs. Interestingly, most of these findings compliance with the outcomes of two studies carried out by Ali and Muhammad (2013).

Conclusions and Recommendations

At present, cartoon movies are most favorite program on television among the school going children of the age group 6-13 years. Cartoons have changed drastically over the years but have their lasting effects on children. Behavior of the children in school is influenced by watching different cartoon programs. Since they have being affected by the cartoons that they usually watch, they like dresses and accessories of their favorite cartoon characters. Due to their tender minds, most of the children try to imitate the actions performed by different characters. Consequently, confirming the findings of Hapkiewicz and Aubrey (1971), this study revealed two facts; the first fact is that there exists strong association between watching cartoon movies and the behavior of the school children in class. The second fact is the violence present in the cartoons has strong association with the behavior of the children. Yet, most of parents are completely unaware the effects of cartoon on their children's psychological development, and later on their behavior patterns.

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PHYSICAL, CHEMICAL AND MICROBIAL ANALYSIS OF BOTTLED DRINKING WATER IN JAFFNA PENINSULA

K. Ragila^{1*}, P. Abiman¹, A. C. Thavaranjit², and P. Iyngaran¹

¹*Department of Chemistry, Faculty of Science, University of Jaffna, Sri Lanka*

²*Department of Botany, Faculty of Science, University of Jaffna, Sri Lanka*

Introduction

People in Sri Lanka have been using bottled drinking water few years. They expect bottled drinking water to be free of microbial contamination and health hazards. People in Jaffna peninsula, depend mainly on ground water for their drinking and other domestic needs due to the unavailability of other water sources such as waterfalls and rivers, and fresh water ponds and insufficiency of water available from rainfall (Mageswaran et al. 2004). However, the ground water in Jaffna is in danger due to over exploitation and pollution caused by excessive usage of agrochemicals and fertilizer (Velauthamoorthy 2001; Balasanthiran 2005). When A9 road was opened after three decades of war, the people in Jaffna started to use bottled drinking water. With the increase of its popularity and usage of its, questions have been raised about it's the quality and safety of bottled drinking water. Hence this study was conducted to analyze the microbial contamination, physical properties and chemical contents in different brands of bottled water being sold in Jaffna peninsula.

Materials and Methods

Five different batch numbers of each brand were brought from randomly selected grocery stores in Jaffna peninsula. Sample bottles from a total of 8 brands (labeled as A, B, C, D, E, F, G, and H) were selected for the study; and stored at room temperature (25-30 °C) and the samples are analyzed for physical, chemical and microbiological requirements as follows within 1-6 months from the date of manufacture. Color was measured by visual comparison method; odor and Taste were measured through sensory evaluation method. The amount of phosphate was measured using a Colorimeter (6051 colorimeter, Jenway, UK). The Electrical Conductivity was measured with a conductivity meter (Orion 4 star pH, Conductivity Bench top, Singapore) for analysis. The pH values of the bottled water samples were measured with a pH meter (PHS-3BW pH/mV/Temperature Meter). Alkalinity and Chloride were determined by the titrimetric method. Calcium, Hardness and Magnesium were measured by EDTA titration method. The values of Potassium were measured with the Flame photometer. Number of coliforms and faecal coliforms were determined by the standard membrane filter

* kragila88@gmail.com

method and the occurrence of *Escherichia coli* was also tested. Finally results were compared with the recommended Sri Lankan Standard (SLS) values (Theivendirarajah 1990). The obtained results were analyzed by the t-test ($\alpha=0.05$ level) and mean values and standard deviations were calculated.

Results and Discussion

Physical and chemical properties related to analysis are shown in Table 1. The pH values of the 8 different brands of bottled water samples varied from 6.1 to 7.4 and the recommended range is 6.5 to 8.5. Electrical Conductivity content varied from 46.8 to 349.0 $\mu\text{S cm}^{-1}$, which is far below compares to the SLS recommend value of 750 $\mu\text{S cm}^{-1}$. Alkalinity of the water samples varied from 76.5 to 284.8 ppm and the SLS recommended value is 400 ppm. Calcium concentration of the water samples varied from 11.4 to 26.6 ppm, which lies within the permit margin of SLS, which allow the calcium concentration up to 100 ppm. Magnesium concentration of the water samples varied from 14.5 to 24.5 ppm and the SLS permitted the magnesium concentration up to 150 ppm. Hardness of the 8 different brands of water samples varied from 99.8 to 145.5 ppm and the SLS recommended hardness is 400 ppm. Potassium concentration of the water samples varied from 0.3 to 1.6 ppm and the SLS permitted the potassium concentration up to 20 ppm. Phosphate concentration of the water samples varied from 1.3 to 2.4 ppm and the SLS permitted the phosphate concentration up to 2.0 ppm. Chloride concentration varied from 101.2 to 172.7 ppm and the SLS permitted the chloride concentration is up to 200 ppm. Salinity varied from 35.1 to 38.8 and the SLS recommended the salinity is up to 150 ppm. Only one batch of the brand C (2nd batch), E (2nd batch) and F (1st batch) had very high number of coliforms (data not shown) beyond the accepted level of the SLS. *Escherichia coli* was present in only one batch of the brand A. Faecal coliforms were observed only in two batches of the brand A among the brands tested, which was not accepted according to the standard. *Escherichia coli* and faecal should be absent. Coliforms should be less than 10 per 100 mL of sample.

Conclusions and Recommendations

The results of this study revealed that the values of electrical conductivity, chloride, salinity, alkalinity, calcium, magnesium, total hardness, and potassium in all of the water samples were within the acceptable limits of Sri Lankan Standards for drinking water. The mean value of phosphate ion of E, F, G and H brands exceeded the Sri Lankan Standard limit. The mean value of pH of A, D, F and G exceeded the Sri Lankan Standard limit. First and fourth batches of brand A had high faecal contamination and not suitable for consumption where as one batch of C, E and F had a risk of contamination and not suitable for drinking. When all the parameters are considered, two brands (B and C) out of eight brands analyzed are suitable for human consumption.

Table 1. Physical and mineral constituents of different brands of bottled drinking water (mean±st.dev)

Sample		A	B	C	D	E	F	G	H
Physical properties	Colour	Colourless	Colourless	Colourless	Colourless	Colourless	Colourless	Colourless	Colourless
	Odour	No Smell	No Smell	No Smell	No Smell	No Smell	No Smell	No Smell	No Smell
	Taste	Tasteless	Tasteless	Tasteless	Tasteless	Tasteless	Tasteless	Tasteless	Tasteless
	pH	6.5+0.1	7.1+0.4	6.7+0.1	6.4+0.2	7.4+0.4	6.1+0.2	6.6+0.2	7.4 +0.3
	Conductivity (µS/cm)	79.7+3.6	81.0 +10.9	48.6 +7.2	46.8 +1.3	122.0 +5.4	104.0+14.9	63.8 +1.9	349+122.1
Mineral constituents	Alkalinity (ppm)	159.6+11.9	159+25.3	109 +18.9	90.1+13.2	110.9+19.4	100.4+8.4	76.5 +16.7	285 +86.5
	Calcium (ppm)	14.3+1.3	17.3+0.9	11.4+0.7	15.8 +0.9	26.6 +0.6	17.1+1.1	12.2+0.7	18.0+4.4
	Magnesium (ppm)	18.0+0.9	16.1+1.0	19.6 +1.1	14.7 +0.7	14.5+1.1	21.5+1.5	17.8 +0.3	24.5 +2.1
	Hardness (ppm)	109.8+2.0	109.2+5.7	108.8+2.9	99.8+3.8	126.0+3.5	130.7 +3.4	103.4+2.1	145.5+14.3
	Potassium (ppm)	1.3+0.2	1.6+0.2	0.3 +0.2	1.2+0.1	1.5 +0.3	1.4+0.2	0.9+0.2	0.9 +0.2
	Phosphate (ppm)	1.5+0.3	1.3+0.3	1.5+0.3	1.7+0.0	2.3+1.1	2.1+0.5	2.4 +0.6	1.9 +0.7
	Chloride (ppm)	151.8+3.6	133.5+6.3	115.9 +5.8	115.9 +4.8	131.7 +6.4	156.4+11.2	101.2+7.7	173 +14.5
	Salinity (ppm)	37.7+0.2	36.8+0.3	35.9+0.3	35.9+0.3	36.7 +0.3	37.9 +0.6	35.1 +0.4	38.8+0.7

Note: 1ppm=1mg L⁻¹

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COGNITIVE DISSONANCE AS A FACTOR OF INFLUENCE FASHION

G.M. Ranathunga¹⁴¹

¹*Department of Textile and Clothing Technology, Faculty of Engineering,
University of Moratuwa*

Introduction

The Kandyan period (15th Century to 1815), in the annals of Sri Lanka is very unique since it was a melting pot of cultural fluctuations both local and foreign and as a result marked a signpost of the evolution of the country's fashion. The Portuguese, The Dutch, The British, South Indian and Siamese dress influences were mixed altogether. Rather than representing the hierarchy, the Sri Lankan sartorial etiquettes were juxtaposed during the reign.

The most prominent characteristic of the Kandyan dress was the hybrid formation together of different foreign dress influences. The royalty, especially the king had an exotic taste in foreign fashions; thus he displayed a unique individual personality. Elites also adopted a completely different foreign dress from that of the king; the dress conveyed dignity, their status and occupation. The king, the queen, the elite were in contrast to each other, thus reluctant to keep one line of a continuous tradition. Their dress provoked the viewer's curiosity, whether they were guided by their beliefs or behavior.

The aim of the present research is to identify influencing patterns in fashion from characters of king and queens of the Kandyan kingdom and incubating unique fashion concepts. It is important to study well established and well-practiced concepts in order to incubate new fashion concepts. The vision of their fashion gives new ideas to the field of fashion design.

Materials and Methods

Reliability of the data which was incorporated in the research is of much concern. Pictorial data were cross checked with different literary sources such as temple murals, particular literature and interviews. Original written sources and true pictorial evidences were used for the research. Temple murals were compared with contemporary sketches which were produced by foreigners (true observations). Finally the reinforced theory was compared with formally established theory in order to verify the reliability of findings.

Results and Discussion

Most of the early royalties and elites of the Kandyan kingdom had grown up under Portuguese protection due to the instability of sovereignty;

⁴¹ gayathrir@uom.lk

subsequently they were given Christian education and introduced to Christian society. Later they were baptized and took Western names such as did Queen Kusumasana Devi as Dona Chatherine, Prince Yamasinghe Bandara as Dom Philip and Konappu Bandara as Dom Juwan. The Dutch admiral Spilbergen who first visited the Kandyan territory recorded that king Vimala Dharma Suriya I (previously Dom Juwan) his queen Dona Catherine and their children wore Western dresses (Ferguson 1927). Prince Vijayapala of Matale and king Rajasimha II's dressed as Western gentlemen. Royalties who had direct foreign influences changed their dress. But however much they adopted a Western lifestyle their tradition, culture and religion were affected to a minimum in those etiquettes. Kandyan kingdom became the sole surviving link in the age long connection between Sinhalese power and the Buddhist religion. It was the heir to the traditional relationship between state and religion (Silva 2005). The society expected a Sinhalese Buddhist king. Therefore, it is obvious that mental tension existed between their desires/beliefs and their behavior. As a result a conflict arose between their beliefs and behavior. According to the convincing evidences some royalties of the Kandyan era had mental tension with their Western dressing habits. Some royalties changed their dress being inspired by foreign dresses. But later they found that foreign influences did not suit their culture. This is evident from prince Vijayapala's confession: "though I am a Chingala by blood I am a Portuguese in my way and my affection; it may well be that this is the chief reason for my losing my kingdom, treasures, the queen my wife, my son, and all that I possessed, at least reaching this state in which I see myself" (Pieris 1927).



Figure 1. King Vimala Dharma Suriya I (1591-1604), queen Dona Catherine (1602-1613 AD) and king Rajasimha II (1635-1687) clad in Western dresses

Pictorial and written evidences show that when such a situation arose, these concerned either changed their beliefs or behavior to reduce this conflict. Prince Vijayapala changed his beliefs to be able to be suited to his behavior of wearing foreign costumes and accepting the Christian faith. He

appreciated and considered western dresses as good and that a little cultural impact and western dress may be beneficial. The idea strengthened his adopted behavior influencing the power of foreign dress habits.

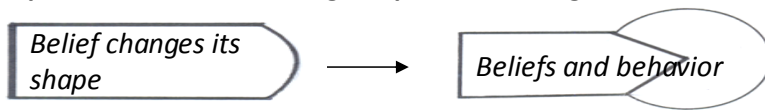


Figure 2. Change of beliefs to suit to behavior

In some instances, Kandyan Sinhalese Queens did not change their behavior. It was found that during the Nayakkar period Sinhalese queens never followed superior South Indian Queens' dress of *mottakkiliya*. Although Nayakkar queens were given a high position compared to Sinhalese queens the latter never wore *mottakkiliya* but followed the Sinhalese Buddhist traditional way of dressing. Sinhalese queens did not change their behavior as king Vijayapala did. This process maintained their original behavior of dressing habits and reduced the influencing power.

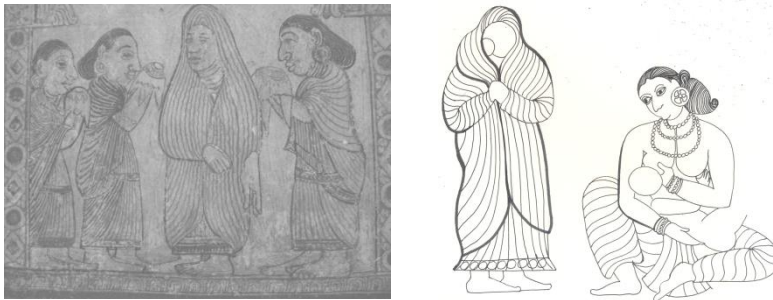


Figure 3. Murals of Degaldoruwa (Nayakkar queens clad in Mottakkiliya and Shinhalese queen clad in jacket and a cloth)

This process of influence is similar to learning theory describes in psychology, cognitive dissonance found by social psychologist Festinger 1957 (Puri and Tyler 1998). Festinger (1998) defines that 'cognitive dissonance which is a state of tension which arises when we realize that two or more of our cognitions are consistent with each other. People seek to reduce cognitive dissonance'. In this situation Kandyan royalty undergo tension if they find that their beliefs are different from their behavior. They tend to reduce the tension by completely changing their beliefs or their behavior .This process can be identified as a theory which strengthens or diminishes influencing fashion of dress. Although, Festinger describes that either behavior or belief changes, there are examples from the Kandyan era that do not purely confirmed to it. There are some instances, where people have changed their beliefs and behavior only half way.

Even though, king Vimala Dharma Suriya I, queen Dona Catherine, Prince Vijayapala and Rajasimha II had similar political, religious and social forces to follow the Western way of life, king Vimala Dharma Suriya I controlled his beliefs and behavior only half way in order to suit the needs of the society which he represented. In Spilbergens' travel records it is recorded that, "Dona Catherine visits no pagodas, Don Joan the king what he does is mostly to please the Sinhalese' (Ferguson 1927). Therefore his costume shows the hybrid formation of Eastern and Western styles. He rationalized his behavior.

The influence pattern which was derived through the analysis was Kandyan kings tend to change their beliefs or behavior when a mental tension arose. Often they selected one option to solve the conflict. Besides, they denied selecting either beliefs or behavior but tend to rationalize behavior then change belief and behavior half way. It was clearly proven that changing behavior or changing perception of behavior facilitate influencing fashion. The reinforced theory addresses the pervasive human tendency to rationalize, and gives three fundamental assumptions.

1. Humans are sensitive to inconsistencies between beliefs and behavior.
2. Recognition of this inconsistency will cause dissonance, and will motivate an individual to resolve the dissonance.
3. Dissonance will be resolved in one of three basic ways.
 - a. Change belief
 - b. Change behavior
 - c. Change perception of behavior and rationalize the behavior.

Conclusions and Recommendations

The influencing process of 'Cognitive dissonance' is able to change the attitude of fashion. The process is activated in three ways, exerted considerable influences. The key of this process is to build a mental tension between beliefs and behavior. As a result, people automatically tend to solve the conflict in between, consecutively are influenced by various fashion.

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EFFECT OF SILICON APPLICATION ON DOWNY MILDEW IN BITTER GOURD (*Momordica charantia* L.) LEAVES

R.M.R.N.K.Ratnayake^{1,42}, W.A.M. Daundasekera², H.M. Ariyaratne³
and M.Y.U. Ganehenegge⁴

¹Postgraduate Institute of Science, University of Peradeniya, Sri Lanka

²Department of Botany, Faculty of Science, University of Peradeniya, Sri Lanka

³Horticultural Crop Research and Development Institute, Gannoruwa, Sri Lanka

⁴Department of Chemistry, Faculty of Science, University of Peradeniya, Sri Lanka

Introduction

Silicon (Si) has been exploited for its beneficial effects in terms of disease control in many plants including wheat, rice, cucumber, cherry, muskmelon, and potato (Rodrigues et al. 2004). Two hypotheses for the Si-enhanced resistance to diseases have been proposed; i. Silicon deposited on the tissue surface acts as a physical barrier and makes the plant cells less susceptible to enzymatic degradation by fungal pathogens, ii. Silicon functions as a signal to induce the production of phytoalexin or expression of pathogenesis-induced host-defense responses in the infected plants. Bitter gourd (*Momordica charantia* L.; Family Cucurbitaceae) is a popular vegetable in south-east asia due to its unique taste and medicinal properties. However, bitter gourd plants are highly susceptible to many fungal diseases leading to loss of yield and product quality. Downy mildew caused by *Pseudoperonospora cubensis* is one of the major diseases in bitter gourd leaves. The very little published work related to silicon-related work on bitter gourd indicate that this crop is a moderate-silicon accumulator and Si application develops resistance in the plant against *Pythium aphanidermatum*, a root rot fungus. In this context, this study was conducted to determine the effectiveness of soil-amendment with Si on controlling downy mildew in bitter gourd leaves at field level and to determine if any cytological changes were associated with fungal restriction.

Materials and Methods

Bitter gourd (*Momordica charantia* L.) cv. Matale green was seeded on 20 cm diameter plastic pots with 18 L capacity filled with sand: top soil: compost at 2:1:1 ratio as the growing medium and they were arranged in a plant house at Department of Botany, Faculty of Science, University of Peradeniya according to complete randomized design (CRD) with 1m x 1.5 m plant spacing. Level of plant-available Si and pH in soil were assessed before amendment with silica. Potassium silicate (K_2SiO_3) (20 % SiO_2 , 8 % K_2O ; Daejung, Korea) in three concentrations; 100 mg Si kg^{-1} soil, 200 mg Si kg^{-1} soil, 300 mg Si kg^{-1} soil was added to the growing medium once a week

⁴² ruwankaratnayake@yahoo.com

starting from four leaf stage and continued up to 28 days. Potassium fertilizer was added to control pots as well at appropriate amounts to compensate the effect of added potassium in the form of K_2SiO_3 . Each treatment consisted with 15 pots. Fertilizer application, irrigation, staking, pruning and trellising were done according to Department of Agriculture (DOA) recommendations. Plants were allowed for natural infection since downy mildew is the most prevailing disease in bitter melon. Downy mildew was identified by disease symptoms and descriptions of International Mycological Institute. Disease severity was rated through external observations on weekly basis using a self-prepared scale (0=no disease, 1 = 1 to 20 %, 2 = 21 to 40 %, 3 = 41 to 60 %, 4 = 61 to 80 %, and 5 = 81 to 100 % of upper leaf surface covered with disease). Disease ratings were used to calculate area under disease progress curve (AUDPC) for each treatment by the midpoint rule method (Campbell and Madden 1990).

Number of leaves, flowers, and fruits per plant were counted in seven days interval beginning from four weeks after flowering. Silicon content was determined by a modification of the autoclave-induced digestion procedure of Elliot and Snyder (1991) at 5 weeks and 10 weeks after seedling emergence and the concentration of Si in leaf tissue was determined by a colorimetric analysis. Data were analyzed using ANOVA by SAS for windows version 9.0. Differences between treatment means were obtained by DMNRT at $p < 0.05$. Microtome sections of leaves were observed under light microscope and cuticle – epidermal layer thickness was measured to determine silicon deposition in leaves. Leaf sections were stained with silver amine chromate to observe any Silicon depositions in cells.

Results and Discussion

Average soluble Si concentration (plant available Si) in the soil suspension was 2.87 mg L^{-1} and Soil pH of the experimental field was 7.8. When pH levels are too high or too low, minerals are bound to soil particles and thus, unavailable to plants. Si presents in the soil solution as solubilized monosilicic acid $[Si(OH)_4]$ at a $pH < 9$, and as the silicic anion SiO_3^{2-} at a $pH > 9$ (Raven 2003). Therefore, soil pH Level of the experimental field was suitable for Si application.

Downy mildew symptoms were observed in the field at 7 weeks after seeding. Angular chlorotic lesions in the infected leaves, eventually, turned necrotic and curled upwards showing the typical symptoms of the disease. Further, Gray-brown to purplish-black fungal growth was observed at the lower surface of the leaf. Disease severity was increased with the time in all treated and control plants although it was lower in silicon treated plants than that of control plants during the entire period. Thus, area under

disease progress curve (AUDPC) was highly significant with treatments but there was no correlation between AUDPC and added Si content (Table 1). Among the three silicon treatments the highest Si accumulation in leaves (2.8 % of the leaf dry matter) was shown by 200 mg Si kg⁻¹ soil and also it showed a highly significant reduction ($p < 0.001$) in fungal growth and Si content in all control plants was less than 1 % (Table 1). However, accumulated silicon content in 300 mg kg⁻¹ soil was lower than that of the other two silicon treatments. This may be due to the pH of potassium silicate solution which was 10.5, thus addition of increasing amount of Si into the soil may increase soil pH hence Si absorption may be reduced. When pH levels are too high minerals are bound to soil particles and become unavailable to the plant (Raven 2003). Further, this may be the reason for the higher disease appearance linked with 300 mg Si kg⁻¹ soil compared to the other two treatments.

Si treatment had no effect on leaf flower or fruit number in plants (Table 2). Cuticle - epidermal layer thickness in Si treated plants ranged from 17.2 – 21.3 µm where in control plants it ranged from 10.3-13.81 µm. Further, it was noted that when bitter melon plants get diseased cuticle - epidermal layer thickness was decreased thus in the Si treated plants average thickness was 15.8 µm where in the control plants it was 10.2 µm.

Table 1. Effect of soil amendments with silicon in different concentrations on leaf silicon content and disease severity expressed as area under disease progress curve (AUDPC)

Si application rate (mg kg ⁻¹ soil)	Si content accumulated in leaves (% leaf dry matter) *	AUDPC*
100	1.9 b	50.05 c
200	2.8 a	43.45 d
300	1.7 c	61.62 b
0	<1.0 d	>75.6 a
P value	0.0001	0.0001

** Data based on observation of 70 days after seeding. Any two means in the same column followed by different letters differ significantly according to Duncan's multiple range test ($P < 0.05$).*

Note: AUDPC was calculated using disease severity data (on a 0 to 5 scale, where 0 = no disease symptoms and 5 = 81 to 100 % of upper leaf surface covered with disease symptoms of downy mildew).

According to the results, protective role of Si can at least be partly due to the accumulation of Si in the leaves, enhancing their strength and rigidity, which may create a physical barrier to pathogens as described by Rodrigues et al. (2004) and many researchers.

Conclusions and Recommendations

Soil amendments with silicon in bitter gourd plants reduced the severity of Downey mildew disease although it did not affect on growth stages. Cytological evidences suggest the physical barrier concept, however, further investigations are needed to explain the mechanism linked with Si application and enhanced disease resistance.

Table 2. Variation in number of leaves, flowers and fruits with different silicon treatments

Si level	Leaf Number * (no/plant)	Flower Number * (no/plant)	Fruit Number * (no/plant)
100ppm Si	77.80 ± 6.14	6.32 ± 0.32	4.65 ± 0.08
200ppm Si	82.61 ± 5.32	7.61 ± 0.85	5.12 ± 0.03
300ppm Si	80.61 ± 5.02	6.35 ± 1.32	4.48 ± 0.57
Control 1	81.52 ± 3.32	5.95 ± 2.45	4.52 ± 0.25
Control 2	80.61 ± 2.56	6.32 ± 1.32	4.80 ± 0.64
Control 3	78.61 ± 4.34	5.98 ± 1.58	5.30 ± 0.65

Note: * Observations made 70 days after seeding.

Control 1= No addition of silicon but K fertilizer was added to compensate the effect of added K in the form of potassium silicate in 100 mg Si kg⁻¹ soil and **Control 3=** No addition of silicon but K fertilizer was added to compensate the effect of added K in the form of potassium silicate in 300 mg Si kg⁻¹ soil

Acknowledgement

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APPLICATION OF WEAP MODEL FOR ASSESSING IRRIGATION WATER AVAILABILITY—CASE STUDY OF DEDURU OYA RESERVOIR

D.S. Sampath^{1,43}, S.B. Weerakoon², B.K. Mishra³ and S. Herath³

¹*Department of Civil Engineering, Faculty of Engineering, University of Jaffna, Ariviyal Nagar, Kilinochchi, Sri Lanka*

²*Department of Civil Engineering, Faculty of Engineering, University of Peradeniya, Sri Lanka*

³*Institute for Advance Study of Sustainability, United Nations University, Tokyo, Japan*

Introduction

Deduru Oya basin which has an area of 2620 km² is the sixth largest river basin in Sri Lanka. Rainfall in the basin has a temporal and spatial variation, and is only source of water. Deduru Oya carries flash floods during rainy season and very low flows during dry season. It releases about 1600 MCM of

water to the sea annually without being useful purposes. There are several anicuts across it to divert water for irrigation, but there is no single reservoir intercepting the Deduru Oya except the reservoir at Thunmodara which is being constructed under Deduru Oya project (Fig. 1). There is a strong need to regulate Deduru Oya flow for its optimum use especially for irrigation during dry season.

The Deduru Oya reservoir project which is under construction is aimed to improve the livelihood of farmers in parts of the North-Western Province by increasing the productivity of its land and water resources by regulating and diverting the water by two main canals. The proposed Left Bank (LB) canal which is 44.1 km long, will supply water to augment about 136 existing storage-based minor irrigation systems in the left bank of the Deduru Oya (Fig. 1). The Right bank (RB) canal is a trans-basin canal to augment water supply to Iginimitiya reservoir which is located in Mee Oya basin (Fig. 1). The proposed irrigable area under LB canal including new developments is about

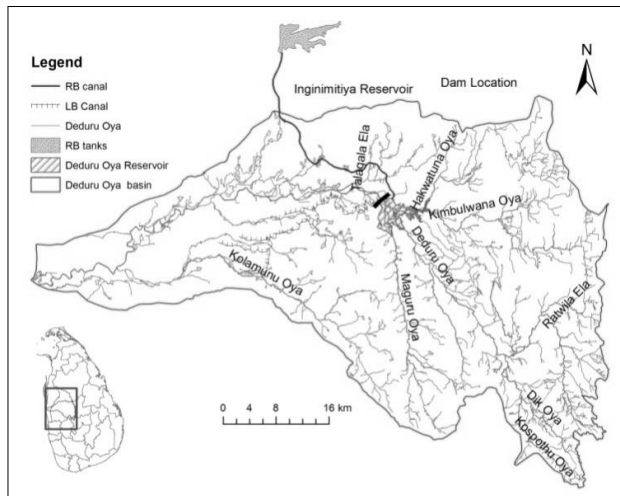


Figure 2. Study area

⁴³ pe.saliya@gmail.com

3000 ha. Total irrigable area under Deduru Oya basin is 7000 ha and under Mee Oya basin is 4115 ha (Pre-feasibility Report 2000).

This study is focused on the setting up WEAP21 version 3.43 (Water Evaluation And Planning) model to study the impact of annual change in river flow on irrigation water availability under different water management scenarios with the reservoir project. They are the operation of LB canal with the small tanks and without small tanks.

Materials and Methods

Rainfall or runoff data are not available for the rain-fed small tanks in the minor irrigation system of the LB region. Hydrologic Engineering Center – Hydrologic Modeling System (HEC-HMS) version 3.0.1 developed by US Army Corps of Engineers in USA was used to develop rainfall runoff model for each of the catchment. HEC-HMS model was calibrated and verified for the Tittawella Tank in Kurunegala (Sampath et al. 2014a). Calibrated HEC-HMS model for the Tittawella Tank and rainfall data at Nikaweratiya and Ridi Bendi Ella stations were used to develop inflows to the minor tanks. Also HEC-HMS model was used to simulate flow in the Deduru Oya. Thirty years daily rainfall data from 6 rain gauge stations in the Deduru Oya basin and runoff data at Moragaswewa from 1984 to 1989 together with monthly evaporation data at Batalagoda agro-meteorological station were used in the simulation. Diversions, reservoir storages and losses were also accounted in the study (Sampath et al. 2014b). Calibrated HEC-HMS model of Deduru Oya and rainfall data at Millawa, Kurunegala, Ridibendiella and Wariyapola stations were used to develop inflows to Deduru Oya reservoir.

For each of the 136 rain-fed small tanks that will be supplied water by the LB canal, the relevant catchment areas, storage areas, natural streams, land use patterns and cascades were identified for modeling the system. Also 12 directly feeding demand sites were identified. Topography, geology and land use details were collected from the digital data of the Survey Department of Sri Lanka and Arc GIS 9.2 was used as a tool for spatial analysis.

For WEAP model application, four types of data viz: Area cultivated annually, Annual water use rate, Monthly variation and Consumption are required at each of the demand sites. Crop water requirement was calculated assuming 105 day low land paddy will be cultivated in an area of 3000 ha under LB canal. Crop water requirement was calculated on monthly basis. Rainfall data at Nikaweratiya and Ridi Bendi Ella station in year 2000 to 2010, Mahailluppallama reference crop evapotranspiration rates and crop factors for each growth stages were used for the calculation of crop water requirements. Computations of irrigation water requirements were made using 60 % application efficiency and 75 % conveyance efficiency. Land

soaking and tiling requirement were also taken into account (Pre-feasibility Report 2000). Calculated crop water requirements and irrigable area were used to develop demand site 'Annual water use rate' and 'Monthly variation' of the WEAP model. Different 'Supply preference' and 'Priorities' were used to model the diversion link and transmission links.

In WEAP model, two Flow Requirement elements were used to model RB trans-basin diversion and downstream environmental and irrigation requirements. Annual RB trans-basin diversion water requirement was fixed as 90 MCM. The minimum value of environmental flow requirements were used as $3.2 \text{ m}^3\text{s}^{-1}$ (Pre-feasibility Report 2000). Total downstream water requirement was calculated by considering both Ridi Bendi Ella scheme and mandatory releases for downstream irrigation flow requirement of Deduru Oya.

Results and Discussion

WEAP Model results provide the reservoir storage, possible releases to LB canal and to RB canal in a given month during the simulation period against the inflow to the reservoir, downstream mandatory release and actual demand for LB canal and RB canal during the month. Analysis showed that Year 2005 was the year with the greatest water scarcity during the period of simulation from 2001 to 2010.

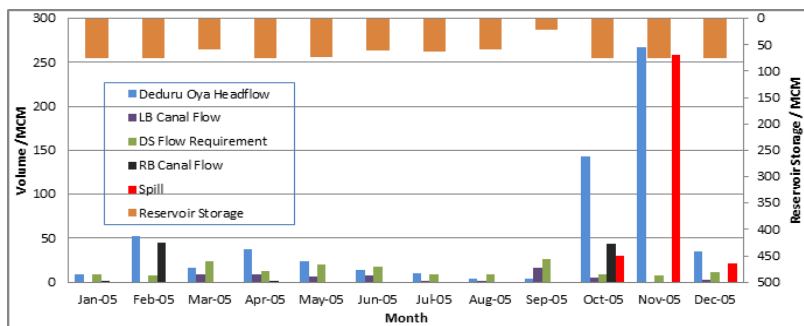


Figure 3. Water Diversion in 2005 with tank scenario

Figure 2 shows the different diversion quantities for the different water needs from the reservoir in year 2005 for the scenario where the small tanks are also in operation in the LB canal system to receive flow from their own catchments and the LB canal. The Deduru Oya reservoir project is able to satisfy all the water demands for different water needs in the area without failure even during year 2005 which was the driest year during the selected period. Figure 3 shows the different diversion quantities for the different water needs from the reservoir in year 2005 for the other scenario where small tanks are abandoned and not used for water management in the LB region. The simulation shows that the Deduru Oya reservoir satisfies all the

water demands for different water needs in all months of the simulation period but it fails in September 2005. This is depicted by the plot in Figure 3. This indicates that the reservoir will fail in some months even in the future if the existing small tanks are not incorporated in to the water management of the LB canal irrigation system. The model developed is a useful tool for planning of water issues in the reservoir operation to optimize the water use in the Deduru Oya reservoir. This shows the importance of incorporating existing distributed small irrigation tanks to the modern irrigation development to improve the resilience of the system.

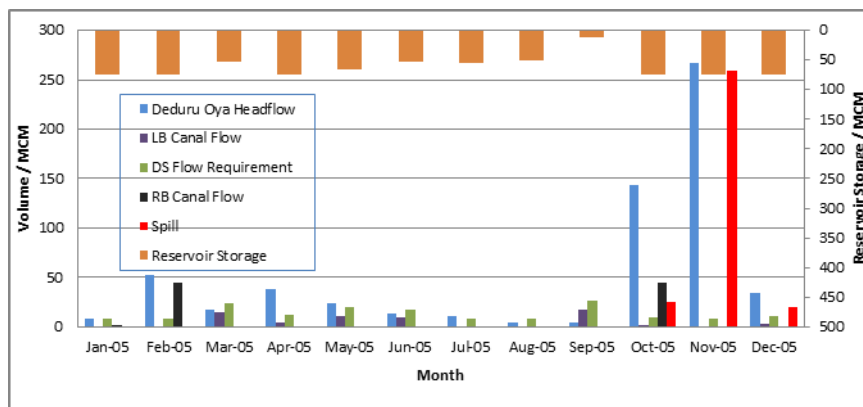


Figure 4. Water diversions in 2005 without tank scenario

Conclusions and Recommendations

The simulation carried out for past ten years reveal that the Deduru Oya reservoir project which has planned to operate LB canal irrigation management incorporating the existing small irrigation tanks will be able to supply the water demand for LB development area for paddy cultivation without failure. The study emphasizes the importance of incorporating existing distributed small irrigation tanks to the modern irrigation development. The irrigation water management model is a useful tool for planning of water management in the Deduru Oya reservoir project

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EVALUATION OF THE ANTAGONISTIC NATURE OF EXTRACELLULAR COMPOUNDS SECRETED BY SOME SELECTED BACTERIAL ANTAGONISTS AGAINST *Colletotrichum truncatum*, THE CAUSAL AGENT OF ANTHRACNOSE DISEASE IN CHILLI

H.B.P. Sandani¹, N.P. Ranathunge^{2,44} and P.L.N. Lakshman³

¹*Board of Study of Faculty of Agriculture, Faculty of Graduate Studies*

²*Department of Agricultural Biology, Faculty of Agriculture*

³*Department of Food Science and Technology, Faculty of Agriculture
University of Ruhuna, Mapalana, Kamburupitiya, Sri Lanka*

Introduction

Chilli, fruit is popular for its spicy nature, in both fresh and dried forms. According to Pakdeevaporn et al. (2005), anthracnose is a threat which causes up to 50 % loss in chilli yield. *Colletotrichum truncatum* is one of the most destructive agents of anthracnose disease. It is highly worth to pay attention on biological control means as a reliable alternative in controlling anthracnose when considering the adverse effects of fungicides. The use of microbial antagonists as an alternative is becoming popular. However, one of the possible exceptions is to utilize the extracellular compounds secreted by such antagonists that can effectively control the pathogens. As this is a long term process that leads to separation and identification of responsible compounds, the primary step is to identify the antagonistic properties of the extracellular metabolites of potential candidates. The objective of this study was to evaluate the nature of antifungal compounds secreted by five selected bacterial antagonists against *C. truncatum* causing anthracnose disease in chilli.

Materials and Methods

Production of diffusible antifungal substances: The diffusible nature of antifungal compounds secreted by antagonists was evaluated using the cellophane overlay technique (Nourozian et al. 2006). The results were expressed as percent inhibition of radial growth (PIRG) values according to Sariah (1994) compared to control.

$$\text{PIRG} = (R1-R2)/R1 \times 100$$

R1- Radial growth of *C. truncatum* in control plate; R2- Radial growth of *C. truncatum* interacting with antagonistic bacteria. Experiment was a randomized design with three replicates and repeated once for better accuracy.

Production of volatile antifungal substances: Volatile nature of antifungal compounds was evaluated according to agar strip removal technique

⁴⁴ nalika@agbio.ruh.ac.lk

described by Choi et al. (2006). Experiment was repeated once in completely randomized design with three replicates. The results were expressed as PIRG values compared to a control.

Mycelial growth test: First, small agar plugs of seven days old *C. truncatum* was dipped in each overnight antagonistic bacterial suspension (10^8 CFU mL⁻¹) for thirty minutes in sterile test tubes. The treated *C. truncatum* mycelial plugs were air dried in laminar air flow and transferred onto fresh PDA media separately. Fungal plugs dipped in sterile broth medium were used as the control. Plates were incubated at 28 °C for seven days. Then crude supernatants of overnight nutrient broth cultures of each bacterial antagonist were incorporated with PDA in 2:3 (V/V) ratio and 5 mm size mycelial blocks of *C. truncatum* was introduced into those media and the radial growth of the fungus was measured.

Spore germination test: A volume of 100 µL of *C. truncatum* spore suspension (10^6 spores mL⁻¹) was spread over PDA medium using a bent glass rod. Two sterilized filter paper discs (0.5 cm diameter) were placed 3 cm apart from each other on PDA. A volume of 50 µL of each bacterial suspension was pipetted onto filter papers in each plate. Filter paper discs containing 50 µL sterilized nutrient broth served as the control. After 24 hours of incubation at 28 °C, the diameter of the inhibition zones around filter paper discs were measured.

All the collected data was analyzed using Dunnet's test in SAS software.

Results and Discussion

Volatile and diffusible antifungal substances: According to the results obtained, antifungal substances produced by selected antagonists were diffusible in nature. As shown in Table 1, the mycelial growth of *C. truncatum* has reduced significantly ($p < 0.05$) in the cellophane overlay technique. Except C31 isolate, all the other antagonists have been able to completely inhibit the radial growth of *C. truncatum* through their diffusible antifungal substances. As there weren't any significant inhibition in radial growth of *C. truncatum* at significant level in agar strip removal technique, the antifungal substances secreted by the antagonists can be considered as non volatile (Table 1). As C31 antagonist doesn't produce either diffusible or volatile antifungal substances, its bio controlling mechanism may not be antibiosis.

In both experiments of mycelial growth test, inhibition of radial growth of *C. truncatum* by all the selected antagonists was 100 % compared to the control.

Table 1. Growth inhibition of *C. truncatum* by diffusible and volatile antibiotics produced by antagonistic bacteria at seven days after incubation

Antagonist	Inhibition of radial growth (%)	
	Diffusible antibiotics	Volatile antibiotics
F2	100	0
F35	100	0
F65	100	0
F79	100	0
F80	100	0
C31	12 *	0
Control	0	0

Note: * PIRG isn't significant at $p < 0.05$ level.

Mycelial plugs of *C. truncatum* have lost their viability due to the effect of antagonistic bacteria when dipped in an overnight culture. In the PDA incorporation method where the media composition was antagonists' supernatant: PDA in 2:3 ratio didn't allow the growth of *C. truncatum*. As shown in Table 2, all the antagonists significantly inhibited the mycelial growth of *C. truncatum* at $p < 0.05$ level.

Table 2. Effect of antagonistic bacteria on the growth of *C. truncatum* after seven days of incubation

Antagonist	Inhibition of radial growth (%)	
	Dipping method	Incorporation method
F2	100	100
F35	100	100
F65	100	100
F79	100	100
F80	100	100
C31	80	100
Control	0	0

Note: *all Inhibition of PIRG as significant at $p < 0.05$ level

In spore germination test 50 μ L of each antagonist which were added to filter paper discs inhibits spore germination significantly giving a clear zone around each disc. But C31 antagonist didn't show any inhibition on spore germination on the spore lawn. The radius of each inhibition zone is indicated in Fig. 1.

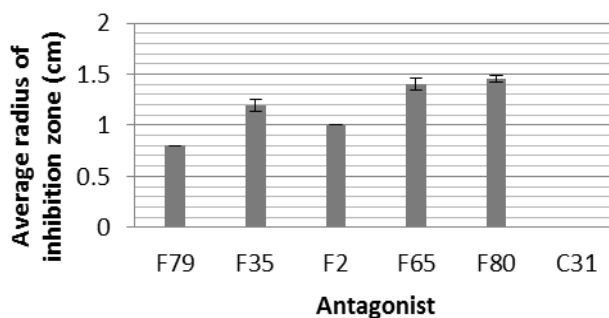


Figure 1. Radius of the inhibition zone on *C. truncatum* spore lawn by each antagonist

Analysis of all the selected antagonists showed a great antagonism against *C. truncatum* in many ways. It is clear that mycelial growth and spore germination of *C. truncatum* are strictly controlled through extracellular antifungal substances F2, F35, F65, F79, F80 antagonists. Rahman et al. (2007), have also stated the antagonistic effect of *Burkholderia cepacia* and *Pseudomonas aeruginosa* against *Colletotrichum gloeosporioides* as discussed above. However C31 antagonist deviates from others, probably because C31 displays fungistatic properties by competition rather than antibiosis. Further studies will be required to confirm this assumption.

Conclusions and Recommendations

F2, F35, F65, F79, F80 antagonists produce extracellular antifungal compounds which strongly inhibit the mycelial growth and spore germination of *C. truncatum*.

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LAND SUITABILITY EVALUATION FOR RUBBER CULTIVATION USING MULTI CRITERIA DECISION APPROACH AND GIS: SPECIAL FOCUS ON THE AMPARA DISTRICT

J.K.S. Sankalpa¹⁴⁵, S. Karunaratne² and W. Wijesuriya¹

¹ *Rubber Research Institute of Sri Lanka*

² *Faculty of Agriculture & Plantation Management, Wayamba University of Sri Lanka, Makandura, Gonawila(NWP), Sri Lanka*

Introduction

A considerable extent of rubber lands in the Wet Zone has been converted to other crops due to unstable price and high cost of production in the rubber sector. Conversion of rubber lands into industrial sites and dwelling units was also common in these areas. Hence, to meet the production targets in the rubber sector, the Government has taken an initiative to expand rubber cultivation into drier areas due to low availability of lands for further expansion in traditional areas. These areas include; Ampara, Anuradhapura, Vavuniya, Kilinochchi and Mullaitivu districts (Anon 2011a). Among these areas, the most extensive and rapid change in expansion of rubber farming occurred in the Eastern province of Sri Lanka. In order to perform in an optimal manner, rubber plantations require certain climatic and physiographical characteristics as listed in the handbook of rubber (Anon 2011b). Hence, it is a basic necessity to identify areas which are suitable for rubber cultivation based on the requirements to reap the maximum benefits. On the other hand, areas with limiting characteristics also should be identified to propose recommendations to minimize adverse impacts. In this regard, crop-land suitability analysis is a most important step in achieving optimum utilization of the available land resources.

In this paper we examine the resource quality and accessibility for rubber farming in Ampara district of Sri Lanka using multi criteria and GIS approach of land suitability evaluation. The aim of this multi criteria evaluation with Geographical information systems (GIS) is to provide more flexible and superior mechanism to the decision makers for efficient planning and management in rubber farming in Ampara district.

Materials and Methods

Study Area and Data used: The study area is bounded within 80° 58 and 81° 52 E longitudes and 7° 44 to 6° 30 latitudes. Different types of data sets were used for rubber land suitability analysis. Various physical properties of the land (Soil and topography) and ecological parameters (temperature and

⁴⁵ ssankalpa2@gmail.com

rainfall) of the area were used for this purpose. Economic suitability is measured using accessibility to the market *via* main roads. Soil texture map provided by European commission joint research center and 1:50,000 topographical maps provided by DIVA.org free GIS data sources were used for the study. Temperature and rainfall maps used were developed by Natural Resources Management Centre (NRMC).

Preparation of topographic database: USGS SRTM DEM (90 m resolution) was used to produce the slope map for the study area. Slope values were divided into three classes and rating for land suitability was done as proposed by Karunaratne et al. (2011).

Preparation of soil database: Physical properties of soils were considered in the analysis. There were 9 soil series in the soil map and was prepared by digitizing the soil texture map of Ampara district. Rating was done for the soil classes according to Karunaratne et al. (2011).

Climate data: Temperature and rainfall are the two climatic factors which affect the performance of rubber cultivations. Ratings for these variables were assigned based on the requirements for rubber planting (Anon 2001b) and as proposed by Karunaratne et al. (2011).

Market access data base: In this research, in addition to the variables considered by Karunaratne et al. (2011), Market access was determined by the distance from major roads in the district. Euclidean distance road map was prepared using GIS spatial analyst tool. The accessibility attribute also divide into three levels-good, moderate and poor based on distance from road. Obviously the time spent in transporting produce to the roadside increases as accessibility declines due to the greater distance and poorer conditions of the roads

Land suitability analysis: Arc GIS 10 software was used for preparing the slope map, temperature maps, mean rainfall map, soil series map and the Euclidean distance map. Those maps were combined as inputs for the land suitability model according to the equation given below. Spatial multi criteria suitability analysis is a process where geographical data is combined and transformed into a decision. Analytical Hierarchical process (AHP) was used to evaluate the priority weights of each factor (Satya 2008). A final value of the weight for each criterion was decided compared with available literature.

Table 1. Weight for the model parameters

Parameter	Weight (%)
Rainfall	55
Maximum and minimum temperature	2
Slope	15
Elevation	6
Soil type	15
Market access	7

Suitability raster model for cultivation,

$$S = \sum_{i=1}^n w_i c_i \prod_{j=1}^m r_j$$

S = suitability for cultivation of rubber, w_i = weight

for criteria i , c_i = criteria for suitability and r_j = restriction

Results and Discussion

All inputs were used in index model according to the suitability criteria. Maps were prepared after reclassifying all attributes for each variable. The proposed suitability map for rubber cultivation in the Ampara district and generated criteria are depicted in Fig. 1.

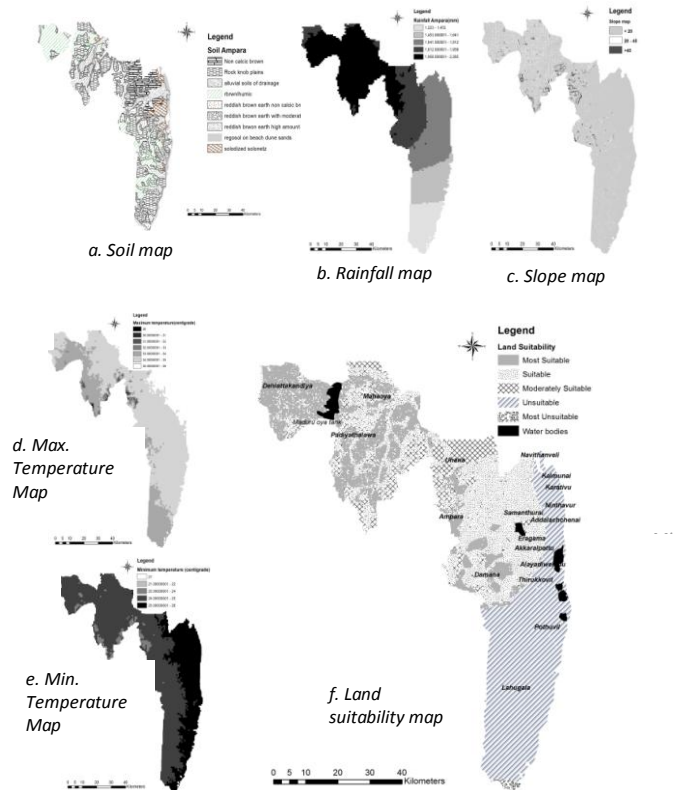


Figure 1. Main input parameters and generated Land suitability map.

The area which is identified as most suitable can be effectively used for rubber cultivation. Areas having moderate suitability should be given a package of conservation practices to control moisture and nutrient while increasing the market accessibility. Areas having Poor resource quality should be given more attention when converting them into rubber cultivation. Validation of the derived land suitability model was done using

available Land Use map for the study area (year 1981). Most of the suitable land areas were overlaid with high dense to low dense vegetation cover.

Conclusions and Recommendations

Areas falling into most suitable classes can be effectively used for rubber cultivation with minimum conservation and technical effort which is effective for minimizing the cost of production in rubber farming. Percentages of land extent under most suitable, suitable and moderate suitable are 18 %, 35 % and 10 % respectively. Suitable land for cultivation rubber under most suitable, suitable and moderate suitable land areas are 75,996 ha, 147,770 ha and 44,331 ha respectively. The outcomes of this research can be effectively used for better decision making in ground planning and for further expansion schemes for rubber expansion within the district. The analysis could be refined further by incorporating more suitability indicators such as soil chemical properties and present land use condition in the district.

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EFFECTS OF BIOCHAR PRODUCED FROM DIFFERENT FEED STOCK MATERIALS ON SOIL PROPERTIES IN SANDY REGOSOL OF JAFFNA, SRI LANKA

S. Senthuran¹⁴⁶ and N. Gnanavelrajah¹

¹Department of Agricultural Chemistry, Faculty of Agriculture, University of Jaffna

Introduction

Soil fertility plays a vital role in agricultural productivity. Sri Lanka's sandy regosol is one of the major soil groups in dry zone found along or near the coastline of the Jaffna Peninsula covering 25 % of land area. Low nutrient retention and high hydraulic conductivity of sandy regosol soils contribute to the large amount of water and nutrients to be lost beyond the rooting zone of plants and is low in agricultural value. Biochar does increase crop productivity through an improved water holding capacity of the soil, along with improved crop nutrient availability, especially in sandy soils. Biochar is the charred by-product of biomass pyrolysis, the heating of plant-derived material in the absence of oxygen in order to capture combustible gases. Therefore application of biochar could be a solution to low productivity of sandy regosols. With this background, a field experiment was conducted to study the effect of biochar produced from different feed stocks namely paddy husk, coconut nut shell and palmyrah nut shell with combination of departmental recommended fertilizer (DRF) on growth and yield of brinjal and soil properties. Hence objective of this study was to find the effects of three different biochar on soil properties.

Materials and Methods

Study area: The study area is a low country dry zone located in the Vidaththatpallai located in Thenmaradchi DS Division, Northern Province of Sri Lanka. The soil belongs to sandy regosol.

Charred biomass production: Different biomass namely paddy husk, coconut shells and palmyrah nut Shells were charred using two drum method. In this method one small drum filled with biomass was kept inside a big drum so as to put firewood in between the available space and firewood was lit. Charred biomass was broken separately into powder form, sieved by 2 mm and 4 mm sieves and both were mixed at 1:1 ratio.

Field experiment: Field experiment was designed with four treatments (Table 1) arranged in a randomized complete block design (RCBD). Details about fertilizer application are given in Table 2. Each treatment was replicated three times. One week after land preparation char was applied at the rate of 10 t ha⁻¹. Char was mixed well with soil at a depth of 15 cm and a

⁴⁶ senthu1000@gmail.com

diameter of 60 cm around each planting hole. Twenty one day old brinjal seedlings were transplanted after two weeks of char incorporation. Irrigation was done manually using buckets made up of palmyrah leaf which is practiced in the sandy soil villages. After the transplantation irrigation was done daily until twenty days and thereafter once in two days.

Table 1. Treatments application (DOA)

Treatment No	Experiment procedure
T ₁	DRF
T ₂	Paddy husk char + DRF
T ₃	Coconut shells char + DRF
T ₄	Palmyrah nut shells

Table 2. Fertilizers

	DRF-Basal : 7.07 kg (15:65:17)	DRF-1 st top dressing
Urea (N)	1.09 kg	1.09 kg
TSP (P)	4.74 kg	
MOP (N)	1.24 kg	

Note: DRF-Department of Agriculture Recommended Fertilizer

Analysis of soil sample collection and preparation: Sixty days after applying char soil samples were collected from each plot (12 plots) in the experimental field. From each treatment plot three plants were randomly selected and samples were collected within 30 cm radius from plant and up to 15 cm depth. Then collected samples were air dried, crushed and sieved to pass through 2 mm sieve. Available phosphorus, potassium, nitrogen (ammonium and nitrate), CEC, organic matter content, and soil moisture content were measured. Results were analyzed by SAS (9.1) package and the mean separation was done by Duncan multiple range test at p=0.05

Results and Discussion

Results showed that biochar application increases the N availability than the fertilizer application alone (Fig. 1.1). Ding et al. (2010) have reported that application of biochar reduced overall cumulative losses of NH_4^+ -N via leaching which could have been due to increased ammonium adsorption by biochar treated soils. Available P of initial soil was 17.91 mg kg⁻¹. All treatments had higher P availability compared to initial soil.

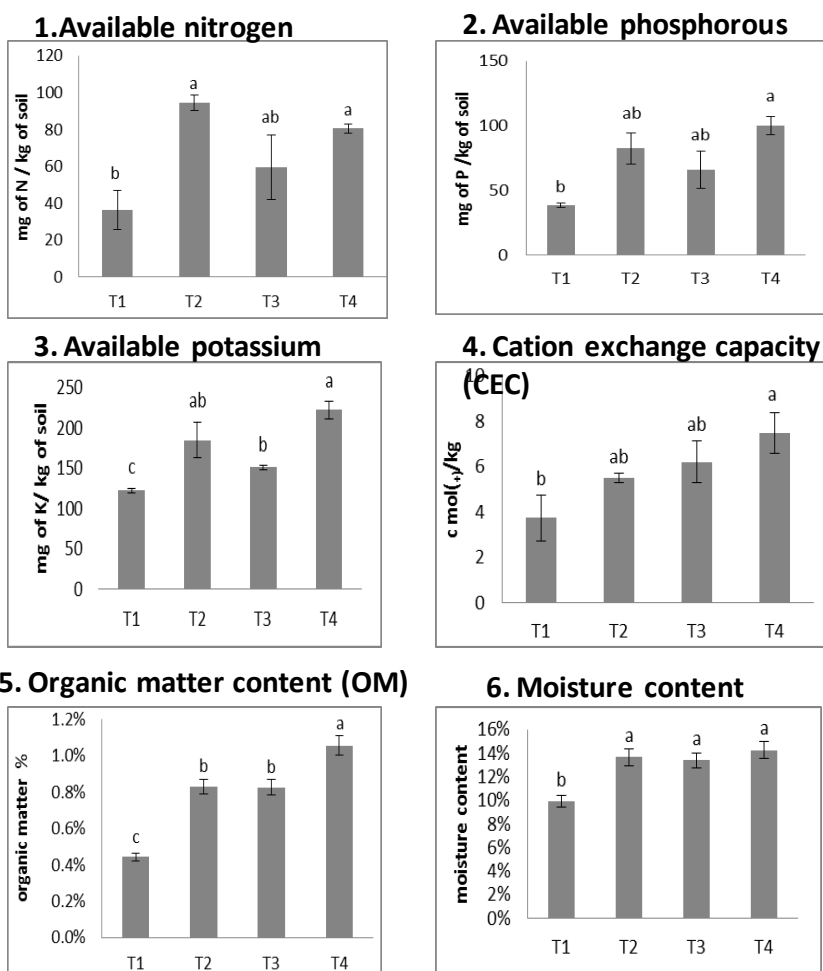


Figure 1. Effect of different treatments on N availability, P availability, K availability, CEC OM and moisture contents with 40 DAP soil. The columns marked with same lowercase letters do not differ significantly ($p < 0.05$). Error bars, $n=3$).

Note: T_1 - DRF

T_2 - Paddy husk char + DRF

T_3 - Coconut shells char + DRF
+ DRF.

T_4 - Palmyrah nut shells char

T_4 had the highest significant available P compared to all other treatments (Fig. 1.2). Lehmann et al. (2003) has reported that the immediate beneficial effects of bio-char additions for nutrient availability are largely due to the availability of higher K and P. T_4 (palmyrah nut shells char + DRF) had the significantly highest available K compared to all other treatments except T_2 (paddy husk char + DRF) (Fig. 1.3).

All char treatments had higher CEC compared to without char treatment. T_4 had the highest CEC, while T_1 had the lowest CEC compared to all other

treatments (Fig. 1.4). Biochar has high surface area which increases soil CEC. T₄ gave the highest organic matter compared to all other treatments and T₁ had the lowest (Fig. 1.5). Lehmann (2007) has also reported that high organic carbon was in soils treated with biochar. Moisture content (Fig.1.6) was significantly increased in char added treatments (T₂, T₃ and T₄) compare to without char treatment (T₁). Increased moisture content could be attributed due to increased water holding capacity by char.

Conclusions and Recommendations

Palmyrah nut shell char significantly increased available N, P, K, CEC, organic matter content, and moisture content compared to without char treatment. Though coconut shell char and paddy husk char also improved these properties, significant increases were observed only in available K, CEC, organic matter content and moisture content.

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**ACCEPTANCE OF A CLINICAL PHARMACY SERVICE BY OTHER
MEMBERS IN THE HEALTHCARE TEAM: EXPERIENCE IN A TERTIARY
CARE HOSPITAL IN SRI LANKA**

**L.G.T. Shanika^{1,47}, S. Jayamanne^{1,2}, N. Wijekoon², J. Coombes¹, I. Coombes¹,
D. Perera¹, N. Mamunuwa¹, V. Pathiraja¹, H. A. De Silva² and A. Dawson¹**

¹*South Asian Clinical Toxicology Research Collaboration*

²*Faculty of Medicine, University of Kelaniya*

Introduction

Clinical pharmacists are part of a multidisciplinary team whom intends to provide a patient-oriented pharmaceutical care to enhance the quality use of medicines (QUM) of patients. Clinical pharmacist takes an accurate medication history of patients' pre-admission drugs and reviews it for safety, appropriateness and efficacy and provides medicine information to health professionals and patients. Ward based clinical pharmacists interact with other healthcare professionals as a source of drug information and educate them when necessary.

Many studies done in different countries in the world have shown the benefits received by patients when a clinical pharmacy service is incorporated to the multi-disciplinary healthcare team. In Sri Lanka, the pharmacists' direct involvement in patient management is limited and their duty is restricted to drug dispensing. The concept of clinical pharmacy is relatively new to the current healthcare system in Sri Lanka. In Sri Lanka, no research has been previously carried out to evaluate the impact of a ward based clinical pharmacy service in optimizing QUM and the perspectives of the healthcare team on this service.

The objectives of this study were to evaluate the rate of acceptance of the clinical pharmacist's recommendations regarding Drug Related Problems (DRPs) by the healthcare team, to determine the quality and quantity of drug information queries directed to the clinical pharmacist and to assess the level of acceptance of the clinical pharmacist's service by the other members of the ward staff.

Materials and Methods

This is a part of a large controlled trial conducted to assess the effectiveness of a clinical pharmacy service in patient management for the first time in Sri Lanka. It was done over a seven month period, in the Professorial Medical Unit of a tertiary care hospital. Patients with defined chronic non-

⁴⁷ thushani85@gmail.com

communicable diseases who needed long term medications were included in the trial. The control group received current standard care, while the intervention group received a clinical pharmacist's service in addition to the standard care. DRPs were classified according to the adapted PCNE classification system V5.01. Through prospective (intervention group) or retrospective (control group) medication review the pharmacists identified the DRPs. In the intervention group the clinical pharmacist discussed the hospital related DRPs with the healthcare team (doctors and nurses).

The quality and quantity of medicine - related questions directed to the clinical pharmacist by healthcare staff during the study period were also recorded. A staff survey was carried out at baseline before introducing the clinical pharmacy service and at the end of the study to determine the views of doctors and nurses regarding their perceptions of the role of clinical pharmacist as a member of the healthcare team. The survey was conducted through a pre-tested self-administered questionnaire. The questionnaire was peer reviewed prior to the survey. The questionnaire was composed with 9 closed-ended questions. Final question was about assessing the attitudes of staff on specific tasks of the customized clinical pharmacy service in Sri Lanka. An individual consent form was attached with the questionnaire and it clearly explained the purpose of this survey and the participants' right to grant or withhold the consent voluntarily. Moreover the investigator verbally explained all the information before distributing the questionnaire.

Ethics approval for the study was obtained from the Ethics Review Committee of the Faculty of Medicine, University of Kelaniya. The data were input into SPSS, V.21. Descriptive statistics were used to interpret the results. P values less than 0.05 ($P > 0.05$) were considered to be statistically significant.

Results and Discussion

In the intervention group a total of 270 (268 doctors and 2 nurses) recommendations regarding DRPs were directed to the healthcare team. Eighty three percent [221/268 (95% CI=77% - 87%; $P < 0.001$)] of the recommendations were accepted by the doctors. Previous acceptance rates reported in the literature range between 69% - 90%. The rate of implementation of the pharmacist's recommendations was 74.3% [199/268 (95%CI=69% - 79%; $P < 0.001$)] consistent with previous studies. A few DRPs (69/268) were not resolved even though discussed with prescribers. A majority happened when interventions were directed to intern house officers (58/69). This may be due to lack of experience of intern house officers as they started the carrier recently. Prescribers identified some DRPs were not significant to take an action. Almost all pharmacist

recommendations on drug-drug interactions (3/3) and prescribing errors (17/17) were accepted by doctors. A large number of recommendations were suggested on untreated indications or missing drugs from prescription (159/268). Out of 159 interventions, 116 were acted upon by doctors [73.0% (95%CI=65% - 80%; $P<0.001$)]. Two DRPs about missing the drugs on drug chart when recording were discussed with nursing staff were resolved completely. We believe that the development of a good inter-professional relationship with mutual trust greatly contributed to the high rate of acceptance.

A total of 17 medication-related questions were directed to the clinical pharmacist during the study period. Nine were from doctors, 4 were from nurses and 4 were from the medical students. The majority of questions from doctors and medical students were related to the doses, indications and available generic substitutions in the market. The nature of the questions directed from nurses' was using the pharmacist as a source of information about indication, side effects and generic name of the drugs.

At the end of the study period the rate of the response to the questionnaire from doctors was higher compared to the baseline survey; 67% (8/12) at baseline and 92.3% (12/13) at the end of the study. The attitudes of the medical staff regarding incorporation of clinical pharmacy service to the current healthcare system were satisfactory at the baseline period.

At the end of study, 100% of consultant physicians, 100% of senior registrars, 100% of registrars and 80% of house officers admitted that the incorporation of a clinical pharmacist to the existing health care system would be useful and this collaboration could improve the current standards especially in the public sector hospitals. Ninety-two percent of medical staff would be happy to receive the services from a competent clinical pharmacist. Similarly, Gillespie, et al. (2012) reported that 95% of physicians were satisfied with the incorporation of a new clinical pharmacy service to University Hospital in Uppsala, Sweden. Sixty-seven percent (95%CI=35% - 90%; $P>0.05$) of medical staff accepted that the pharmacist could advise medical staff and nurses regarding the issues related to medications. One (8%) prescriber disagreed with that statement and 25% didn't express any view. Majority [91.7% (95%CI=61.5% - 99.8%; $P=0.006$)] of medical staff agreed that adding a pharmacist to the team isn't a waste of money and is necessary to the current healthcare system. Sixty-seven percent of prescribers agreed that pharmacists can play an important role in tailoring drug therapy for individual patients. Ninety-two percent of medical staff acknowledged that pharmacists can play an important role in improving patients' medication taking behaviour. These data highlight that the clinical

pharmacy service has potential to be well accepted and utilized by the medical staff in Sri Lanka.

The survey response rate of nursing staff was 80% at baseline. However the nursing staff did not consent to participate in the questionnaire after the study. The perspectives of nursing staff at baseline (before introducing the clinical pharmacy service to the ward) was negative; 58.3% (95%CI=27.7% - 84.8%; $P>0.05$) stated that there is no need of clinical pharmacy service to the ward and 67% (95%CI=35% - 90%; $P>0.05$) weren't happy to welcome this service. Although a wide proportion of doctors acknowledged the importance of clinical pharmacy service, the response received from the nursing staff was negative in the baseline period. Lack of awareness and knowledge about the importance of this service could be a likely reason for this negative response. Moreover, there were fewer opportunities to interact with nurses regarding patients' drug management and this also could have contributed to the results of the survey. This data emphasize that there is a need to build trust and rapport with nursing staff in order to promote a clinical pharmacy service in Sri Lankan hospitals.

Conclusions and Recommendations

There was high acceptance and implementation rate of clinical pharmacist's recommendations regarding DRPs by the medical team. The medical practitioners were satisfied with the inclusion of a ward based pharmacist to the healthcare team. Doctors recognized that this collaboration improves the rate of resolution of DRPs which in turn improves QUM in patients. However there is a need to improve rapport and relationships between the clinical pharmacist and nursing staff with regards to patient drug management.

Acknowledgement

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FOREIGN DIRECT INVESTMENT AS A CATALYST FOR ECONOMIC GROWTH IN SRI LANKA

P. Sivashankar¹⁴⁸

¹*Sabaragamuwa University of Sri Lanka, Beliholoya, Sri Lanka*

Introduction

Foreign direct investment (FDI), "refers to an investment made to acquire lasting or long-term interest in enterprises operating outside of the economy of the investor." The investment is direct because the investor, which could be a foreign person, company or group of entities, is seeking to control, manage, or have significant influence over the foreign enterprise. It involves participation in management, joint venture, transfer of technology and expertise. Foreign investment can be a significant factor for development of less developed countries. It provides an inflow of capital and funds. Also it contributes to economic growth through an increase in productivity by providing better technologies and managerial skills to the host countries. The relationship between FDI and economic growth has long been a subject of great interest in the field of international development. Developing and transition economies respectively accounted for 45 per cent and 6 per cent of global FDI.

FDI is often seen as an important catalyst for economic growth. Empirical studies on the FDI-growth relationship have reported conflicting results. Some studies have found that FDI exerts a positive growth effect on the recipient countries, while others have discovered no such evidence or even a negative effect on growth. Few studies have been conducted on FDI and growth with relevance to developing economies. Yet, the results reveal that the conclusions are country specific and do not support generalization. Given this context, this study investigates the role of FDI in Sri Lankan economic growth. Thus, the objective of the research is to examine the relationship and the direction of causality between FDI and economic growth in Sri Lanka. Trade openness is crucial for acquiring the potential growth impact of FDI. FDI has no impact on long run growth after controlling for the country specific differences in some studies. Although several studies on FDI and growth in developing countries exist, very few studies have been done on Sri Lanka.

Methods and Materials

Assume a production function of the form of a conventional Cobb-Douglas function,

$$Y = f(FDI, K, L) \dots \dots \dots (1)$$

⁴⁸ sivashankar.p@hotmail.com

Where, Y is aggregate real output; K, capital stock; and L, labour force. Addition of more variables and logarithmic transformation yields the following equation (2).

$$\log GDPPC = \alpha + \beta_1 \log FDI + \beta_2 \log CAP + \beta_3 \log EXP + \beta_4 \text{Trend} + \beta_5 \log Open + \varepsilon \dots \dots \dots (2)$$

Where Log depicts the logarithmic form, GDPPC is GDP per capita, FDI is the foreign direct investment, CAP is the amount of capital, EXP is the exports, trend is the time variable for capturing technology improvements, and Open is the dummy for trade openness (0 for 1970 to 1977 and 1 for thereafter). α is the intercept of the full linear model and β 's are the respective elasticities. Data for the analysis were obtained from world bank and UNCTAD. The time period taken into account was from 1970 to 2011. The following tests were done to overcome the effects and avoid spurious regression.

Unit root test: The objective of the unit root test is to empirically examine whether a series contains a unit root or not. If the series contains a unit root, this means that the series is non-stationary. Otherwise, the series will be categorized as stationary. **Co-integration test:** This test was used to find out the long-term relation between the variables. **Granger causality test (GCT):** FDI and GDP growth are, in fact, interlinked and co-related through various channel. There is no theoretical or empirical evidence that could conclusively indicate sequencing from either direction. For this reason, the GCT will be carried out on FDI and GDPPC.

Results and Discussion

Initially, a GCT was conducted for FDI as dependent variable and GDP per capita as independent variable. Next, another GCT was performed where GDP was the dependent variable and FDI as independent variable.

Table 1. Results of granger causality test

Direction	F value	Probability
GDPPC does not granger causes FDI	1.36	0.2805
FDI does not granger cause GDPPC	2.61	0.0646*

Note: *significant at 10 % significant level

It is concluded that FDI can cause GDP per capita at 10% significant level. This causality is not very strong. Therefore, there is no robust unidirectional relationship of cause and effect. Since both test revealed that probability values were not significant, it can be concluded, that both can granger cause each other and lead to each other. Thus, there is bidirectional causality present between the variables, GDPPC and FDI.

To overcome serial correlation and to make data white noise, lagged variables or differenced variables can be used. First order derivatives of the variables were used to remove the non-stationary effect. By differencing, autocorrelation is removed and made to white noise. GCT was performed using Wald test. According to the results, a weak bi-directional causality between differenced logarithmic variables of GDP per capita and FDI inflows were obtained. A robust relationship between these two variables cannot be inferred, as they are only significant at 10 % significant level. Studies on other countries reveal most of the time a unidirectional or bidirectional relationship between these two variables exists in many instances.

Table 2. Results of the granger causality test after differencing

Equation	Excluded	χ^2	df	Prob> χ^2
Dln_gdp	Dln_FDI	8.8761	4	0.064 *
Dln_gdp	ALL	8.8761	4	0.064 *
Dln_FDI	Dln_gdp	7.9601	4	0.093 *
Dln_FDI	ALL	7.9601	4	0.093 *

*Note: *significant at 10 % significant level*

A regression analysis was employed with GDPPC as the dependent variables as in equation 2. It suggests that the model captures 98 % of variation of independent variables to GDP.

Table 3. Multivariate regression results

Variable	Coefficient	T value
logFDI	0.0477***	4.46
logCAP	0.4516 ***	7.29
logEXP	-0.0634	-0.83
open	- 0.8093***	-9.67
logtrend	0.5948***	9.07
Constant	12.0114***	9.6

$R^2 = 98\%$

*Note: ***significant at 1 %, **significant at 5 %, and *significant at 10%*

A 1 % increase in FDI increases GDP by 0.047 %, and a 1 % increases in Capital increase GDPPC by 0.45 %. Export coefficient is not significant. When the country is open for trade the GDP reduces by 0.80 %, this suggests that the exports composition might be mostly on raw materials and unprocessed items. This will reduce the income earned and rupee depreciation with the US dollar could have had a serious effect on the income earnings from exports. The dummy also captures only 7 years of closed trade. The dependent variable is on per capita basis, which suggest that the increase in FDI, Exports, and openness will not directly affect the per capita income, it

would take some time to materialize in the income per capita. Time increments increase GDPPC by 0.59 %, which also accounts for the technological improvements over the years. The host country situations, especially on human capital, infrastructure, technical know-how and governance issues matter a lot in properly utilizing the Benefits of FDIs.

Conclusions and Recommendations

This study examines the relationship between FDI and GDP in Sri Lanka using data from 1970-2011. The results indicated that FDI is one of the key determinants of Sri Lankan economic growth after the 1977 period. The empirical results suggest that one long-run equilibrium relationship exists between GDP, FDI, CAP, Trend and negative relationship with variable, OPEN. The Granger causality approach is then used to investigate the direction of causality flow in the short-run and long-run. The FDI inflows cause an independent influence on economic growth and direction of Granger causation is towards FDI to GDP growth and GDP growth to FDI. Thus, there is bidirectional Granger causality between FDI and economic growth. A unit increases in FDI causes GDP to rise by 0.047 units. GDP increases by 0.45 units per unit increment in capital. In general, the study appears to support the notion of foreign direct investment on GDP growth of Sri Lanka. This finding confirms the relevance of the economic reform programmes in Sri Lanka to reduce macroeconomic instability, remove economic distortions, promote exports and restore sustainable domestic investment for economic growth. However, the country's protectionist trade policies, direct and indirect regulatory barriers may have impeded foreign investment.

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INTER-RELATIONSHIP BETWEEN THE PREVALANCE OF TICK INFESTATION ON CATTLE AND THE FACTORS RELATED TO BREED, PHYSIOLOGICAL STAGE, MANAGEMENT AND DIVERSIFICATION SYSTEMS

E. Subalini¹⁴⁹, S. Dasinaa¹ and S. Thanuejah¹

¹*Faculty of Agriculture, Eastern University of Sri Lanka, Chenkalady*

Introduction

Ticks are economically the most important ectoparasite of cattle and other domestic species in tropical and subtropical countries (Jongejan and Uilenberg 1994). The ticks cause harm to animals through blood loss, general stress and irritation, depression of immune function and damage to hides and skin (Ghost et al. 2007). Although, economic losses due to ticks are mainly due to the diseases which they transmit (Garcia 2003), financial losses associated with nagging, irritation and depreciation of the value of skins and hides (up to 20 to 30 %) are also significant (Biswas 2003). Susceptibility and Resistance of animals to tick infestation have been influenced by several factors including; species, age, sex, season, breed, photoperiod and management.

In Sri Lanka, ticks are the most important of all ectoparasites. The economic loss incurred when they infest livestock particularly, cattle are enormous. To control the tick infestation efficiently well-documented information on the occurrence of ticks is utmost important. To develop a proactive program for ticks' control at smallholder and commercial level, it is important to determine prevalence of ticks and assessing the risk factors of tick infestation are important to recommend best tick control measures. In this context, the present study was designed with the objective to report the prevalence status of ticks in different breeds of cattle at various physiological stages under different management and diversification systems and to analyzing the associations related with the risk factors of the tick prevalence.

Materials and Methods

The study was conducted in four selected veterinary ranges such as Vantharumoolai, Kokkaddicholai, Kaluwanchikudy and Thumpankeni in Batticaloa district during the period from January 2014 to September 2014. The selected veterinary ranges had comparatively higher cattle population in Batticaloa district. For the analysis, 311 cattle were randomly selected irrespective to their age, sex, breed and physiological stages. Equal numbers of cattle were selected from each veterinary range. The parameters

⁴⁹ subalini123@yahoo.com

recorded were type of breed, sex of animals, physiological stage of animals, management and diversification system.

The raw data recorded from the study were entered into Microsoft excel data base system and using SPSS version 22 computer program, data were summarized and analyzed. Chi-square (χ^2) test was used to determine the variation in infestation prevalence between different peasant associations. A 5 % significant level was used to determine whether there were significant differences in the parameters measured between the groups.

Results and Discussion

Out of 311 animals, 124 cattle were tick infested among the study areas. The overall prevalence was 40 %. Chi-square value indicated no significant ($P>0.05$) association in tick infestation among different breeds of cattle. Higher prevalence of tick infestation was recorded in local cattle (51.61 %) as compared to exotic crossbred cattle (0.8 % to 16.12 %). The higher tick infestation in local cattle might be attributed to differences in management system, lack of supplementary feeding to local cattle breed, lack of control measures against tick on local cattle breeds. Furthermore, it can be assumed that it might be due to lack of interest of farmers about local cattle as well as taking more care to cross breeds than local cattle. Among the exotic crossbred cattle, higher tick infestation was recorded in crossbreds of temperate cattle (65 %) such as Jersey and Friesian as compared to crossbreds of tropical cattle such as Sahiwal, Sindhi and Tharparkar. This finding was supported by the report made by Sajid et al. (2009) where Pakistan's Sahiwal cattle is more resistant to tick infestation than European breeds. Tick resistance is a hereditary trait in *Bos indicus* cattle (Jongejan and Ulenberg 2004).

According to the Chi-square value ($\chi^2=20.64$, $df=4$, $P>0.05$) a significant association was observed in tick infestation among different physiological stages of cattle. Regardless of the locations under study, the higher prevalence of tick infestation (70.90 %) was recorded in adult cow (heifer, lactating, pregnant and dry) as compared to adult males (4.84 %) and calves (24.19 %). However, the findings of the present study was not agreed with the observation made by Hostis et al. (1996) and Swai et al. (2005) where the depicted that the tick infestation was higher in calves as compared to adult animals. Among the adult females higher tick infestation was observed in lactating animals (54.54 %). Moreover, it was observed that higher number of ticks were localized in an around the udder and teats. During the lactating stage, the lack of immunity in animals and soft tissues and thinner skins of udder would help in the penetration of mouth parts for feeding (Sajid 2007).

The results of the study further revealed that a significant association ($\chi^2=21.39$, $df =2$, $P<0.05$) was observed in tick infestation of cattle among

different management systems. Comparatively higher tick infestation was observed in cattle rearing under semi- intensive system (78.22 %) whereas 17.74% intensively operated and 4.03 % extensively operated farms were infested with ticks. Lower tick infestation observed in extensive management system in the study areas indicated that the extensive management system is not favor for the prevalence of ticks in cattle which was well agreed with the report of Durrani (2008) where it was stated that the grazing helps in lesser tick infestation and maximum tick anorectic effect. However, contradictory observation was made in another study (Belew et al. 2011) where it was recorded that the susceptibility of cattle for tick infestation is higher under extensive management system as the extensive cattle are moved anywhere for grazing.

The significant association was observed ($\chi^2= 7.94$, $df=3$, $P<0.05$) in tick infestation of cattle among different diversification system. Higher tick infestation was observed in cattle rearing in monoculture system (80.64 %) whereas under mixed farming system in which the cattle was a major component in crop and livestock farming, the lower tick infestation was recorded (2.42 %). It was observed that under mixed farming system the existence of natural predators for ticks such as crow, minah, and crane is high.

Conclusions and Recommendations

The results of the study revealed that there were significant association between the prevalence of tick infestation and sex, physiological stage of animal, management and diversification systems. However, no association was recorded between prevalence and type of breed. For further conclusion it should be needed to identify specific tick species and sub species prevailed in cattle under different farming system.

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POSSIBILITIES OF SHARING MULTI SPECIES IN A CAGE BASED ON THEIR FOOD PREFERENCE USING GRAPH COLORING TECHNIQUE

G. K. H. Thondilege¹⁵⁰, G. H. J. Lanel² and P. D. A. G. D. De Almeida³

¹*Institute of Engineering Technologies, Katunayake*

²*Department of Mathematics, University of Sri Jayewardenepura, Gangodawila, Nugegoda*

³*ShipXpress INC, No 247 1/2, Stanley Thilakarathne Mw., Nugegoda*

Introduction

During the evolution of zoos, from the mere presentation of animal collections, separately exhibited in single species cages, to the presentation of exhibits, set up to resemble natural habitats, the idea of mixing different species in the same enclosure became increasingly important. This study was set up to support the exchange of expertise in establishing and maintaining mixed species exhibits in zoos of Sri Lanka, as well as to promote the outstanding value of this concept for future animal keeping and exhibition facilities.

Graphs are structures formed by a set of vertices and a set of edges that are connections between pairs of vertices. Graph clustering is the task of grouping the vertices of the graph into clusters taking into consideration the edge structure of the graph in such a way that there should be many edges within each cluster and relatively few between the clusters. Graph clustering in the sense of grouping the vertices of a given input graph into clusters, by using any graph theoretical technique. In this research, graph coloring technique was used in each step of the clustering. The technique behind the vertex coloring for clustering cannot be explained simply as follows. Consider the famous example of sorting fish: A tropical fish hobbyist had six different types of fish designated by A, B, C, D, E, and F. Because of predator-prey relationships, water conditions, and size, some fish cannot be kept in the same tank. Table 1 shows which fish cannot be together. Aim is finding the smallest number of tanks needed to keep all the fish.

Table 2. Data of the Sorting fish example

Type of Fish	Type A	Type B	Type C	Type D	Type E	Type F
Type of Fish Can't be with	B, C	A, C, D	A, B, D, E	B, C, F	C, F	D, E

⁵⁰ ganesha193@gmail.com

Vertex coloring algorithm was applied for the graph and Table 2 shows the solution for the problem.

Table 2. Final solution of the sorting fish problem obtained using graph coloring technique

Tank 1	Tank 2	Tank 3
A & D	F & C	B & E

Similarly in this research, the aim was to investigate the possibilities of multi species sharing the same cage (multi species exhibits) at the National Zoological Gardens, Dehiwala and propose an algorithm that can be used to achieve that target for any zoological garden using graph coloring technique.

Materials and Methods

During the research the following previous works which were related, but not similar to this work were studied thoroughly in order to come up with a better output.

a) *Dynamic Graph Clustering Using Minimum Cut Trees (2006)* by Robert Gorke, Tanja Hartmann, and Dorothea Wagner, Faculty of Informatics, Karlsruhe Institute of Technology, KIT.

b) *Hazard rates for clustered populations of David's deer (2008)* by Marta Molinska Glura, Tomasz Szwaczkowski and Krzysztof Molinski.

Mainly this research was based on some attributes of the animals in the National Zoological Gardens, Dehiwala. A complete set of data relevant to the following attributes at the animals was collected considering 220 animals.

a) Animal Inventory (class, order, common name, and the initial group structure)

Eg. Mammalia, Primates, Toque Monkey, Male-1, Female-2, Unknown-1

b) Diet list of each animal in the inventory (common name, meal time, food type, and amount in g) and Table 3 shows an example for it.

Table 3. Example for the data collected relevant to the animal's diet list

Common name	Meal time	Food type	Provided Amount (g)
Toque Monkey	Morning	Amberalla	7
		Gram	57
		Plantain	121
		Water melon	100
		Apple	11
		Guava/Mango	64
		Orange	14
		Cabbage	7

Using the above data a graph was modeled by considering all the animals and all the food types as vertices and edges as the attribute which food type they are consuming. Then the observed clusters (See Fig. 1 for an example and three clusters were observed as shown) can be extracted as a sub graph at each step below by considering the animals as the vertices and considered attribute in the relevant step as the edges using the modeled graph above. A database was created to store the above data and all the partitioning parts were preceded using the graph coloring technique. Since the original graph consists with end vertices of each edge sharing the common attribute, graph coloring was applied for the compliment graph.

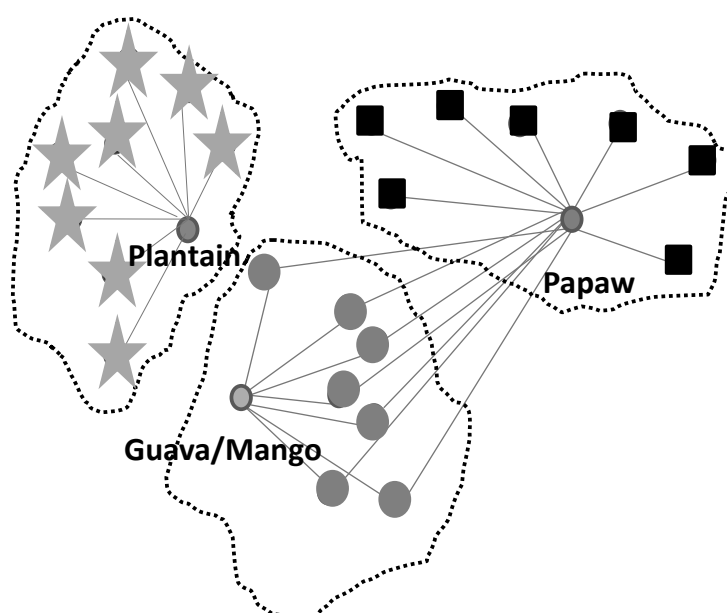


Figure 1. Part of the Initial graph that used for the analysis: (*Squares*: Two Humped Camel, Reticulated Giraffe, African Elephant, Slender Tailed Meerkat, Hooded Capuchin, White Throated Capuchin, Purple Faced Leaf Monkey), (*Stars*: Greater Kudu, Arabian Oryx, Scimitar Horned Oryx, Hog Deer, Sambar, Wild Boar, Domestic Ass, Mule), and (*Circles*: Red Giant Kangaroo, Red Necked Wallaby, Mouse Deer, Squirrel Monkey, Long Haired Spider Monkey, White Fronted Brown Lemur, Slender Loris)

The basic steps of the data analysis are given below and the number of exhibits obtained in each step is given within brackets compare to diet and using JAVA programing proceed all the steps on whole population of animals.

1) Grouped all the animals into three classes namely Mammals, Aves and Reptiles.[3 exhibits] (to avoid the predator-prey effect)

- 2) Grouped by carnivore order (Separated those mammals into different exhibits). [25 exhibits] (to avoid the predator-prey effect)
 - 3) The popular feed/feed combination as Papaw, Plantain and Guava/Mango was identified using the percentages of each food type. Overall 45% of all provided food types represent from that combinations. (Papaw – 19.5 %, Plantain – 14.1 %, Guava/Mango – 10.9 %)
 - 4) Again grouped the each exhibit obtained in step 1 into two exhibits as having papaw and not having papaw for feeding. [6 exhibits]
 - 5) Again grouped the each exhibit obtained in step 4 into two exhibits as having Plantain and not having Plantain for feeding. [12 exhibits]
 - 6) Again grouped the each exhibit obtained in step 5 into two exhibits as having Guava/Mango and not having Guava/Mango for feeding. [24 exhibits]
- Then finally 49 mixed species exhibits were obtained instead of 220 individual exhibits.

Results and Discussion

As a result after the step 6 there were 49 exhibits. For example,

1. For class Mammalia and relevant to the feed combination papaw, plantain and not having guava/mango.
Red giant kangaroo, Red necked wallaby, Mouse deer, Squirrel monkey, Long haired spider monkey, White fronted brown lemur, and Slender loris.
2. For class Mammalia and relevant to the feed combination papaw, guava/mango and not having plantain.
Silver pheasant, Malay great argus pheasant, Chinese ringed necked pheasant, Silky bantam, Eurasian collared dove, Diamond dove, victoria crowned pigeon, Green imperial pigeon, Domestic pigeon, and Spotted dove.

Conclusions and Recommendations

In this study the method of graph coloring technique was used to a real world problem and it has been successfully completed up to a certain level with 49 exhibits of mixed animals instead of 220 individual cages. This result can be used practically to build up a zoological garden within a limited area. For further studies we are planning to use other attributes such as height/length, and gestation, dominant character etc. in the analysis.

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BIODEGRADATION OF *Eichhornia crassipes* TOWARDS BIOFUEL PRODUCTION

A. Thurairajah¹, R.R. Ratnayake²⁵¹, N. Gnanavelrajah¹ and M. Kathirgamanathan²

¹ *Faculty of Agriculture, University of Jaffna, Sri Lanka*

² *Institute of Fundamental Studies, Hantana Road, Kandy, Sri Lanka*

Introduction

Energy is an essential input to our day-to-day activities. In the year 2012 the imports of fuel was 26.3 % of all imports (Central Bank of Sri Lanka 2013). Usage of fossil fuels results in the net increase of carbon dioxide in the atmosphere. Therefore, alternative sources of energy with high efficiency and low cost of production is important to meet the energy requirements. Biofuels can be defined as a fuel produced using biological material as a raw material and/or catalysts. Ethanol produced from lignocellulosic materials is considered as a potential transportation fuel. In Sri Lanka, the spread of invasive aquatic plant species is an environmental problem which needs to be controlled. This study aims to examine the appropriateness of water hyacinth (*Eichhornia crassipes*) as a low cost substrate for biofuel production.

Materials and Methods

Isolation of cellulolytic microorganisms: Decaying plant materials, compost samples, cow dung and elephant droppings were collected from different places in Kandy. Samples were inoculated on to Bushnell Haas agar with 1 % cellulose and carbendzim (1.7 mgL^{-1}) for bacteria and potato dextrose agar with the antibiotics streptomycin (30 mgL^{-1}) and chloramphenicol (50 mgL^{-1}) for fungi. For the isolation of anaerobic microorganisms, samples were enriched in a modified medium with cellulose for a week at 37 °C (Scott and Dehority 1965).

Production of enzymes: Liquid medium from Mandels and Weber (1969) was prepared for culturing isolated aerobic fungal strains with incubation for seven days at 28 °C with shaking at 100 RPM. The bacterial cultures were inoculated into Bushnell Haas broth with 1 % cellulose and 0.05 % proteose peptone and incubated at 35 °C for 3 days with shaking at 100 RPM.

Enzyme assays: Total cellulase assay was performed by using Whatman No.1 filter paper as the substrate, according to Mandels et al. (1976). Reducing

⁵¹ reukaratnayake@yahoo.com

sugars formed by the degradation were measured by using di-nitro salicylic acid reagent (Miller 1959), against D-glucose standards. Xylanase activities were measured by a method modified from Gottschalk et al. (2010) using beech wood xylan (Sigma) as the substrate. The above assays were carried out at 50 °C/pH 4.8 for fungi and 40 °C/pH 6.0 for bacteria.

Substrate collection and preparation: Water hyacinth (*E. crassipes*) was used as the substrate for degradation. It was collected from a site near Mahaweli river in Kandy district and was washed with flowing tap water, sundried for few days and kept in an oven at 50 °C until constant weight was obtained. Dried substrate was cut into small pieces and ground by using plant grinder and sieved through 50µm sieve. The sieved powder was used for the experiment.

Degradation of E. crassipes with enzymes from selected micro organisms: Isolates with high cellulase activities were chosen for the degradation. Enzymes were produced by submerged cultivation in a liquid medium (Mandels and Weber 1969). The pH of the enzymes was adjusted to 4.8 for fungi and to 6.0 for bacteria and they were filter sterile using sterile 0.45 µm nylon membrane filters. Five hundred milligram of *E. crassipes* powder was taken in to separate conical flasks. To each flask, 2 mL of distilled water was added and autoclaved at 121 °C for 20 minutes. 10 mL of sterile crude enzyme was added to each flask. 10 mL of distilled water was added to the controls. Degradation was carried out at 40 °C in with shaking at 100 rpm.

Measurement of reducing sugars: Total reducing sugars formed by each enzyme were measured at 24 hour intervals up to 7days of decomposition with di-nitro salicylic acid reagent (DNS) against D-glucose standards.

Data analysis: Statistical analysis was done using with SAS package (Complete Randomized Design design).

Results and Discussion

Isolation of cellulose degrading microorganisms: In the present study, 89 aerobic fungi and 22 aerobic bacteria were isolated from different substrates. Five strains of anaerobic bacteria were isolated from cow dung, elephant dropping and compost samples.

Enzymatic degradation of E. crassipes: Accessibility to enzyme is one of the major factors determining the degree of degradation of lignocellulosic materials. Smaller the particle size the better the hydrolysis in terms of the extent and rate of reaction. Therefore, the ground plant materials were sieved by using the smallest available sieve size (50 µm). The dried substrates were not subjected to any other significant pre-treatment than

size reduction. The fungal isolates F5, F22, F37, F41, F85 and F86 were selected for degradation of *E. crassipes*. In addition, the bacterial isolate B9 was also used for degradation.

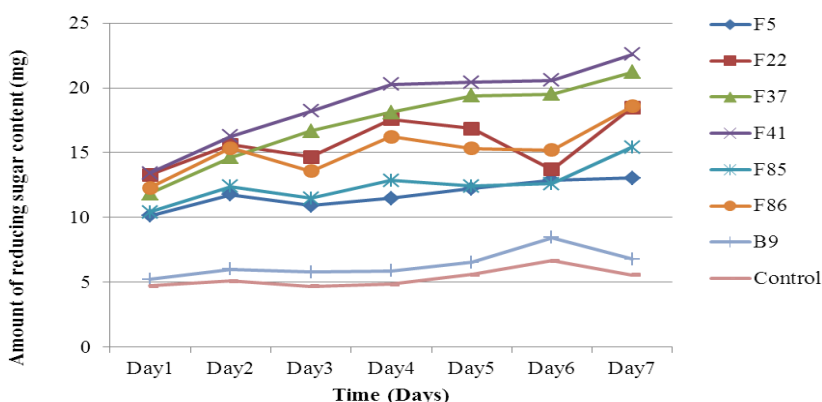


Figure 1. Cumulative quantities of reducing sugars released from *E. crassipes*

The rate of degradation was higher for the first day compared to that of subsequent days (Fig. 1). This may be due to hydrolysis of amorphous regions of cellulose by endoglucanases. The subsequent degradation of crystalline region occurs only at the ends of cellulose molecules therefore rates of formation of reducing sugar decreased (Lynd et al. 2002). This general pattern is observed with most of cultures. After 7 days of degradation, the highest amount of sugars released was 22.58 mg by F41, which was 4.5 % of the total dry weight of *E. crassipes* powder.

Identification of selected fungal isolates: The identified fungal isolates are summarized in Table 1 below.

Table 1. Identification of fungal isolates selected for degradation

Isolate	Genus
F5	<i>Aspergillus sp.</i>
F37	<i>Penicillium sp.</i>
F41	<i>Penicillium sp.</i>
F85	<i>Trichoderma sp.</i>
F86	<i>Trichoderma sp.</i>

Conclusions and Recommendations

Among the 89 isolated cellulolytic fungal strains, F5, F22, F37, F41, F85 and F86 were found to have high cellulase activity. Among these, F41 had the

highest sugar yield. In addition to these six fungal strains one bacterial strain B9 was also used in this study. The bacterial strain had lower cellulase activity compared to that of fungal strains. The highest amount of sugars released was 22.58 mg by F41, which was 4.5 % of the total dry weight of *E. crassipes* powder. Thus, the isolate F41 could potentially be used for degradation of *E. crassipes* for biofuel production.

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ASSESSMENT OF SOIL MICRONUTRIENT AVAILABILITY IN DIFFERENT CULTIVATED AND UNCULTIVATED SOILS IN JAFFNA PENINSULA

S. Vijayakumar¹⁵², R. R. Ratnayake² and N. Gnanavelrajah¹

¹ *Faculty of Agriculture, University of Jaffna, Jaffna*

² *Institute of Fundamental studies, Kandy*

Introduction

Soil fertility is the foremost important factor to maintain sustainable agriculture. It refers the ability of the soil to supply essential plant nutrients and soil water in adequate amounts and proportions for plant growth and reproduction in the absence of toxic substances which may inhibit plant growth (Foth and Ellis 1997). In the present commercialized world, increasing population has increased the demand on food and land. Farmers are forced to supply the required food through cultivation within the limited land. Thus, they are doing intensive agriculture with the use of high levels of fertilizers to provide the required nutrients and using hybrid varieties for increased production. Farmers mainly focus on macronutrients. Micronutrients are not supplied generally with fertilizers unless through organic manures. Therefore, continuous cultivation could lead to the depletion of micronutrients in soils. There are limited studies carried out on soils of Jaffna regarding micronutrients availability. Therefore, this study has planned to investigate the micronutrient availability in cultivated and uncultivated upland and low land soils in Jaffna district.

Materials and Methods

Study area: The study was carried out in Jaffna district, Northern Province of Sri Lanka. Jaffna experiences average annual rainfall of 696 mm to 1125 mm and mean temperature is 26 °C to 33 °C. Soil samples were collected by categories of upland and lowland and cultivated and uncultivated land. Here, lowland means paddy cultivated and its adjacent uncultivated lands where as upland means vegetable and field crop cultivated and its adjacent uncultivated lands. Three locations were selected in each category, though 12 locations were selected for sample collection. In this study, concentrations of micronutrient cations were investigated in cultivated and uncultivated lands of both upland and lowland areas at two depths (0-15 cm, 15-30 cm). The design was Completely Randomized Design in three factor factorial with three replicates.

Sample analysis: Soil samples were air dried at room temperature and crushed to pass through a 2 mm sieve. Soil suspension was prepared with 1:

⁵² sowthini@gmail.com

5 Soil : water ratio and pH was determined by using pH meter (Anderson and Ingram 1993). The available micronutrients Zn, Cu, Mn and Fe were determined by extracting the soil with DTPA- Ammonium bicarbonate extract and measured by using atomic absorption spectrophotometer (Soltanpour and Schwab 1977).

Table 1. Sampling sites in Jaffna district

Upland			Lowland		
Location	Cultivated/ Uncultivated	Abbreviation	Location	Cultivated/ Uncultivated	Abbreviation
Uduvil	Cultivated	JUC1	Valukaiyaru	Cultivated	JLC1
	Uncultivated	JUU1		Uncultivated	JLU1
Kondavil	Cultivated	JUC2	Moolai	Cultivated	JLC2
	Uncultivated	JUU2		Uncultivated	JLU2
Urumpirai	Cultivated	JUC3	Ponnaalai	Cultivated	JLC3
	Uncultivated	JUU3		Uncultivated	JLU3

Results and Discussion

Soil pH: Soil pH varied between 5.19 to 8.33 (Table 2) and was not significantly differed between soils, upland/lowland, cultivated/uncultivated lands and depth. However, mean pH was high in upland (7.54). In lowland 15-30 cm shows higher pH than the top soil. Root distribution of paddy occurs in the top soil. Since, most of the nutrient cations are being basic cations, when the crops uptakes the nutrients, that ultimately reduces the pH of top soil. This may be possible reason for this situation.

Table 2. Soil pH in different locations in Jaffna district

Place	Depth (cm)	pH	Place	Depth (cm)	pH
JUC1	0 - 15	7.2	JLC1	0 - 15	7.05
	15 - 30	7.32		15 - 30	7.08
JUU1	0 - 15	7.25	JLU1	0 - 15	6.24
	15 - 30	7.36		15 - 30	7.24
JUC2	0 - 15	7.79	JLC2	0 - 15	5.19
	15 - 30	7.89		15 - 30	6.88
JUU2	0 - 15	7.26	JLU2	0 - 15	6.4
	15 - 30	7.34		15 - 30	7.36
JUC3	0 - 15	8.09	JLC3	0 - 15	6.15
	15 - 30	7.3		15 - 30	6.14
JUU3	0 - 15	8.33	JLU3	0 - 15	6.5
	15 - 30	7.43		15 - 30	7.62

Available micronutrients:

(1) Iron- Iron availability showed significant difference among the different sampling locations. The highest value of $227.33 \text{ mg kg}^{-1}$ was obtained in surface layer of lowland uncultivated areas (JLU1) and the lowest value of 3.57 mg kg^{-1} was obtained in subsurface layer of upland cultivated soils. Soil Iron content was significantly differed with lowland/upland. It was higher in lowland than upland, which is due to the low pH in lowland than upland and hence the iron becomes more available in low pH (Ponnamperuma 1972). According to Benton (2001) classification, availability of iron can vary from very low ($0-5 \text{ mg kg}^{-1}$) to very high ($> 25 \text{ mg kg}^{-1}$). Only the samples taken from sub-surface layer of JUC2, JUU1 and surface layer of JUC2 soils experienced very low, low and medium level iron availability respectively. Both layer of JUU2, JUC3 and surface layer of JUU1, JLC3 falls under the high ($17 - 25 \text{ mg kg}^{-1}$) iron availability and other soils had very high iron availability (Table 3).

(2) Manganese- Mn availability ranged between 6.11 and 35.93 mg kg^{-1} , which lies in low ($4 - 8 \text{ mg kg}^{-1}$) to very high ($>30 \text{ mg kg}^{-1}$) availability category according to Benton classification in 2001. Manganese availability varied significantly only with elevation which was high in upland. JUC1 had very high amount of Mn availability where as only the upper layer of JLC1 resulted as low Mn availability area. Mn availability in other areas experienced medium and high amount (Table 3).

(3) Zinc- Zinc availability ranged between $4.8 - 46.1 \text{ mg kg}^{-1}$. Zinc availability in soils of lowland/upland and cultivated/uncultivated was significantly differed and those were high in upland and uncultivated areas respectively. Zinc availability showed significant different with in the different sampling places (Table 3). According to Benton (2001) Zn availability relies in high ($3.1-6 \text{ mg kg}^{-1}$) to very high ($>6 \text{ mg kg}^{-1}$) category in Jaffna soils.

(4) Copper- Cu availability ranged from 0.67 to 10.38 mg kg^{-1} and showed significant difference within the sampling places (Table 3). These soils falls between low ($0.3-0.8 \text{ mg kg}^{-1}$) to very high ($>2.5 \text{ mg kg}^{-1}$) range in Cu availability according to Benton classification, 2001. Except JLC1 and JLU1, other soils had very high Cu content. Low Cu content was observed in surface layer of JLC1. There was no significant difference observed in Cu availability between soils of lowland/upland, cultivated/uncultivated and depth-wise.

Note: Means with same letter along the columns are similar

Table 3. Mean available micronutrients concentrations in different places in Jaffna

Place	(mg kg ⁻¹ of soil DW)							
	Fe		Mn		Zn		Cu	
	0 - 15 cm	15 - 30 cm	0 - 15 cm	15 - 30 cm	0 - 15 cm	15 - 30 cm	0 - 15 cm	15 - 30 cm
JUC1	47.57 ^c	30.53 ^{de}	35.93 ^a	34.07 ^a	16.50 ^{bc}	15.43 ^{ef}	9.4 ^b	6.86 ^{ab}
JUU1	17.46 ^d	6.52 ^{fg}	22.35 ^d	19.2 ^c	25.10 ^a	26.30 ^b	5.59 ^e	5.06 ^{abc}
JUC2	13.48 ^d	3.57 ^g	20.77 ^e	12.92 ^e	25.56 ^a	18.97 ^{cd}	5.53 ^e	4.03 ^{abcd}
JUU2	18.92 ^d	22.77 ^e	24.38 ^c	23.26 ^b	26.24 ^a	20.62 ^c	5.56 ^e	4.81 ^{abc}
JUC3	20.88 ^d	18.07 ^{ef}	26.81 ^b	14.58 ^d	13.31 ^c	11.69 ^{gh}	8.72 ^c	7.87 ^a
JUU3	45.92 ^c	43.55 ^{cd}	14.41 ^f	15.25 ^d	20.63 ^b	46.09 ^a	5.45 ^e	5.17 ^{abc}
JLC1	224.84 ^a	48.95 ^c	6.11 ^j	8.15 ^g	6.04 ^d	5.73 ⁱ	2.38 ^g	0.67 ^{cd}
JLU1	227.33 ^a	148.83 ^a	9.02 ^h	10.97 ^f	4.81 ^d	11.90 ^{gh}	1.1 ^h	1.61 ^{cd}
JLC2	182.72 ^b	119.5 ^b	20.02 ^e	14.23 ^{de}	14.75 ^c	13.41 ^{fg}	10.38 ^a	7.41 ^a
JLU2	214.68 ^a	121.18 ^b	10.83 ^g	10.0 ^f	15.74 ^c	16.88 ^{de}	8.1 ^d	5.83 ^{ab}
JLC3	23.65 ^d	37.71 ^d	7.52 ⁱ	10.06 ^f	4.86 ^d	9.01 ^h	2.9 ^g	3.15 ^{bcd}
JLU3	48.45 ^c	38.86 ^{cd}	8.52 ^{hi}	8.44 ^g	16.77 ^{bc}	25.41 ^b	4.05 ^f	3.86 ^{abcd}

Conclusions and recommendations

There was a significant variation in micronutrients namely Fe, Mn, Zn and Cu among different places in Jaffna soils. Fe content in soils varied between 3.57 - 227.33 mgkg⁻¹, even though except one place other places had very high amount. Mn, Zn and Cu content ranged between 6.11 – 35.93, 4.81 – 46.1 and 0.67 – 10.38 mgkg⁻¹, respectively. As the nutrient availability varies significantly from place to place in the study area site specific soil testing and fertilizer recommendation for micro nutrients will be useful.

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ANALYSIS ON THE RISE OF RELIGIOUS CRISIS AND NATIONAL INTEGRATION OF POST WAR SRI LANKA

N. Weerasekara¹⁵³ and R.K.S. Navodani¹

¹Department of Social Sciences, Sabaragamuwa University of Sri Lanka, Belihuloya

Introduction

Nation building and national integration are necessary conditions for the development of a country. The main development goal of Sri Lanka is to be a miracle of Asia. As a democratic country, Sri Lanka has accepted to protect the equality, human rights, and minority rights without any discrimination. But Sri Lanka has been facing the problem of nation building from the independence to date. Within post war context, ethnic conflict has been transformed into a new form of religious crisis in Sri Lanka. There has been a domination of Sinhala Buddhist nationalism for pre-colonial period to post war period in Sri Lanka. The priority is given for the Sinhala Buddhist identity from policy formulation to party representation and to civil society activities. Sinhala Buddhist identity was represented even in the political leadership (Phadnis 1989), and they used the Sinhala Buddhist nationalist theme in their election movements (Phadnis and Ganguly 2001). Even the legislature tried to give more opportunities for the Sinhales excepting the minorities, formulating the Sinhala Buddhist only acts such as 1947 citizenship act, 1956 Sinhala Language act and 1972 and 1978 constitutions has been given the more priority to the Buddhism (Warnapala 1991). 1957 Bandaranaike-Chelvanaiyagam agreement had to tear because of the Sinhala Buddhist protest. Especially after the civil war, to build up the ethnic harmony and national integration became as a vital goal of present Sri Lanka. But although the government has implicated many steps for that, a new form of religious crisis has arisen. Thus, it is worthy to study the reasons and factors that has affected for the rise of religious nationalism and to examine its impact to the national integration of Sri Lanka.

Materials and Methods

The study mainly based on the secondary data gathered from the secondary sources such as books, articles, party constitutions and Sri Lanka constitution, contemporary reports, contemporary newspapers, acts, magazines, internet articles and books, and the qualitative data analysed by the Descriptive Data Analysis Method. In addition to that the study used primary data selecting 50 respondents to investigate the public opinion on the religious crisis and national integration in Sri Lanka covering farmers,

⁵³ weerasekeranelum@gmail.com

civil society activist, party activists, academic scholars, religious society activists and public using interview method. The study was limited to the time period from 2000 to date.

Results and Discussion

Today ethnic based crisis has become as a religious base crisis. There are many of reasons for that. The first reason is the impact of the Sinhala Buddhist Nationalism to the politics. The government and the political leaders are depended on the Buddhist instructions of the Malwathu and Asgiri Mahanayaka Theroes for the important political decisions and policies. As the examples; the president Mahinda Rajapaksha met the Maha theores to discuss the problems in the Bhuddha Sasana on 2013.5.22 (<http://sinhala.theindependent.lk>), The Former president Mrs. Chandrika Bandaranaike met the Thibbatuwawe Siddhartha Thero on 2014.01.18, Rawana Balaya and Bodu Bala Sena met the Maha Thoreos to discuss on the Casino business regulation act and the problems in the Law Faculty on 2014.04.23 (<http://www.hirunews.lk>), Minister Dinesh Gunawardena met the Maha Theores to get instruction on the North provincial election and the provincial powers on 2013. 05. 21 and Minister G.L Peeris met the Maha Theoros to get instruction on the rehabilitation and the educational issues on 2012.08.28. And also the present government has established the Buddhist Advisory Committee and *Gihi Pewadi Anusashaka Sabhawa* for the religious advice for the political issues on 2014. 06. 18. Another main reason is the endowment of the main political leadership only to the Sinhalese from D.S. Senanayake to Mahinda Rajapaksha. Another factor is to use the political power against to the minority norms and customs by politicians. Some political leaders tried to disturb the Hindu religious tradition in Munneshwaram Kovil.

Another main factor for the rise Sinhala Buddhist nationalism is to maintain the Buddhist religion by the Ministry of Bhuddha Sasana. And also Sri Lankan constitution and the political party constitutions have given the foremost position for the buddhism. Article 10 and article 14(1) have given the foremost position for the Bhuddism with respecting to other religions (Sri Lanka Constitution 1978). The article 2 of the Sri Lankan Freedom Party constitution has formed the Sri Lanka Freedom Monk Society under the Sri Lankan Freedom Party. And also article 11, 5, has implied that it should be the main part in the Executive Committee (Sri Lanka Freedom Party Constitution 1951). Although United National Party hasn't given the foremost position for the Bhuddism, both parties are engaged the Buddhist monks with their political activities. Another main problem is to use the religious nationalist ideas for the electoral victories by the political leaders. Although the main political parties haven't used the religious and nationality based names to address their parties, in practice they are totally religious alleged

and many of minority political parties have based on their religion and nationality. As the examples; National Heritage Party, All Ceylon Tamil Congress, Illankai Tamil Arasu Kachchi, Sri Lankan Muslim Congress and Eelam *National* Democratic Liberation Front. And also when the political coalitions form they have addressed only the political advantageous not ethnic harmony. There is no a unity even among the political leader on the contemporary religious issues within the UPFA.

Another main factor for the emergence of the religious crisis is to emergence of the Bhuddist political parties and the Bhuddist civil societies against to the minority religious extremism and to protect the Bhuddism such as National Heritage Party, National Monk Front, Desha Hithishee Jathika Viyaparaya, Sinhala Rawayana Organization, and Bodu Bala Sena Organization for the cases such as to aginst to Watareka Wijitha Thero, Pitiduwe Siri Dhamma Thero, Ruwanwelle Sobitha Thero (Rivira 12.3.2012), to Remove the Muslim Church form the Dambulla Secred Lanad and to release the Kuragala secred site on 2012.01.08 (Lakbima 19. 01.2012). The steps taken by the government for the nation building has mainly addressed on the language problem. It has formed the ministry of National Language and Social Integtration and has given its leadership to a minority leader. Although the government has taken some steps to prevent the religious conflicts such as to formulate a National plan to practice the Lessont Learnt Commission Report, to held *Maha Samuluwa* for the national harmony on 2014.4.7 (Divaina 8.4.2014), to eshtabish a special police task on the religious conflicts, to held the *Wishwasaye Agamika Sammuthi Wadasatahana* and to form a committee for the religious harmony by the South Asian Policy and Research Centre, the practice and the outcomes are under questioned. According to the reseach and the investigation of the State Language Commission, there are many of weaknesses of the second language training programme(Perera 2008) and there are many of problems and challenges of language policy practicers in the public administrative institutions in Sri Lanka (Ranjith and Weerasekara 2013). According to Lanka Law Society, new police task is a threat for the spirit of law (<http://lankafocusnews.com>).

Conclusions and Recommondations

In the whole, the factors and reasons such as, to give the priority to the Sinhala Buddhist identity by the tradition, law, policies, political parties and political leaders, and rise of the Sinhala Buddhist civil societies and political parties against to the radical minority religious prevalence and mistrust that the government has evaded in representing the Buddhism after the civil war have affected to rise of the religious crisis among the religious groups and it has affected to infract the national integtration of Sri Lanka because the government hesitated and feared to implicate a strong stept to prevent the

religious crisis withing the post war context. Therefore the political leaders have used the religious nationalism to fulfil their electoral purposes. Therefore the state constitution, party constitutions and ministry of Buddha Sasana and Religious Affairs should accept the equal position of all religions without limiting to a one religion. And also it is better to join the all religions to get advisors to the politcs. Coalitions should be used for nation building purposes other than the power purposes. Minority political parties should address the every citizens of the country without limiting to only their nation. It is better to implicate a strong stept and laws to prevent the prevalence of the radical religious hejoman. And also the government should have the stable vision on the buiding the national integtration. When the government hasn't a stable vision and a plan to build up the national integration of Sri Lanka, it will be provided a path to rise of a religious conflict in Sri Lanka. Therefore the government should have a stable solution to build up the national intergtation without hesitating in Sri Lanka.

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