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## **PATHOGENESIS OF PULLORUM DISEASE (PD) IN CHICKENS BY LOCAL ISOLATE OF *Salmonella pullorum* IN BANGLADESH**

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### **ABSTRACT**

Pullorum disease is caused by *Salmonella enterica* subspecies *enterica* serovar Pullorum. Chickens are the natural host of this pathogen. In the present study experimental pathogenesis was studied. Twenty pullets (*Salmonella pullorum* seronegative) of Isa Brown breed of 18 weeks and 10 cocks (*Salmonella pullorum* seronegative) of RIR breed of 26 weeks of age were experimentally infected orally with  $2 \times 10^7$  (CFU) dose of *Salmonella Pullorum* organisms and in control group no bacteria was given. Birds were observed for clinical signs, gross pathology, and reisolation of *S. Pullorum* from different organs and blood, histopathological study, detection of antibody levels and detection of *S. Pullorum* by PCR at different time intervals of experimental period. Four hens and one cock were randomly selected and sacrificed on 6 hr before inoculation and 1 wk, 2, 3 and 4 wks of post infection (PI). Samples were collected for bacteriological, serology and histopathological examinations. Liver, lungs, ovarian follicles and testis were also collected in 50% buffered- glycerol and preserved in  $-80^{\circ}\text{C}$  for PCR. The clinical signs of infected hens were found at 72 hrs of PI, which continued up to 4 wks. 15.81% reduction in egg production was observed. The highest mean CFU ml<sup>-1</sup> of *Salmonella Pullorum* from blood was  $13.55 \times 10^3$  at 1 wk PI and the lowest was  $13 \times 10^2$  at 4 wk PI. Gross lesions were variable in different birds at different time interval. The highest gross lesion was 93.75% as swollen and congested spleen and the lowest lesion was 43.75% as pericarditis and necrotic foci/ nodules in the heart. Microscopically, the liver showed congestion; hepatitis with infiltration of inflammatory cells, and focal necrosis with nodule formation. The antibody titre increased gradually and the highest titer was at 4 wks PI in hens ( $4712 \pm 1851$ ) than that of cock ( $3059 \pm 903$ ). *S. Pullorum* was

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detected by PCR in all liver and lung samples from 1 wk to 4 wks PI. *S. Pullorum* was reisolated from male and female reproductive organs after experimental infection. *S. Pullorum* was detected by PCR at 1 wk to 3 wks PI from testicular tissues. *S. Pullorum* was also reisolated from 50% eggs of experimentally infected birds.

**Keywords:** Chickens, pathogenesis, pullorum disease

## INTRODUCTION

Pullorum disease is one of the major constraints of poultry industries in Bangladesh (Das et al., 2005). Khan et al. (1998) recorded 12% morbidity and 75% mortality at the age of 5 wks in broiler breeder replacement pullets of Shaver Red Bro breed at Bangladesh Agricultural University Poultry Farm. They also recorded 100% morbidity and 75% mortality in local birds at the age of 8 wks with oral dose of 0.5 ml of  $10^7$  CFU of *S. Pullorum*. A total of 33,204 birds at different groups were investigated for Salmonella infection in Bangladesh. The morbidity and mortality rate due to salmonellosis among the birds were 4.90% and 1.83%, respectively. The mortality rate was highest (68.53%) in the early age groups (0-3 months). It decreased with age and was reduced to 0.25% in the twelve months and above age group (Khan et al., 1998).

PD causes great economic losses every year in poultry farms and it has also public health significance (Shivaprasad, 1997). The diseases can spread via meat and eggs. A few investigations on natural cases of Salmonella infections have been completed in Bangladesh using the methods of necropsy, histopathology and isolation of bacteria by culture; staining and sugar fermentation tests (Haider et al., 2008). However, no investigations have yet been performed by locally isolated *Salmonella Pullorum* organisms in respect of pathogenesis, pathology and vertical transmission in chickens. For this reason, the present study was taken for better diagnosis, prevention and control of this economically important pullorum disease in Bangladesh.

## MATERIALS AND METHODS

The proposed research work was conducted at the Department of Pathology, Faculty of Veterinary Science, Bangladesh Agricultural University, Mymensingh during the period from March 2007 to March 2009.

### Experimental hens and cocks

A total of 40 pullets (*Salmonella Pullorum* seronegative) of Isa Brown breed of 18 wks were purchased from Nourish Hatchery Ltd., Shreepur, Gazipur, Bangladesh and 10 cocks (*Salmonella Pullorum* seronegative) of RIR breed of 26 weeks old were taken from BAU Poultry farm. The birds were vaccinated against MD, IB, IB, Fowl Pox and ND obtained from Intervet, Holland. The birds were divided into two groups in which one group remained as control.

### **Bacterial infection**

25 birds (20 hens at the age of 21 wks and 5 cocks at the age of 29 wks) were experimentally infected orally with  $2 \times 10^7$  (CFU) dose of *Salmonella enterica* sub. *enterica* serovar Pullorum (Isolate no. 5) organisms in 0.5 ml broth culture with 0.5 ml of sterile phosphate buffer saline (PBS), pH 7.2, using sterile syringe (Roy et al., 2001; Wigley et al., 2005). Control birds were given only 0.5 ml of nutrient broth without bacteria with 0.5 ml of PBS.

### **Samples collection**

Five birds (4 hens and 1 cocks) in each case were randomly selected and sacrificed on 6 hrs before inoculation and 1, 2, 3 and 4 wks of post infection. A total of 25 (20 hens and 5 cocks) birds were used for the control group, and necropsied in similar way along with the infection groups. Different types of samples were collected as described early. Eggs, ova and parts of female reproductive organs were collected for the isolation of organisms (Wigley et al., 2005).

### **Clinical signs**

Clinical signs of chickens after experimental infection were observed and recorded up to necropsy (Okamura et al., 2000; Wray and Davies, 2001).

$$\text{Feed consumption was calculated by} = \frac{\text{Total feed supplied} - \text{Amount of residues}}{\text{Total number of birds}}$$

$$\text{Loss of egg production (\%)} \text{ calculated by} = \frac{\text{Eggs control group} - \text{Eggs in infected group}}{\text{Eggs in control group}} \times 100$$

### **Gross Pathological study**

At necropsy, gross tissue changes were observed and recorded carefully (Roy et al., 2001; Wray et al., 2001).

### **Reisolation of *Salmonella Pullorum* from different organs**

Collected samples (crop, liver, lung, heart, duodenum, cecum, kidney, bile and spleen) were weighed and placed in a tube containing 1ml of sterile phosphate – buffered saline (PBS) solution. The colony-forming units of *S. Pullorum* were counted followed by standard methods (Haider et al., 2008).

### **Reisolation of *Salmonella Pullorum* from the blood**

For each bird of each group of 6 hrs before inoculation and 1, 2, 3 and 4 wks of post infection, 1 ml of blood was collected and *S. Pullorum* was reisolated as previously described (Haider et al., 2008).

### **Histopathological study**

The formalin-fixed tissues were trimmed, processed, sectioned and stained as per standard procedure (Haider et al., 2008). Samples of specific lesions from each group were used in histopathological study.

### **Immunological responses at the onset of egg laying**

Serum samples were collected at the onset of egg production and ELISA test was performed using early described standard procedure (Barrow et al., 1992).

### **Detection of *Salmonella Pullorum* in the hens by PCR**

Genomic DNA of *S. Pullorum* was extracted from liver, lung and follicle tissue samples using DNA extraction kits (Promega Corp. Madison, WI, USA). Extracted DNA amplification was carried out using commercial PCR kits (PCR Master Mixture Kits, GeNei™, Bangalore, India) in Gene amplification PCR system 9600 Thermocycler (Eppendorf, Germany). Amplified products were separated by electrophoresis on 1.5 % agarose gel containing 5 µg ml<sup>-1</sup> ethidium bromide with a 100 bp ladder as molecular weight marker (Desai et al., 2005; Olivera et al., 2003).

### **Statistical analysis**

Significance of different groups was determined by the application of chi-square or Fisher's exact probability test using SPSS or MSTAT-C computer software program. Repeated measures analysis were performed with the data of mean feed intake and body weight gain of pullet of different groups at different weeks in a Completely Randomized Design (CRD) for significant variation using the SPSS package program version 10.0. Pair wise comparison of means was done by Least Significant Difference (LSD). The differences in the increase or decrease of the ELISA antibody titre of hens and cocks of different groups at different weeks were analyzed for analysis of variance in a Completely Randomized Design (CRD) using an MSTAT-C computer package program. Significant differences between means were identified by Least Significant Differences (LSD).

## **RESULTS**

### **Clinical signs**

All hens and cocks were depressed at 1 and 2 wks PI and showed loss of appetite at 1 and 2 wks of PI. Their feed intake dramatically was reduced significantly ( $p < 0.01$ ) in infected group. The mean body weights of hens and cocks differed significantly ( $p < 0.01$ ) between infected and control group. Cumulative clinical signs of hens were loss of appetite 60%, depression 60%, diarrhea 20% and emaciation 12% and loss of egg production 15.81%. The mean egg production differed significantly ( $p < 0.01$  and  $p < 0.05$ ) in infected hens. All clinical signs were not found in all birds. The morbidity (clinical signs) was found 90% in overall 90% birds were found to be worked for experimental infection group. No mortality was

found in infected group during study period. Recovery from clinical signs began at 2 wks PI. No clinical signs and mortality were seen in control group.

### **Gross pathological study**

The haemorrhagic and congested liver and congested, edematous and brown coloured lungs were observed at 1wk of PI. The gross lesions were reduced gradually at 3 wks of PI. The gross findings were hemorrhage and congestion in 68.75% liver and necrotic foci in 56.25% liver; pericarditis and necrotic foci/ nodules in 62.5% heart; congested, edematous and brown colour in 68.75% lungs; caseous materials in 81.25% intestine, semi-solid, cheesy material and button like ulcer in 87.75% ceca (Figure 1); swelling and congestion in 93.75% spleen; enlargement in 56.25% kidney; misshapen discoloured cystic and congestive in 81.75% ova (Figure 2); and hemorrhage and congestion in 75% oviducts. The affected ova contained oily and caseous material enclosed in a thickened capsule. These degenerative ovarian follicles were closely attached to the ovary and they were pedunculated and detached from the ovarian mass.

Cocks were infected with  $2 \times 10^7$  CFU of *S. Pullorum* at 29 wks of age. Pericarditis, haemorrhagic and congested liver, congested and caseous lungs, and white foci in the testis were observed in cock at 2 wks PI. The size of the infected testes were reduced compared to control cock. The lowest weight of testis was 9.53 gms at 3 wks PI in infected cock. The mean weight of testes reduced significantly ( $p < 0.05$ ) in infected cocks. Gross lesions were variable in different birds at different time interval. The highest percentage of birds showed (93.75%) in swollen spleen, while the lowest lesion was reported 56.25% in enlarged kidney in infected hens.

### **Reisolation of *Salmonella Pullorum* from different organs of hens**

Reisolation of *Salmonella Pullorum* from different organs was variable at different time schedule (Table 1). *S. Pullorum* was reisolated from liver (93.75%) (Figure3), lungs (100%), duodenum (100%), ceca (100%), and spleen (100%) at 1 wk to 4 wks PI (Table 1). The *S. Pullorum* was also reisolated from crop (81.25%), heart (87.5%), bile (18.75%), and kidney (75%) during the study period. The most frequent reisolated *S. Pullorum* was  $64.58 \times 10^5$  in liver at 1 wk PI and the less frequent was  $14.96 \times 10^1$  in crop at 4 wks PI. The details mean CFU/gm of reisolated *S. Pullorum* at different time intervals are shown in table 1.

### **Reisolation of *Salmonella Pullorum* from blood**

The mean CFU/ml of *Salmonella Pullorum* was reisolated from blood and shown in table 1. The blood sample of four hens out of four (4/4) at 1wk and 2 wks PI, three hens out of four (3/4) and one hens out of four (1/4) at 3 wks and 4 wks PI, respectively were positive for *S. Pullorum*. The highest number of reisolated *S. Pullorum* was  $13.55 \times 10^3$  at 1 wk PI and the lowest number of reisolated *S. Pullorum* was  $13 \times 10^2$  at 4 wks PI from blood sample. No *S. Pullorum* was found in control group.

### **Reisolation of *S. Pullorum* from female reproductive organs**

*Salmonella Pullorum* was reisolated from ovary (100%) (Figure 4), ovarian follicle (100%), oviduct (68.75%), uterus (56.25%) and vagina (75%) of female reproductive organs after experimental infection (Table 2). *S. Pullorum* was not found in control group.

### **Reisolation of *S. Pullorum* from different organs of cocks**

Reisolation rate of *Salmonella Pullorum* from different organs was variable in different time schedules (Table 3). *S. Pullorum* was reisolated from liver (100%), lungs (100%), heart (75%), cecum (100%), spleen (100%), and testes (75%) at 1 wk to 4 wks PI. Control group was free from *S. Pullorum* in culture during the study period.

### **Histopathological study**

The liver showed (81.25%) congestion, hepatitis, infiltration of inflammatory cells; multifocal necrotic foci with infiltration of histiocytes in (56.25%) liver parenchyma; nodule formation with the infiltration of heterophils, lymphocytes, macrophages and plasma cells in (12.5%) liver (Figure 5). 75% lungs developed pneumonia and bronchopneumonia which were characterized by hemorrhage, infiltration of neutrophils and lymphocytes, heterophils and mononuclear cells in lung alveoli, and lumen and wall of the bronchus. In 62.5% heart and pericardium, infiltration of heterophils and lymphocytes was found while nodule was formed with infiltration of heterophils, macrophage and lymphocytes in 31.5% heart of hens (Figure 6). Focal necrosis and infiltration of heterophils and RE cells were found in 93.75% spleen. The intestinal (81.25%) and cecal mucosa (87.5%) exhibited necrosis and infiltration of mononuclear cells in the submucosa. Congestion and infiltration of heterophils and lymphocytes were seen in (56.25%) kidneys. Ovary (87.5 %) and oviduct (75%) showed haemorrhage and congestion with infiltration of macrophages, plasma cells, heterophils and lymphocytes (Figure 7). In males, degeneration and necrosis of spermatogonia and infiltration of inflammatory cells (heterophils, lymphocytes and plasma cells) in the somniferous tubules of testes were (75%) found (Figures 8). No lesion was found in cocks of control group.

### **Immune response in hens after experimental infection with *S. Pullorum***

Antibody titre (Mean $\pm$ SD) against *Salmonella Pullorum* was determined in sera collected at different time intervals and the results are shown in table 4. The antibody titres of infection group increased significantly ( $p<0.01$ ) and the values were 1211.82 at 1 wk PI and reached 4712.39 at 4 wks PI. The antibody titres increased slightly in basal level in control group ranging from 151.69 at 1 wk to 171.29 at 4 wks but the values were statistically insignificant.

### **Immune response in cocks after experimental infection with *S. Pullorum***

Antibody titre (Mean $\pm$ SD) against *Salmonella Pullorum* was determined in sera collected at different time intervals and the results are shown in table 5. The range of antibody titre in experimental *S. Pullorum* infected cocks was 508.20 to

3059.27 at 1 wk PI to 4 wks PI. In control cocks, the antibody titre varied from 202.74 to 172.56 at similar time intervals with infected group, but these values were insignificant. The antibody titres of infection group increased significantly ( $p < 0.01$ ) while the antibody titres decreased gradually in basal level in control group.

#### **Detection of *Salmonella Pullorum* by PCR**

*Salmonella Pullorum* was detected by PCR at different time intervals. No *S. Pullorum* was detected by PCR 6 hrs BI and control birds during study period. *S. Pullorum* was detected from four out of four liver (4/4), lungs (4/4) and ovarian follicle (4/4) from 1 wk to 3 wks PI in experimental group (Figure 9). Three liver (3/4), and four lungs (4/4) and four ovarian follicles (4/4) out of four at 4 wks PI were positive for *S. Pullorum* by PCR.

### **DISCUSSION**

In the present study, the oral route of experimental inoculation demonstrated that *S. Pullorum* caused bacteremia and colonized in the liver, lungs, heart, kidney, intestine, spleen, and ceca of chicks to various degrees that corresponded with the findings of Wigley, et al. (2005). Clinical signs appeared in chicks after 12 hrs PI. These were loss of appetite, depression, droopiness, ruffled feather, and diarrhea, labored breathing, loss of body weight, and pest vent. The clinical findings in this study were similar to the natural/experimental findings of other authors (Shivaprasad, 1997; Wary and Davies, 2001). No mortality was found in adult birds in this study, which was similar to the findings of others (Roy et al., 2001; Wary and Davies 2001). Clinical signs of infected hens were limited to slight depression (100%) and diarrhoea (75%) that lasted for 3 days after inoculation with the field isolate of *Salmonella Enteritidis* phage type 4 (Kinde et al., 2000; Okamura et al., 2001).

The gross findings were haemorrhagic and congested liver, necrotic foci in the liver, pericarditis and necrotic foci/ nodules in heart, congested, edematous and brown colour lungs, semi-solid, cheesy material in ceca, unabsorbed and coagulated yolk, swollen and congested spleen and enlarged kidney. In this investigation, the gross findings described above corresponded with slight variation with the findings of other authors (Chauan and Roy, 2007).

In the present study, after experimental infection *S. Pullorum* was reisolated from different organs at 1 wk PI. But the maximum number of *S. Pullorum* was reisolated from 1wk to 2 wk PI. From 3 wk PI reisolation of *S. Pullorum* organisms reduced gradually. The findings of the reisolation and identification of *Salmonella Pullorum* organisms from different organs have also been described by other investigators (Hoop and Pospishil, 1993; Okamura et al., 2000). The highest percentage of reisolation of *S. Pullorum* from liver (100%), lungs (100%), cecum (100%) and spleen (100%) in hen at 7 days PI corresponded with the reisolation of Roy et al, (2001). Wigley et al. (2001) also recovered *S. Pullorum* after experimental inoculation at 7 days PI from all tissue samples, which did not correspond with the

findings of present study. The finding of the reisolation from liver, spleen and cecum with infection of *S. Pullorum* in the present study was similar to the finding with infection *S. Enteritidis* of Okamura et al, (2001). The findings of reisolation from liver, heart, crop and intestine with the infection of *S. Enteritidis* phage type of Kinde et al., (2000) did not correspond with the findings of present study. Liver and spleen were 100% positive at 1 wk PI, and 60% and 80% at 18 wk PI, and 37.5% and 12.5 % at 22 wk PI with the infection of *S. Pullorum* at 1 wk old male and female commercial brown –egg- laying chickens which agreed with the findings of Wigley et al, (2005). Wigley et al. (2005) also reported that the reisolation rate gradually decreased from 1 wk onwards while the reisolation had been found decreases from 2 wks onwards in the present study. They infected the birds at 0 days old but in the present study birds were infected at 21 wks of age for female and 29 wks of age for male. The findings in the present study, the reisolation of *Salmonella Pullorum* in blood culture was the highest at 1 wk PI ( $13.55 \times 10^3$  CFU) which was similar to the finding of Okamura et al, (2000).

All experimental hens showed histological lesions in different internal organs at 7 days PI. The histological lesions began from 1 wk PI and decreased gradually at 3 wks PI to onward. The highest lesion was found in spleen, followed by lungs and liver in the present study. In this investigation, hepatitis, typhilitis, bronchitis and pneumonia were recorded while Roy et al. (2001) also found similar result. In the present study, the histopathological lesions were hepatitis and infiltration of inflammatory cells in liver; multifocal necrotic foci in the liver; nodule formation in the liver; pneumonia and bronchopneumonia; nodule formation in the lungs; focal necrosis and inflammatory cells in the spleen; infiltration of inflammatory cells in the intestine; ulcer in the cecal tonsils; typhilitis and infiltration of inflammatory cells in ceca; congestion in the kidneys and infiltration of inflammatory cells in gizzard; congestion and infiltration of inflammatory cells in the ovary; infiltration of inflammatory cells in the oviduct. The above types of histological lesions with mild variation in severity and infiltration of inflammatory cells were supported for *Salmonella* infection by different investigators (Chauhan and Roy, 2007; Haider et al., 2008).

In this investigation, experimental infection by *S. Pullorum*, the immune response increased gradually from 1wk PI (titre 1211.82) and the highest titre (4712.39) was found at 4 wks PI which was also similar to the findings of Barrow et al. (1992) and was dissimilar to the findings of Hoop and Pospishil (1993). However, other investigators used other *Salmonella* organisms to determine the immune response by ELISA and RPA test. Their findings were slightly varied with the present findings due to the strain variation, management, age etc. Skov et al. (2002) was found have highest ELISA titre at 3 wks PI with the infection of *S. Typhimurium* in Chickens.

All *Salmonella* strains screened by PCR resulted in visualization of the predicted 457-bp amplified product in ethidium bromide-stained gels (Stone et al., 1994) and in the present study *invA* gene also visualized and produced 284-bp amplicon. In this investigation, *Salmonella Pullorum* was detected from tissue (liver, lung and ovarian follicle) samples by PCR with the amplification of *invA* gene at 12 hrs PI to onwards. Desai et al. (2005) and Olivera et al. (2003) also detected 284-bp amplicon from tissue samples in experimental infected chickens.

In present study in male, their feed intake and body weights were reduced, which was supported by the finding of Shivaprasad (1997). In the present study, pericarditis, haemorrhagic and congested liver, congested and caseous lungs, and white foci in the testis were observed in cock at 2 wks PI and histopathologically, degeneration and necrosis of spermatogonia, and infiltration of heterophils, lymphocytes and plasma cells in the somniferous tubules of testes were (75%) found. These gross and histopathological lesions corresponded with the findings of Chauhan and Roy (2007). In the present study, *S. Pullorum* was reisolated from 75% testes of cocks but Wigley et al. (2005) could not reisolate *S. Pullorum* in the testes of male birds.

### CONCLUSIONS

In this study, it is clear that after oral route of infection with infective dose of *S. Pullorum*, the bacteria invades digestive epithelia and ultimately enters into blood called bacteremia. From blood, bacteria are seeded into cells and tissues of different organs such as liver, lung, spleen, kidney, different parts of reproductive tracts of hens and testes of male and other tissues producing pathological lesions. It is also confirmed that the bacteria invade ovary and egg follicles, and this infection persists in ovary and egg follicles and transmits into laid eggs then to hatched chicks. No chronic lesion especially arthritis was recorded in this experiment. In this study pathogenesis and pathology are known in poultry birds of different age groups. In future for the control of *Salmonella* infections in poultry, vaccine production and sequencing of vaccine candidate in association with phylogenetic analysis of circulating *Salmonella* organisms should be performed in Bangladesh.

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**Table 1: Mean CFU/gm of isolated and identified of *Salmonella Pullorum* from different organs of experimentally infected hens**

Organs	BI 6 hr	PI 1 wk	PI 2 wk	PI 3 wk	PI 4 wk
1. Crop	00 (0/4)	58.72 x 10 <sup>4</sup> (4/4)	25.81 x 10 <sup>3</sup> (4/4)	25.68 x 10 <sup>2</sup> (3/4)	14.96x 10 <sup>1</sup> (2/4)
2. Liver	00 (0/4)	64.58 x 10 <sup>5</sup> (4/4)	65.79 x 10 <sup>4</sup> (4/4)	55.98 x 10 <sup>3</sup> (4/4)	44.28x 10 <sup>3</sup> (3/4)
3. Heart	00 (0/4)	21.34 x 10 <sup>3</sup> (4/4)	40.16 x 10 <sup>3</sup> (4/4)	48.66 x 10 <sup>2</sup> (3/4)	14.71x 10 <sup>2</sup> (3/4)
4. Lungs	00 (0/4)	44.85 x10 <sup>4</sup> (4/4)	39.02.2x10 <sup>4</sup> (4/4)	66.95 x10 <sup>3</sup> (4/4)	24.26 x10 <sup>2</sup> (4/4)
5. Duodenum	00 (0/4)	20.98 x10 <sup>5</sup> (4/4)	47.51x10 <sup>4</sup> (4/4)	36.54 x10 <sup>4</sup> (4/4)	43.56 x10 <sup>3</sup> (4/4)
6. Cecum	00 (0/4)	81.11 x 10 <sup>5</sup> (4/4)	73.93 x 10 <sup>5</sup> (4/4)	64.47 x 10 <sup>4</sup> (4/4)	80.77x 10 <sup>3</sup> (4/4)
7. Bile	00 (0/4)	47.88 x 10 <sup>3</sup> (2/4)	14.29x 10 <sup>3</sup> (1/4)	00 (0/4)	00 (0/4)
8. Kidney	00 (0/4)	39.72 x 10 <sup>3</sup> (4/4)	13.58 x 10 <sup>3</sup> (4/4)	18.84 x10 <sup>2</sup> (2/4)	19.26 x10 <sup>1</sup> (2/4)
9. Spleen	00 (0/4)	89.25 x 10 <sup>5</sup> (4/4)	93.34 x10 <sup>5</sup> (4/4)	62.21 x10 <sup>3</sup> (4/4)	51.25 x10 <sup>3</sup> (4/4)
10. Blood	00 (0/4)	13.55 x 10 <sup>3</sup> (4/4)	8.43 x 10 <sup>3</sup> (4/4)	33.76x 10 <sup>2</sup> (3/4)	13 x 10 <sup>2</sup> (1/4)

Note: Percentage calculated from 1 wk PI to 4 wks PI.

**Table 2: Mean CFU/gm of isolated and identified *Salmonella Pullorum* from female reproductive organs in experimentally infected hens**

Organs	BI 6 hr	PI 1 wk	PI 2 wk	PI 3 wk	PI 4 wk	Total (%)
1.Ovary	00 (0/4)	41.05 x 10 <sup>5</sup> (4/4)	64.6 x 10 <sup>4</sup> (4/4)	33.05 x 10 <sup>3</sup> (4/4)	40.93x 10 <sup>4</sup> (4/4)	100
2.Ovarian follicle	00 (0/4)	67.75 x 10 <sup>4</sup> (4/4)	47.61 x 10 <sup>4</sup> (4/4)	31.67 x 10 <sup>3</sup> (4/4)	49.5 x 10 <sup>3</sup> (4/4)	100
3. Oviduct	00 (0/4)	39.65 x 10 <sup>4</sup> (4/4)	25.17 x 10 <sup>3</sup> (3/4)	40.67 x 10 <sup>3</sup> (2/4)	27.54x 10 <sup>3</sup> (2/4)	68.75
4. Uterus	00 (0/4)	40.11 x10 <sup>5</sup> (3/4)	14.36x10 <sup>3</sup> (3/4)	19.16 x10 <sup>3</sup> (2/4)	30.81 x10 <sup>3</sup> (1/4)	56.25
5. Vagina	00 (0/4)	51.15 x 10 <sup>3</sup> (4/4)	07.2 x 10 <sup>3</sup> (4/4)	43.12 x 10 <sup>3</sup> (3/4)	13.13x 10 <sup>3</sup> (1/4)	75

Note: Percentage calculated from 1 wk PI to 4 wks PI

**Table 3: CFU/gm of isolated and identified *Salmonella Pullorum* from different organs of cocks after experimental infection**

Organs	BI 6 hr	PI 1 wk	PI 2 wk	PI 3 wk	PI 4 wk	Total (%)
1. Liver	00 (0/1)	69.83 x 10 <sup>3</sup> (1/1)	158.62 x 10 <sup>3</sup> (1/1)	37.05 x 10 <sup>3</sup> (1/1)	40.93x 10 <sup>2</sup> (1/1)	100
2. Lung	00 (0/1)	60.89x 10 <sup>3</sup> (1/1)	86.90 x 10 <sup>3</sup> (1/1)	31.67 x 10 <sup>3</sup> (1/1)	49.5 x 10 <sup>2</sup> (1/1)	100
3. Heart	00 (0/1)	12.24 x 10 <sup>3</sup> (1/1)	23.31 x 10 <sup>3</sup> (1/1)	41.67 x 10 <sup>2</sup> (1/1)	00 (0/1)	75
4. Spleen	00 (0/1)	105.41 x10 <sup>3</sup> (1/1)	163.36x10 <sup>3</sup> (1/1)	89.16 x10 <sup>3</sup> (1/1)	30.81 x10 <sup>2</sup> (1/1)	100
5. Cecum	00 (0/1)	236.88 x 10 <sup>3</sup> (1/1)	118.68 x 10 <sup>3</sup> (1/1)	103.12 x 10 <sup>3</sup> (1/1)	83.13x 10 <sup>3</sup> (1/1)	100
6. Testis	00 (0/1)	35 x10 <sup>2</sup> (1/1)	16.4 x 10 <sup>3</sup> (1/1)	19.32 x 10 <sup>2</sup> (1/1)	00 (0/1)	75

Note: Percentage calculated from 1 wk PI to 4 wks PI

**Table 4: Antibody titre in hens infected with *Salmonella Pullorum* detected by indirect ELISA**

Time schedule of PI	Infection group (Mean±SD)	Control group (Mean±SD)
BI 6 hr	137.37±103.66*	138.85±72.34
PI 1 wk	1211.82±585.07*	151.69±162.54
PI 2 wk	3189.90±1383.26*	154.24±129.93
PI 3 wk	4312.58±1476.23*	155.26±172.18
PI 4 wk	4712.39±1851.94*	171.29±169.68

\*The mean difference is significant at p<0.01

**Table 5: Antibody titre in cocks infected with *Salmonella Pullorum* detected by indirect ELISA**

Time schedule of PI	Infection group (Mean±SD)	Control group (Mean±SD)
BI 6 hr	182.24±56.54*	192.05±38.62
PI 1 wk	508.20±98.77*	202.74±45.75
PI 2 wk	1167.61±162.39*	193.87±143.72
PI 3 wk	2114.94±612.03*	189.99±51.63
PI 4 wk	3059.27±903.40*	172.56±200.45

\*The mean difference is significant at  $p < 0.01$

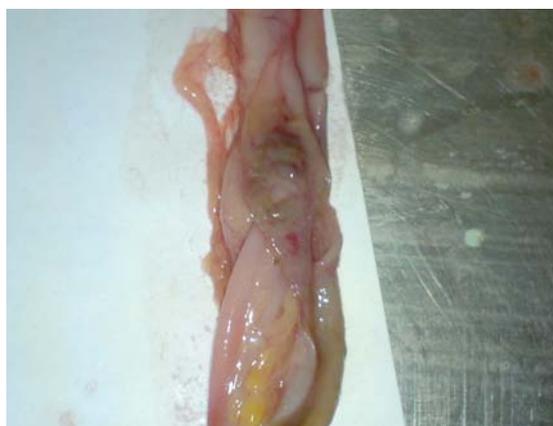


Figure 1: Semi-solid, cheesy material in ceca and button like ulcer at 3 wks PI with experimental infection of *S. Pullorum*.



Figure 2: Misshapen, discoloured, cystic and congested ova at 2 wks PI with experimental infection of *S. Pullorum*.

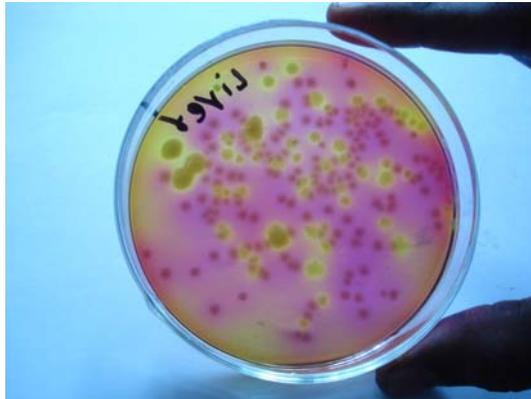


Figure 3: Experimentally infected with *S. Pullorum* at 1wk PI showing 76 CFU of *S. Pullorum*/gm of tissue of liver at  $10^5$  dilution.

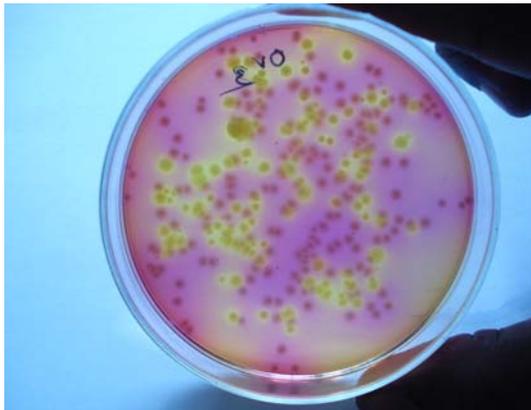


Figure 4: Experimentally infected with *S. Pullorum* at 1wk PI showing 86 CFU of *S. Pullorum*/gm of tissue of ovary at  $10^5$  dilution

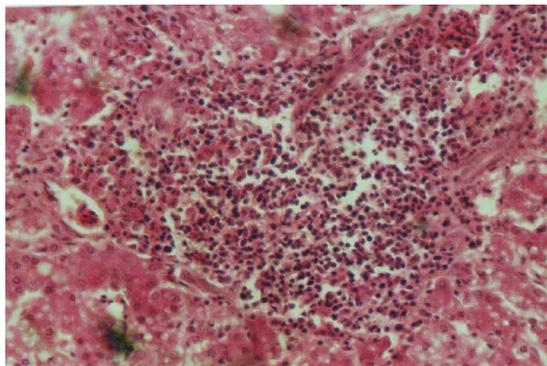


Figure 5: Nodule formation with the infiltration of heterophils, lymphocytes, macrophages and plasma cells in liver in experimental PD in hens at 3 wks PI (H&E, X 333).

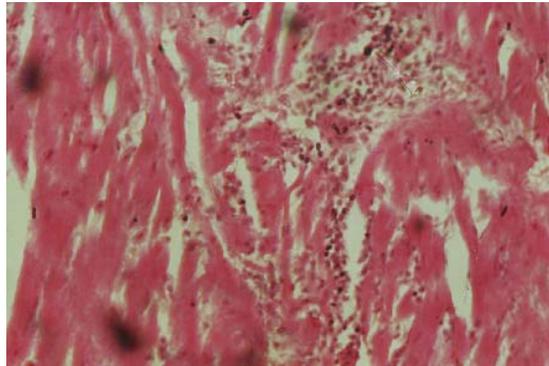


Figure 6: Nodule is formed with infiltration of heterophils macrophage and lymphocytes in heart in experimental PD in hens at 2 wks PI (H&E, X 333).

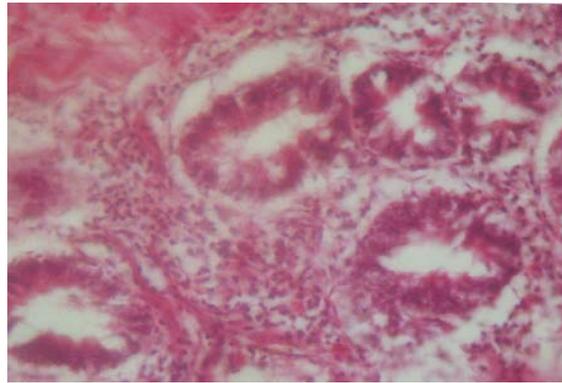


Figure 7: Infiltration of macrophages, heterophils and lymphocytes in the uterus and necrosis in the epithelial cells of the gland in experimental PD in hens at 2 wks PI (H&E, X 333).

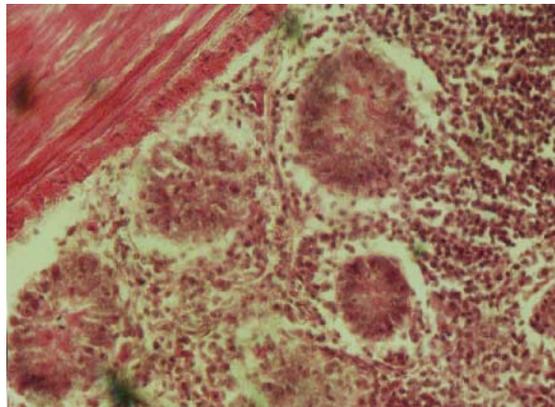


Figure 8: Degeneration and necrosis of spermatogonia and infiltration of heterophils, lymphocytes and plasma cells are found in the somniferous tubules of testes in experimental PD in cock at 3 wks PI (H&E, X 83).

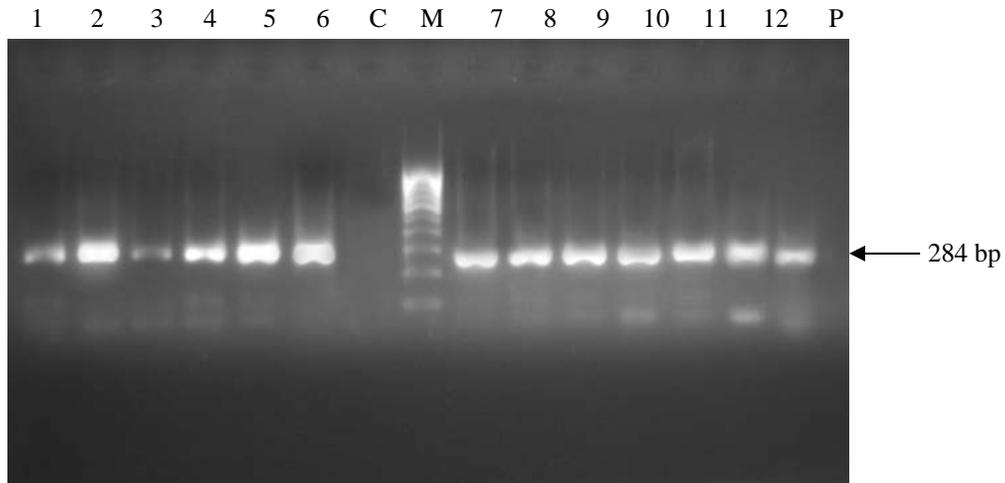


Figure 9: Electrophoresis for *Salmonella Pullorum* at 1wk of PI on 1.5% agarose gel showing band from lane 1 to 4 liver, from lane 5 to 8 lungs and from lane 9 to 12 ovarian follicle samples, lane p showing the 284-bp PCR products as a positive control and lane C showing no band as a negative control after amplification with the primer 139 (F) and 141 (R) targeting the gene *invA* for *Salmonella Pullorum* and lane M showing DNA molecular mass marker (100-bp ladder).

## SELECTION CRITERIA, YIELD RELATIONSHIP WITH COMPONENT TRAITS AND GROUPING OF TROPICAL JAPONICA, INDICA LINES AND DERIVED HYBRIDS OF RICE (*Oryza sativa* L.)

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### ABSTRACT

Forty-five rice lines comprising of thirty derived hybrid lines obtained from ten tropical *Japonica*, three *Indica* and two national checks viz. Pusa Basmati 1121 and Sarjoo-52 were evaluated for selection parameters, yield contributing components and genetic divergence. Fifteen quantitative and three qualitative traits were studied from experimentation with randomized block design during *Kharif* 2011. The phenotypic coefficient of variability was higher than genotypic coefficient of variability for all of the traits. The highest estimates of broad sense heritability coupled with genetic advance in per cent of mean was recorded for spikelets per panicle, plant height followed by L:B ratio, spikelets per panicle, grains per panicle, biological yield per plant, flag leaf area, days to 50% flowering, plant height which might be due to the additive nature of gene action. Such results indicated that these traits will be reliable for the effective selection. Highly positive and significant correlation was observed at both phenotypic and genotypic level between grain yield per plant and biological yield per plant, followed by panicle bearing tillers per plant, spikelet fertility, panicle length, 1000-grain weight, grains per panicle, panicle weight, flag leaf length, spikelet per panicle, flag leaf area, kernel length, flag leaf width, days to 50% flowering, and harvest index. This relationship reflected that grain yield and aforesaid economic traits can be increased simultaneously in breeding programme to develop high yielding *Indica* as well as Tropical *Japonica* rice varieties. Whole genotypes grouped in 8 non-overlapping clusters exhibited maximum genetic diversity between clusters III i.e., TJ-

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64897 × NDR-359, TJ-64897 × CSR36, TJ-64897 × PB-1 and VIII i.e., TJ-11010 × NDR359, TJ-11010 × PB-1, TJ-16081 × NDR-359, TJ-16081 × PB-1. These clusters also stand for early days to flowering, short slender, second highest harvest index and panicle bearing tillers per plant, spikelets per panicle, grains per panicle, spikelet fertility, 1000-grain weight, long bold slender, biological yield per plant, and grain yield per plant. These genotypes showing higher mean performance for aforesaid traits can be exploited for enhancing hybrid vigour of desired New Plant Type with higher number of panicle bearing tillers per plant, spikelet per panicle and grains per spike in *Indica* as well as Tropical *Japonica* rice varieties for achieving higher yield.

**Keywords:** Clustering pattern, Correlations, Genetic diversity, *Indica*, *Japonica*, New Plant Type, Path analysis and Selection parameters.

## INTRODUCTION

Rice as lifeline for a large segment of the world's population, more than 91% of world's rice is produced in Asia and provides 20% of the per capita energy and 13% of the per capita protein worldwide. However, in Asia rice contributes about 35% of the energy and 28% of the protein. In India it plays an integral role for 65 per cent population, food security, income generation, employments and foreign currencies gains for the nation. India has the largest area, about  $4.5 \times 10^7$  ha, under rice in the world and produced  $9.5 \times 10^9$  tone in year 2010-11. In accordance with current population growth, rice requirement by 2025 is estimated to be around  $1.3 \times 10^{10}$  tone (Anonymous 2010). Hence it is very crucial to develop high yielding varieties. It is cultivated on foot hills of the Himalaya in the North Western parts of Indian sub-continent comprising the states of Haryana, Punjab, Uttaranchal, Uttar Pradesh, Himachal Pradesh and Delhi. Current *Indica* rice varieties have higher yield potential of  $100 \text{ t ha}^{-1}$  with harvest index of 0.4-0.5 under tropical conditions.

The hybrid vigour between *Indica* and *Japonica* inter-subspecies can be used to improve the yield of *Indica* and *Japonica* hybrids by introducing parts of the *Indica* genomic segments into the *Japonica* background or vice versa, after several generations of backcrossing. For developing New Plant Type with increased vigour, *Indica* and *Japonica* materials are an effective source of breeding for the many pleasing and desirable traits such as less and productive tillers, canopy structure, increased biomass production, reduced carbon assimilation, increased harvest index, increased sink-source and spikelet fertility.

Crop improvements depend upon the magnitude of genetic variability present in the base population. The expected improvement in yield components primarily depends on the nature and magnitude of heritable total variation. Selection based on a single character may not always be effective while it is impractical for a researcher to consider a large number of component characters simultaneously in a particular selection procedure. Correlation analysis between yield and its components provides

useful information for right choice of characters in the selection programme. Path coefficient analysis provides an effective tool for partitioning the correlation coefficient in to direct and indirect effects of yield attributes on yield, with their cause and effect relationship.

The varieties originated from widely distinct localities or geographical areas were usually presumed to be diverse and were utilized in hybridization programmes because earlier workers regarded this geographical isolation as reasonable index of genetic diversity. However, several workers have found the lack of parallelism between geographical distribution and genetic diversity in different crop species and have also reported that genetic drift and selection in different environments could cause greater diversity than geographic distances. (Murty and Arunachalam, 1966; Bhanumathi et al., 2010; Manikya and Reddy, 2011). For improving genetic diversity of best suited germplasm across the location, it is very important to know the extent of existing genetic diversity. Genetic diversity is the heritable difference among germplasm, is crucial at an optimum level within a population to facilitate and sustain an effective, continuous breeding programme. Exotic introduced materials have specific genetic background and also may play a greater role to enhance heterosis.

### MATERIALS AND METHODS

The reported experiment included forty-five rice lines comprising of ten Tropical *Japonica* as lines and three *Indica as* testers and their thirty crosses along with two checks viz. Pusa Basmati 1121 and Sarjoo-52. The genotypes were evaluated in Randomized Block Design with three replications during *kharif* 2011. The crosses were made during *kharif* 2010 at Crop Research Farm, Nawabganj of C.S. Azad University of agriculture and Technology, Kanpur, situated between 27.24° N latitude, 77.5° E longitudes and at an altitude of 178 meters above the msl. in the gangetic plain of central Uttar Pradesh. The climate of district Kanpur is semi-arid with hot summer and cold winter. Nearly 80 percent of total rainfall occurs during the monsoon (only up to September) with a few showers in the winter. The site of experiment was salt affected clay loam (natrustalf) soil having pH = 10.5, EC =2.03 and low in organic carbon, nitrogen and phosphorus.

Observations were recorded on ten competitive plants randomly selected in each replications for fifteen quantitative and three qualitative traits viz., days to 50% flowering, plant height (cm), panicle bearing tillers per plant, flag leaf length (cm), flag leaf width (cm), flag leaf area (cm<sup>2</sup>), panicle length (cm), panicle weight (g), spikelets per panicle, grains per panicle, spikelet fertility (%), 1000-grains weight (g), kernel length (mm), kernel breadth (mm), L:B ratio, biological yield per plant (g), harvest index (%) and grain yield plant (g). Each genotype was accommodated in three-row of 5 m length following row to row and plant to plant spacing of 20 × 15 cm, respectively. The recommended dose of fertilizers N: P: K @ 120:60:60 kg ha<sup>-1</sup>, cultural packages and practices were followed to raise a good healthy crop.

Standard statistical techniques such as analysis of variance (Panse and Sukhatme, 1967), genotypic and phenotypic coefficients of variation (Burton and de Vane, 1953), estimate of broad sense heritability ( $h^2_b$ ) (Hanson et al., 1956), genetic advance as percent of the mean (Johnson et al., 1955), phenotypic and genotypic correlation coefficient (Searle 1961), path analysis (Dewey and Lu, 1959) and standard statistical techniques such as Non-hierarchical Euclidean cluster analysis (Beale, 1969; Spark, 1973) were applied for assessing variability and genetic divergence present in the germplasm (using MASTA-C software). Grouping and arrangement of genotypes were done by Tocher's method (Rao, 1952).

## RESULTS AND DISCUSSION

### Genetic variability parameters

The analysis of variance revealed presence of highly significant differences among all the genotypes for eighteen characters and indicated considerable amount of variability in the genotypes. Significant variability due to treatment for all the characters (Table 1) was also confirmed by Jayasudha and Sharma (2010). Phenotypic coefficient of variability (PCV) was higher than genotypic coefficient of variability (GCV) (Table 2). A perusal of coefficient of variability indicates that PCV and GCV were quite high for flag leaf area (47.79 & 47.06), panicle bearing tillers per plant (46.94 & 44.65), grains per panicle (45.92 & 45.14), grain yield per plant (44.32 & 43.07), biological yield per plant (40.73 & 39.22), spikelets per panicle (38.88 & 38.33), panicle weight (38.71 & 38.22), flag leaf width (28.86 & 28.36), and flag leaf length (25.16 & 24.26). Moderate PCV and GCV were recorded in panicle length (19.1 & 18.35), L:B ratio (17.18 & 16.56), 1000-grain weight (16.05 & 15.89), kernel breadth (14.82 & 14.21), days to 50% flowering (14.82 & 14.14), spikelet fertility (12.09 & 8.94), plant height (11.68 & 11.29) and kernel length (10.95 & 10.69) by Seyoum et al., 2012, Yadav et al., 2011, Manikaya and Reddy, 2011 and Barber et al., 2009. The highest estimates of broad sense heritability coupled with genetic advance in per cent of mean was recorded for spikelets per panicle (96.60 & 149.86), plant height (98.9% & 65.13%) followed by L:B ratio (99.99 & 39.30), spikelets per panicle (97.20 & 32.39), grains per panicle (96.60 & 141.11), biological yield per plant (92.70 & 42.47), flag leaf area (97.00 & 40.47), days to 50% flowering (91.10 & 26.75), and plant height (93.40 & 25.04). Due to the additive nature of gene action these traits will be reliable for the effective selection. High heritability coupled with moderate genetic advance for grain yield per plant (94.40 & 15.22) and flag leaf length (93.00 & 13.62) indicated preponderance of non-additive gene action, hence selection cannot be rewarded which was also agreed with by Seyoum et al., 2012; Yadav et al., 2011 and Suman et al., 2005. The most preferable genotypes having high variability, heritability and genetic advance for characters other than grain yield per plant can also be used as donor parents in hybridization programme to improve desirable characters for which they showed high value of heritability and genetic advance.

### Relationship analysis

The genotypic correlation coefficient was higher in magnitude than the corresponding phenotypic correlation coefficient in general for most of the characters (Table 3). This is possibly due to the linkage or modifying effect of the gene and environment in genetic association between characters (Swain and Reddy 2006). There were strong positive and significant correlation at both phenotypic and genotypic level, between grain yield per plant and biological yield per plant (0.931 & 0.884) followed by panicle bearing tillers per plant (0.823 & 0.593), spikelet fertility (0.823 & 0.593), panicle length (0.809 & 0.752), 1000-grain weight (0.738 & 0.710), grains per panicle (0.699 & 0.666), panicle weight (0.683 & 0.660), flag leaf length (0.678 & 0.625), spikelet per panicle (0.633 & 0.604), flag leaf area (0.628 & 0.599), kernel length (0.579 & 0.549), flag leaf width (0.516 & 0.499), days to 50% flowering (0.385 & 0.351), and harvest index (0.368 & 0.366). This relationship reflected that grain yield and aforesaid traits can be improved simultaneously (Jaisudha and Sharma, 2010; Manikaya and Reddy, 2011 and Seyoum et al., 2012). Besides, biological yield was positively and highly significantly correlated with panicle bearing tillers per plant (0.881 & 0.821), grains per panicle (0.794 & 0.756), spikelet fertility (0.863 & 0.620) and panicle length (0.828 & 0.771) and 1000-grain weight (0.764 & 0.730). This indicates that vigorous plant population may enhance economic yield. 1000-grain weight was significantly and positively associated with spikelet fertility and panicle length and other yield contributing traits. A highly significant and positive correlation was noted between grains per panicle and panicle length (0.679 & 0.639), panicle bearing tillers per plant (0.597 & 0.570) and panicle weight (0.843 & 0.816). Panicle weight showed highly significant and positive association with panicle per plant (0.528 & 0.501) and panicle length (0.772 & 0.736). Such type of results were reported by Petchiammal and Khan, 2007. This association reflected that above character can be considered as new plant type for enhancing the economic yield.

### Path coefficient analysis

The highest direct effect of grains per panicle (1.208 & 0.235) was noted over grain yield per plant, followed by flag leaf width (1.197 & 0.477), biological yield per plant (1.132 & -0.340), harvest index (0.854 & 0.725), kernel breadth (0.825 & -0.330), flag leaf length (0.818 & 0.373) on grain yield via panicle weight (1.018 & 0.192), spikelets per panicle (1.195 & 0.230), spikelet fertility (1.013 & 0.162), biological yield per plant (0.959 & 0.178), 1000-grain weight (0.852 & 0.160), panicle length (0.820 & 0.150), flag leaf area (0.787 & 0.150), flag leaf length (0.724 & 0.135), flag leaf width (0.723 & 0.137) and panicle bearing tillers per plant (0.721 & 0.134) (Table 4). Almost similar relationships were also reported by Borbora et al., 2005; Veni and Rani, 2007; and Kisore et al., 2007. Therefore, these traits can be used for selection criteria in breeding programme to develop high yielding new plant type rice varieties.

### Clustering pattern and genetic divergence analysis

Non-hierarchical Euclidean cluster analysis grouped the forty-five aforesaid genotypes, into 8 non-overlapping clusters (Table 5 and 6). Cluster II had the maximum 13 genotypes, followed by cluster VIII, I, and cluster III, IV, V, VI which had same number of 4 genotypes. Cluster VII comprised of 3 genotypes which reflected narrow genetic diversity (Dushyantha and Kantti 2010).

Maximum inter-cluster diversity was observed between III & VIII, followed by cluster III & IV, III & VII, II & VIII, III & VI, II & IV, IV & V, I & VIII and I & IV. Similar inter-cluster distances were reported by Vaithiyalingan et al., 2007. The genotypes grouped in cluster I consisted of eleven genotypes, only three genotypes i.e. TJ-25966 × NDR-359, TJ-25966 × CSR36, TJ-25966 × PB-1 showed short bold grain and harvest index. It is big information that genotype TJ-64897 × NDR-359, TJ-64897 × CSR36, TJ-64897 × PB-1 from cluster III and genotype TJ-11010 × NDR359, TJ-11010 × PB-1, TJ-16081 × NDR-359, TJ-16081 × PB-1 from cluster VIII expressed maximum genetic diversity and also stands for early days to flowering, short slender, second highest harvest index and panicle bearing tillers per plant, spikelets per panicle, grains per panicle, spikelet fertility, 1000-grain weight, biological yield per plant, grain yield per plant. Chandra et al., 2007 also observed similar results in their study. Therefore, it is suggested that any superior genotype of cluster III may be crossed with any superior genotypes of cluster VII to produce desirable recombinants with more suitable traits for higher yield.

Maximum genetic divergence within the cluster was estimated for cluster V (3.058) followed by cluster II (2.205), VII (1.995), I (1.892), VII (1.378) and VI (1.288) while cluster IV (1.140) showed minimum intra cluster distance. Almost similar inter-cluster distance was estimated by Suman et al., 2005 in rice. The maximum intra cluster value indicated maximum divergence among various genotypes within the cluster whereas; minimum value reflected minimum diversity among the genotypes within the respective cluster.

Based on cluster mean performance (Table-7), cluster VII showed the highest cluster mean for productive tillers per plant (17.73), flag leaf length (35.91), panicle length (31.23), grains per panicle (245.04), 1000-grain weight (32.54), biological yield per plant (85.02), grain yield per plant (28.24). Gahalain et al., 2006 reported similar type of cluster mean in rice.. From this cluster, the genotypes TJ-11010 × NDR359, TJ-11010 × PB-1, TJ-16081 × NDR-359, TJ-16081 × PB-1 expressed superiority over the checks for aforesaid respective characters. Cluster IV showed maximum mean performance for genotypes TJ-10365 × NDR-359 and TJ-10365 × PB-1 for flag leaf width (2.87), flag leaf area (71.97) and spikelet per panicle (284.33). Cluster VII showed third highest mean performance for days to 50% flowering (87.78), panicle bearing tillers per plant (15.11), panicle length (30.10), spikelet per panicle (275.22), grains per panicle (238.00) and 1000-grain weight (31.00). The highest mean performance for panicle weight (4.16), kernel breadth

(3.25) were recorded in the two most superior genotypes TJ-25892 × NDR359 and TJ-25892 × PB-1 of cluster VII. Earliness (87.78 days) and desirable medium plant height (116.00) of the above mentioned genotypes can be taken under consideration in crossing program. On the basis of eight genetically diverse clusters, superior genotypes listed in table 8 from the different clusters were selected. Interacting from different clusters and diverse genotypes would lead greater opportunity for crossing over which release latent genetic variability by breaking linkage drag and progenies derived from cross are expected to show broad spectrum of genetic variability providing a greater scope for screening transgressive segregants as new plant type for sustaining yield of existing varieties.

### CONCLUSIONS

The study of fifteen quantitative and three qualitative traits ranging for trait to trait among the genotype, indicated that selection can be practiced for background selection to sort-out superior parent with higher spikelets per panicle, grains per panicle, and panicle bearing tillers per plant due to the additive nature of gene action. The direct yield contributing characters showing strong positive association with grain yield reflected that these characters can be enhanced simultaneously. Similarly genotypes falling under cluster III and VIII having high mean performance for respective character can be used for getting better combinations with desirable character in *Indica* as well as in *Japonica* rice varieties for foreground selection to achieve higher yield with desirable new plant type.

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**Table 1: Estimates of variance for 18 characters in *Indica* and Tropical *Japonica* rice lines and their hybrids**

Sl. No.	Characters	Sources of variation				
		Replication (2)	Treatment (44)	Error (88)	CV%	CD
1.	Days to 50% flowering	1.50	573.36**	18.10	4.42	6.89
2.	Plant height (cm)	42.93	485.82**	11.22	3.00	5.42
3.	Panicle bearing tillers/plant	2.61	73.62**	2.49	14.48	2.56
4.	Flag leaf length	2.16	144.59**	3.56	6.67	3.06
5.	Flag leaf Breadth	0.03	0.89**	0.01	5.36	0.16
6.	Flag leaf area	3.41	1206.68**	12.45	8.32	5.72
7.	Panicle length (cm)	1.04	70.18**	1.88	5.28	2.23
8.	Panicle weight (g)	0.02	3.10**	0.02	6.15	0.26
9.	Spikelets/panicle	413.75	16496.41**	158.16	6.53	20.38
10.	Grains/spike	178.37	14739.39**	169.98	8.44	21.15
11.	Spikelet fertility (%)	17.59	187.43**	40.54	8.13	10.31
12.	1000- grain weight (g)	1.96	56.87**	0.39	2.29	1.01
13.	Kernel length	0.01	1.42**	0.02	2.34	0.24
14.	Kernel breadth	0.03	0.35**	0.01	4.22	0.16
15.	L:B ratio	0.02	0.62**	0.01	4.57	0.19
16.	Biological yield/plant (g)	138.51	1410.99**	35.88	10.97	9.72
17.	Harvest index	124.69	106.30**	21.90	14.37	7.59
18.	Grain yield/plant (g)	2.71	176.87**	3.41	10.46	2.98

\*, \*\* Significant at 5% and 1% probability levels, respectively

**Table 2: Estimates of general mean, variability parameters for 18 characters in *Indica* and Tropical *Japonica* rice lines and their hybrids**

Characters	Range		G. mean ±SE(m)	Coefficient of variation (%)			h <sup>2</sup> b (%)	G a (%)
	Minimum	Maximum		PCV	GCV	ECV		
Days to 50% flowering	66.25	114.00	96.19 ± 3.47	14.82	14.14	0.68	91.10	26.75
Plant height (cm)	94.00	143.33	111.41 ± 2.73	11.68	11.29	0.39	93.40	25.04
Panicle bearing tillers/plant	4.65	22.67	10.90 ± 1.29	46.94	44.65	2.29	90.50	9.54
Flag leaf length	17.15	41.70	28.26 ± 1.54	25.16	24.26	0.9	93.00	13.62
Flag leaf Breadth	0.84	2.92	1.91 ± 0.08	28.86	28.36	0.5	96.00	1.10
Flag leaf area	11.06	86.56	42.39 ± 2.88	47.79	47.06	0.73	97.00	40.47
Panicle length (cm)	18.57	34.23	25.99 ± 0.11	19.10	18.35	0.75	92.30	9.44
Panicle weight (g)	1.35	4.37	2.64 ± 0.13	38.71	38.22	0.49	97.50	2.06
Spikelets/panicle	90.67	328.67	192.55 ± 10.26	38.88	38.33	0.55	97.20	149.86
Grains/spike	65.11	292.67	154.38 ± 10.64	45.92	45.14	0.78	96.60	141.11
Spikelet fertility (%)	64.97	89.09	78.25 ± 5.19	12.09	8.94	3.15	54.70	10.66
1000- grain weight (g)	18.29	34.38	27.30 ± 0.51	16.05	15.89	0.16	98.0	8.85
Kernel length	4.77	7.44	6.39 ± 0.12	10.95	10.69	0.26	95.40	1.38
Kernel breadth	1.71	3.56	2.38 ± 0.08	14.82	14.21	0.61	91.90	0.67
L:B ratio	1.93	3.97	2.73 ± 0.10	17.18	16.56	0.62	92.90	0.90
Biological yield/plant (g)	18.27	110.73	54.59 ± 4.89	40.73	39.22	1.51	92.70	42.47
Harvest index	20.86	50.28	32.56 ± 3.82	21.72	16.29	5.43	56.20	8.19
Grain yield/plant (g)	5.59	34.98	17.65 ± 1.50	44.32	43.07	1.25	94.40	15.22

**Table 3: Genotypic and Phenotypic correlation coefficients for 18 characters in *Indica* and Tropical *Japonica* rice lines and their hybrids**

Traits	Level	PH	PBT/P	FLL	FLW	FLA	PL	PW	S/P	G/S	SF	TW	KL	KB	L:B	BY/P	HI	GY/P
DF	rg	0.515**	0.309*	0.647**	0.661**	0.697**	0.518**	0.358**	0.398**	0.430**	0.457**	0.414**	0.618**	-0.049	0.387**	0.571**	-0.462	0.385**
	rp	0.468**	0.298*	0.598**	0.621**	0.658**	0.476**	0.341*	0.370**	0.398**	0.316*	0.394**	0.587**	-0.042	0.363**	0.525**	-0.332	0.351**
PH	rg	1.000	0.101	0.412**	0.654**	0.608**	0.586**	0.708**	0.555**	0.568**	0.445**	0.676**	0.393**	0.236	0.000	0.461**	-0.128	0.349*
	rp	1.000	0.086	0.382**	0.624**	0.579**	0.532**	0.672**	0.524**	0.536**	0.322*	0.642**	0.377**	0.215	0.008	0.426**	-0.108	0.325*
PBT/P	rg		1.000	0.550**	0.239**	0.412**	0.697**	0.528**	0.534**	0.597**	0.741**	0.579**	0.639**	0.250	0.210	0.881**	0.204	0.880**
	rp		1.000	0.515**	0.220**	0.390**	0.638**	0.501**	0.506**	0.570**	0.543**	0.547**	0.595**	0.225	0.189	0.821**	0.138	0.817**
FLL	rg			1.000	0.711**	0.906**	0.708**	0.481**	0.589**	0.599**	0.607**	0.505**	0.642**	-0.004	0.370**	0.759**	-0.087	0.678**
	rp			1.000	0.665	0.892**	0.655**	0.455**	0.563**	0.573**	0.443**	0.486**	0.616**	0.009	0.340	0.711**	-0.105	0.625**
FLW	rg				1.000	0.924**	0.650**	0.665**	0.578**	0.599**	0.555**	0.577**	0.325*	0.226	-0.044	0.628**	-0.194	0.516**
	rp				1.000	0.911	0.608**	0.641**	0.561**	0.581**	0.410**	0.556**	0.302*	0.216	-0.048	0.596**	-0.123	0.499**
FLA	rg					1.000	0.728**	0.620**	0.632**	0.651**	0.622**	0.621**	0.521**	0.076	0.211	0.732**	-0.129	0.628**
	rp					1.000	0.686**	0.599**	0.616**	0.635**	0.462**	0.605**	0.501**	0.080	0.194	0.698**	-0.103	0.599**
PL	rg						1.000	0.772**	0.599**	0.679**	0.855**	0.691**	0.689**	0.238	0.249	0.828**	0.182	0.809**
	rp						1.000	0.736**	0.569**	0.639**	0.606**	0.663**	0.640**	0.210	0.237	0.771**	0.114	0.752**
PW	rg							1.000	0.806**	0.843**	0.759**	0.801**	0.402**	0.528**	-0.181	0.760**	0.034	0.683**
	rp							1.000	0.783**	0.816**	0.558**	0.781**	0.388**	0.490**	-0.164	0.724**	0.028	0.660**
S/P	rg								1.000	0.989**	0.760**	0.641**	0.489**	0.333*	0.014	0.733**	-0.088	0.633**
	rp								1.000	0.977**	0.534**	0.622**	0.471**	0.310*	0.015	0.700**	-0.086	0.604**
G/S	rg									1.000	0.839**	0.705**	0.545**	0.350*	0.038	0.794**	-0.059	0.699**
	rp									1.000	0.686**	0.682**	0.522**	0.326*	0.036	0.756**	-0.055	0.666**
SF	rg										1.000	0.701**	0.651**	0.272*	0.198	0.863**	0.073	0.823**
	rp										1.000	0.511**	0.463**	0.191	0.136	0.620**	0.079	0.593**
TW	rg											1.000	0.520**	0.412**	-0.069	0.764**	0.121	0.738**
	rp											1.000	0.504**	0.391**	-0.066	0.730**	0.092	0.710**
KL	rg												1.000	0.137	0.537**	0.701**	-0.216	0.579**
	rp												1.000	0.126	0.535**	0.656**	-0.159	0.549**
KB	rg													1.000	-0.745	0.308**	-0.276	0.197
	rp													1.000	-0.751	0.284*	-0.188	0.188
L:B	rg														1.000	0.164	0.109	0.178
	rp														1.000	0.152	0.065	0.160
BY/P	rg															1.000	0.021	0.931**
	rp															1.000	-0.068	0.884**
HI	rg																1.000	0.368**
	rp																1.000	0.366**
GY/P	rg																	1.000
	rp																	1.000

\*, \*\* Significant at 5% and 1% probability levels, respectively. Legents: DF= .....GY/P

**Table 4: Genotypic and Phenotypic path of 18 traits on grain yield in *Indica* and Tropical *Japonica* rice lines and their hybrids**

Traits	Level	DF	PH	PBT/P	FLL	FLW	FLA	PL	PW	S/P	G/P	SF	TW	KL	KB	BY/P	HI	GY/P	Corrl. of GY/P
DF	Pg	<b>-0.141</b>	0.063	0.042	0.529	0.791	-1.235	0.097	-0.272	-0.281	0.519	-0.155	0.194	-0.548	-0.041	-0.438	0.488	-0.105	0.385
	Pp	<b>-0.040</b>	-0.028	0.020	0.223	0.294	-0.473	0.003	0.001	-0.065	0.094	-0.021	0.033	0.176	0.014	-0.123	0.381	-0.137	0.351
PH	Pg	-0.073	<b>0.123</b>	0.014	0.337	0.783	-0.078	0.110	-0.537	-0.391	0.686	-0.151	0.316	-0.349	0.195	0.000	0.394	-0.029	0.349
	Pp	-0.019	<b>-0.059</b>	0.006	0.143	0.295	-0.416	0.003	0.002	-0.092	0.126	-0.021	0.054	0.113	-0.071	-0.003	0.309	-0.045	0.325
PBT/P	Pg	-0.044	0.012	<b>0.135</b>	0.449	0.287	-0.731	0.131	-0.401	-0.377	0.721	-0.251	0.271	-0.566	0.206	0.238	0.752	0.046	0.880
	Pp	-0.012	-0.005	<b>0.066</b>	0.192	0.104	-0.281	0.003	0.002	-0.089	0.134	-0.035	0.046	0.178	-0.074	-0.064	0.595	0.057	0.817
FLL	Pg	-0.091	0.050	0.074	<b>0.818</b>	0.850	-1.606	0.133	-0.365	-0.416	0.724	-0.206	0.236	-0.569	-0.003	0.419	0.648	-0.020	0.678
	Pp	-0.024	-0.023	0.034	<b>0.373</b>	0.315	-0.641	0.004	0.002	-0.099	0.135	-0.029	0.041	0.184	-0.003	-0.116	0.515	-0.043	0.625
FLW	Pg	-0.093	0.080	0.032	0.581	<b>1.197</b>	-1.638	0.122	-0.504	-0.408	0.723	-0.188	0.270	-0.288	0.186	-0.050	0.536	-0.044	0.516
	Pp	-0.025	-0.037	0.015	0.248	<b>0.474</b>	-0.655	0.003	0.002	-0.099	0.137	-0.027	0.047	0.090	-0.071	0.016	0.432	-0.051	0.499
FLA	Pg	-0.098	0.075	0.056	0.741	1.106	<b>-1.773</b>	0.137	-0.470	-0.446	0.787	-0.211	0.291	-0.462	0.062	0.239	0.625	-0.029	0.628
	Pp	-0.026	-0.034	0.026	0.333	0.432	<b>-0.719</b>	0.004	0.002	-0.109	0.150	-0.030	0.051	0.150	-0.027	-0.066	0.506	-0.042	0.599
PL	Pg	-0.073	0.072	0.094	0.579	0.778	-1.290	<b>0.188</b>	-0.586	-0.422	0.820	-0.289	0.323	-0.611	0.197	0.282	0.707	0.041	0.809
	Pp	-0.019	-0.031	0.042	0.244	0.288	-0.493	<b>0.005</b>	0.003	-0.100	0.150	-0.039	0.056	0.192	-0.069	-0.081	0.559	0.047	0.752
PW	Pg	-0.050	0.087	0.071	0.393	0.795	-1.099	0.145	<b>-0.759</b>	-0.569	1.018	-0.257	0.375	-0.357	0.436	-0.205	0.649	0.008	0.683
	Pp	-0.014	-0.040	0.033	0.170	0.304	-0.431	0.004	<b>0.004</b>	-0.138	0.192	-0.036	0.065	0.116	-0.162	0.056	0.525	0.011	0.660
S/P	Pg	-0.056	0.068	0.072	0.482	0.692	-1.120	0.112	-0.612	<b>-0.706</b>	1.195	-0.257	0.300	-0.434	0.275	0.016	0.626	-0.020	0.633
	Pp	-0.015	-0.031	0.033	0.210	0.266	-0.443	0.003	0.003	<b>-0.177</b>	0.230	-0.035	0.052	0.141	-0.102	-0.005	0.508	-0.035	0.604
G/S	Pg	-0.061	0.070	0.081	0.490	0.717	-1.155	0.127	-0.639	-0.698	<b>1.208</b>	-0.284	0.330	-0.483	0.289	0.043	0.678	-0.013	0.699
	Pp	-0.016	-0.032	0.038	0.214	0.275	-0.457	0.003	0.003	-0.173	<b>0.235</b>	-0.045	0.057	0.156	-0.108	-0.012	0.548	-0.023	0.666
SF	Pg	-0.064	0.055	0.100	0.497	0.664	-1.104	0.160	-0.576	-0.536	1.013	<b>-0.339</b>	0.328	-0.578	0.225	0.224	0.728	0.017	0.823
	Pp	-0.013	-0.019	0.036	0.165	0.194	-0.332	0.003	0.002	-0.094	0.162	<b>-0.065</b>	0.043	0.139	-0.063	-0.046	0.450	0.033	0.593
TW	Pg	-0.058	0.083	0.078	0.413	0.691	-1.102	0.130	-0.608	-0.453	0.852	-0.237	<b>0.468</b>	-0.461	0.340	-0.078	0.653	0.027	0.738
	Pp	-0.016	-0.038	0.036	0.181	0.263	-0.435	0.004	0.003	-0.110	0.160	-0.033	<b>0.084</b>	0.151	-0.129	0.022	0.529	0.038	0.710
KL	Pg	-0.087	0.048	0.086	0.525	0.389	-0.924	0.129	-0.305	-0.345	0.658	-0.220	0.243	<b>-0.887</b>	0.113	0.608	0.598	-0.049	0.579
	Pp	-0.024	-0.022	0.039	0.230	0.143	-0.360	0.003	0.001	-0.083	0.123	-0.030	0.042	<b>0.300</b>	-0.042	-0.182	0.476	-0.065	0.549
KB	Pg	0.007	0.029	0.034	-0.003	0.270	-0.134	0.045	-0.401	-0.235	0.423	-0.092	0.193	-0.121	<b>0.825</b>	-0.843	0.263	-0.063	0.197
	Pp	0.002	-0.013	0.015	0.003	0.102	-0.058	0.001	0.002	-0.055	0.077	-0.012	0.033	0.038	<b>-0.330</b>	0.255	0.206	-0.077	0.188
L:B	Pg	-0.055	0.002	0.028	0.303	-0.053	-0.374	0.047	0.137	-0.010	0.046	-0.067	-0.032	-0.476	-0.615	<b>1.132</b>	0.140	0.025	0.178
	Pp	-0.015	0.001	0.013	0.127	-0.023	-0.139	0.001	-0.001	-0.003	0.008	-0.009	-0.006	0.160	0.248	<b>-0.340</b>	0.110	0.027	0.160
BY/P	Pg	-0.081	0.057	0.119	0.621	0.752	-1.299	0.155	-0.577	-0.518	0.959	-0.292	0.358	-0.621	0.254	0.185	<b>0.854</b>	0.005	0.931
	Pp	-0.021	-0.025	0.054	0.265	0.282	-0.502	0.004	0.003	-0.124	0.178	-0.040	0.061	0.196	-0.094	-0.052	<b>0.725</b>	-0.028	0.884
HI	Pg	0.065	-0.016	0.028	-0.071	-0.232	0.229	0.034	-0.026	0.062	-0.072	-0.025	0.056	0.192	-0.228	0.124	0.018	<b>0.227</b>	0.368
	Pp	0.013	0.006	0.009	-0.039	-0.058	0.074	0.001	0.000	0.015	-0.013	-0.005	0.008	-0.048	0.062	-0.022	-0.049	<b>0.411</b>	0.366

Residual factors: Genotypic= 0.0072 and phenotypic= 0.0234, Bold figures Indicate direct effects. Legends: DF= PH= .....GY/P



**Table 7: Clusters means, standard deviation and coefficient of variation for 18 traits in *Indica* and *Tropical Japonica* rice lines and hybrids**

Character Clusters	DF	PH	PBT/P	FLL	FLW	FLA	PL	PW	S/P	G/P	SF	TW	KL	KB	L:B	BY/P	HI	GY/P	
	Mean	82.80	106.40	8.73	23.75	2.08	37.31	24.51	3.02	158.87	118.13	73.86	25.42	5.23	2.65	1.97	42.66	36.46	15.48
Cluster I	SD	8.25	2.77	2.24	1.86	0.30	7.38	2.04	0.31	51.01	43.88	8.73	1.37	0.38	0.13	0.05	1.85	3.80	1.57
	CV	9.96	2.60	25.66	7.83	14.42	19.78	8.32	10.26	32.11	37.15	11.82	5.39	7.27	4.91	2.54	4.34	10.42	10.14
	Mean	95.79	105.21	6.98	24.68	1.67	31.04	20.92	1.64	156.40	113.50	72.61	23.57	6.17	2.36	2.62	37.73	27.90	10.46
Cluster II	SD	8.74	8.08	1.56	5.09	0.17	7.18	2.24	0.23	58.66	42.70	5.15	2.10	0.44	0.18	0.24	6.91	4.33	2.48
	CV	9.12	7.68	22.35	20.62	10.18	23.13	10.71	14.02	37.51	37.62	7.09	8.91	7.13	7.63	9.16	18.31	15.52	23.71
	Mean	73.08	103.17	7.18	17.99	0.90	12.10	20.32	1.63	117.50	81.50	69.38	24.78	5.65	1.89	3.02	25.13	39.51	9.77
Cluster III	SD	3.98	5.71	0.19	0.72	0.05	0.84	1.01	0.12	9.48	9.37	3.14	0.92	0.08	0.21	0.32	0.97	3.0	0.47
	CV	5.45	5.53	2.65	4.00	5.56	6.94	4.97	7.36	8.07	11.50	4.53	3.71	1.42	11.11	10.60	3.86	7.59	4.81
	Mean	109.08	140.67	7.30	33.44	2.87	71.97	30.66	4.13	284.33	240.00	84.36	32.16	6.67	2.30	2.90	65.11	32.80	21.22
Cluster IV	SD	5.35	3.03	1.87	1.94	0.08	0.78	1.43	0.25	5.98	13.93	3.60	1.53	0.32	0.15	0.13	5.85	1.73	1.59
	CV	4.90	2.15	25.62	5.80	2.79	1.08	4.66	6.05	2.10	5.80	4.27	4.76	4.80	6.52	4.48	8.98	5.27	7.49
	Mean	92.31	95.26	15.50	30.23	1.49	33.74	28.79	2.02	143.56	118.59	82.63	23.81	6.92	2.05	3.44	61.54	39.64	24.54
Cluster V	SD	17.89	0.83	4.79	4.49	0.20	7.84	1.78	0.20	11.97	12.41	2.29	5.58	0.37	0.27	0.61	14.28	9.75	9.11
	CV	19.38	0.87	30.90	14.85	13.42	23.24	6.18	9.90	8.34	10.46	2.77	23.44	5.35	13.17	17.73	23.20	24.60	37.12
	Mean	110.58	122.25	12.32	34.85	2.15	56.24	29.04	2.71	142.58	111.58	77.62	29.79	6.70	2.31	2.90	65.80	29.17	19.05
Cluster VI	SD	1.20	1.19	1.45	3.24	0.12	7.01	1.32	0.30	21.95	25.85	7.88	0.87	0.07	0.02	0.03	6.85	2.51	2.34
	CV	0.11	0.10	1.18	0.93	0.56	1.25	0.45	1.11	1.54	2.32	1.02	0.29	0.10	0.09	0.10	1.04	0.86	1.23
	Mean	87.78	116.00	15.11	26.32	1.65	32.48	30.10	4.16	275.22	238.00	86.46	31.00	6.96	3.25	2.15	67.44	28.34	18.95
Cluster VII	SD	1.17	3.53	0.69	3.63	0.03	3.91	3.66	0.15	5.72	7.37	1.39	0.97	0.59	0.29	0.07	8.82	0.34	2.42
	CV	1.33	3.04	4.57	13.79	1.82	12.04	12.16	3.61	2.08	3.10	1.61	3.13	8.48	8.92	3.26	13.08	1.20	12.77
	Mean	108.25	115.08	17.73	35.91	2.40	65.52	31.23	3.54	282.46	245.04	86.64	32.54	7.09	2.43	2.92	85.02	33.88	28.24
Cluster VIII	SD	4.72	4.60	3.66	5.32	0.24	15.95	0.56	0.60	24.73	27.05	2.17	1.43	0.21	0.09	0.15	17.63	3.17	4.28
	CV	4.36	4.00	20.64	14.81	10.00	24.34	1.79	16.95	8.76	11.04	2.50	4.39	2.96	3.70	5.14	20.74	9.36	15.16

**Table 8: Important superior genotypes and hybrids identified on the basis of their important traits, and cluster means in *Indica* and *Tropical Japonica* rice lines and hybrids**

Genotypes/hybrids	Cluster	Important traits
TJ-25966 × NDR-359	I	Short bold grain, harvest index
TJ-25966 × CSR36	I	Short bold grain, harvest index
TJ-25966 × PB-1	I	Short bold grain, harvest index
TJ-64897 × NDR-359	III	Fourth early days to flowering, short slender, second highest harvest index
TJ-64897 × CSR36	III	Fifth early days to flowering, short slender, fifth highest harvest index
TJ-64897 × PB-1	III	Third early days to flowering, short slender
TJ-10365 × NDR-359	IV	Third highest flag leaf width, fourth highest flag leaf area, second highest flag leaf length, first highest panicle weight, third highest spikelet grains/panicle
TJ-10365 × PB-1	IV	Fourth highest flag leaf area, fifth highest panicle weight, fourth highest spikelet grains/panicle
NDR-359	V	Medium days to flowering, first harvest index, fourth highest grain yield/plant
Pusa Basmati-1	V	Second dwarf plant height, long slender
Pusa Basmati 1121	V	Fifth dwarf plant height, long slender
Sarjoo 52	V	Third dwarf plant height, first highest tillers/plant, third highest second highest grain yield/plant
TJ-25892 × NDR359	VII	First highest panicle length, third highest panicle weight, spikelet/panicle, long bold slender
TJ-25892 × PB-1	VII	Fifth highest panicle weight, long bold slender
TJ-11010 × NDR359	VIII	Second highest panicle bearing tillers/plant, second panicle weight, spikelets/panicle, first highest grains/panicle, first highest spikelet fertility, second highest long bold slender, first biological yield/plant, third highest grain yield/plant
TJ-11010 × PB-1	VIII	Third highest panicle bearing tillers/ plant, second highest spikelets/panicle, highest grains/panicle, Second highest spikelet fertility, first highest test weight, long bold highest biological yield/plant, first highest grain yield/plant
TJ-16081 × NDR-359	VIII	Fourth panicle bearing tillers/plant, first flag leaf length, fourth flag leaf area, fifth highest grains/ panicle, third highest test weight, long slender, second highest biological yield/plant, highest grain yield/plant
TJ-16081 × PB-1	VIII	Fifth highest panicle bearing tillers/plant, second flag leaf length, first flag leaf area, fourth highest panicle length, fifth highest test weight, long slender, fifth biological yield/plant

## **EFFECT OF ANTIMELANOTIC TREATMENT AND VACUUM PACKAGING ON MELANOSIS AND QUALITY CONDITION OF ICE STORED FARMED TIGER SHRIMP (*Penaeus monodon*)**

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### **ABSTRACT**

The objective of the study was to evaluate the effect of vacuum packaging in combination with antimelanotic treatment on quality characteristics and melanosis in farmed tiger shrimp (*Penaeus monodon*) collected from Ratnagiri region of Maharashtra. The processed shrimps were treated with sulfite mixture in water (30 g kg<sup>-1</sup>) and divided into two lots each with 250 shrimps. One of them was vacuum packed, the other was air packed and stored under ice. The ice stored shrimps were analyzed for quality attributes and melanosis for a period of 27 days. The treated vacuum packed shrimps had a shelf life of 24 days, whereas air packed shrimp had 16 days only. Melanosis was absent in the shrimps from both the lots throughout the storage period. Based on the results, it could be concluded that the combination of antimelanotic treatment (sulphite mixture) and packaging could significantly delay the occurrence of melanosis and increase the storage life of shrimp in ice.

**Key words:** Tiger shrimp, antimelanotic treatment, melanosis, Sodium metabisulphite, treated air packed, treated vacuum packed

### **1. Practical applications**

In farmers practice *Penaeus monodon* will be treated with sodium metabisulphite of high concentrations immediately after the capture to prevent melanosis and for keeping quality. Even after the treatment, the shrimps deteriorate

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rapidly and show melanosis within few hours and may also contain high sulphite residues, which is not desirable. In order to overcome these problems, the tiger shrimp were treated with sulphites of low concentration, than that used in regular farmers practice and was combined with packaging to create more period of contact to get more effect of metabisulphite and to prevent reaction of food with oxygen. The combined effect of packaging and sulphite treatment could delay the occurrence of melanosis and can keep the quality of shrimps for a long period (up to 27 days), which is only few days i.e., 2-8 days in case without treatment and packaging. So the work done is very much useful to farmers as well as processors to maintain the edibility of shrimp for longer periods.

### INTRODUCTION

Shrimp continues to be the most important commodity traded in value terms, accounting for 15.4 per cent of the total value of internationally traded fish products (FAO, 2009). Demand for shrimp is increasing at a rapid rate. During 2008-09, shrimp continued to be the major item contributing more than 50% in value of total export of marine products from India (MPEDA, 2010). Shrimp is more susceptible to spoilage from postmortem autolysis and microbial growth. The high ambient temperature of our country favours rapid growth of microorganisms. There are several methods to preserve fresh fish and shrimp. Icing is a common method for short-term preservation.

The appearance of melanosis or blackspots on prawn, shrimp, and other fresh or raw crustaceans is rapid, even in chilled storage, and involves some important economic losses for the fish industry. One of the cheapest, easiest and most efficient methods of preventing blackspot is the use of sulfites (Smith, 1980). In several studies it was reported that sodium metabisulfite extended the melanosis for considerable periods (Chakrabarti and Gupta, 1998; Chakrabarti et al., 1992; Gomez-Guillen et al., 2005; Martinez-Alvarez et al., 2005; Ogawa, 1987 and Smith, 1980). To promote the marketing of fresh fishery products at retail level, novel methods of packaging and storage are required. One such method of current interest is vacuum packaging. Studies have shown that vacuum packaging in combination with lower temperatures improves the shelf life considerably (Jeyasekaran et al., 2004; Lopez-Caballero et al., 2000; Perez-Alonso et al., 2004; Rajesh et al., 2002 and Shalini et al., 2000).

The black tiger shrimp *Penaeus monodon*, is the most important aquacultured shrimp species in the world, accounting for 46% of the total aquaculture shrimp (Hanpongkittikun et al., 1995). *Penaeus monodon* locally known as 'Tiger kolambi' is one of the most important cultivable species along the entire coast of India. Demand for fresh farmed tiger shrimp is very high and storage life in ice is limited. No literature is available on using of combination of vacuum packaging with antimelanotic treatment for prevention of the quality changes in shrimp. The present

study was therefore undertaken to find out the effect of the combination of antimelanotic treatment and vacuum packaging on the storage characteristics of farmed tiger shrimp in ice, especially on occurrence of melanosis and shelf life.

## MATERIALS AND METHODS

### Harvesting and Treatment of Raw Material

Fresh tiger shrimp, *Penaeus monodon* procured from 'Zadgaon brackish water shrimp farm, Ratnagiri, Maharashtra State, India, were brought to the laboratory in iced condition. The time period from the harvest of shrimps to testing was four hours. The shrimps used for this study had an average weight and average total length of  $25 \pm 5$  g and  $15 \pm 2$  cm respectively. The shrimps were processed to remove rostrum and telson and were given dip treatment in seawater at 1:2 ratio (Shrimp to Sea water) containing sodium metabisulfite, citric acid, ethylene di-amine tetra acetic acid and disodium dihydrogen pyrophosphate at the concentrations of 30 g, 20 g, 0.45 g and 30 g for every kilogram of shrimp treated in every 2 liters of sea water (Gomez-Guillen et al., 2005). The sulfite treated shrimp were divided into two lots-1) treated vacuum packed (TVP): The shrimps were treated with  $30 \text{ g kg}^{-1}$  sulfites and then vacuum-packed. 2) Treated Air Packed (TAP): The shrimps were treated with  $30 \text{ g kg}^{-1}$  sulfites and then packed without vacuum (air pack).

### Packing and storage of treated shrimp

The treated shrimps were packed in laminate pouches of 12  $\mu\text{m}$  polyester and 300 guage polyethylenes Lot I was vacuum packed at -1 bar pressure. Vacuum packaging machine (Dong Bang Machineries Pvt. Ltd., China) was used for packaging of whole tiger shrimps. The physical properties of the packaging materials used are given in table 1. Immediately after packing, all the packs were iced with flake ice in the ratio of 1:1 (shrimp to ice) in an insulated box. The insulated box was kept in a cold room maintained at  $0-4^{\circ}\text{C}$ . Re-icing was done every day to supplement the loss due to melting, after draining the melted ice.

### Quality analysis during period of ice storage

Samples in triplicate were drawn from each lot at regular intervals of 24 hrs for analysis throughout the storage period of 27 days and the average results are presented. All the packs were analysed for sensory, microbiological (Total Plate Count (TPC), *Staphylococcus aureus*, Psychrotrophic count, anaerobic count) and biochemical parameters (pH, TMA-N, TVB-N, AAN and FFA).

Moisture, crude protein, fat and ash were determined according to the methods described in AOAC (2000). The moisture content was measured by drying in a hot air oven (Kumar sales corporation, Mumbai), maintained at  $100^{\circ}\text{C} \pm 2^{\circ}\text{C}$  for 16-18 hours. The weight loss was expressed as percent moisture content of the meat. The protein content was determined by estimating total nitrogen in the sample using Kelplus electrically heating digestion and distillation unit (Pelican Equipment,

Chennai). Crude protein content was calculated by multiplying total nitrogen content by 6.25 and expressed as percentage weight of meat.

The crude fat content of the meat was determined using SOCS plus unit (Pelican Equipment, Chennai). The fat present in the sample was extracted by using petroleum ether (AR grade). Then it was dried initially on a water bath at  $98^{\circ}\text{C} - 100^{\circ}\text{C}$  and then in an oven at  $60^{\circ}\text{C} \pm 5^{\circ}\text{C}$ . The difference in the initial and final weight of receiver beaker was determined and fat content of meat was calculated on wet weight basis. In ash content determination preliminary ashing was done by slow heating on a flame to allow smoking off fat without burning. Once the smoke stopped evolving from sample, it was incinerated in a muffle furnace (Cintex, Mumbai) at  $550^{\circ}\text{C} \pm 10^{\circ}\text{C}$  for 5 hours until a white ash resulted. The pH of the sample was determined with a pHm93 pH-meter and a combined pH electrode (Radiometer, Copenhagen, Denmark).

The Free Fatty Acid (FFA) content in the lipid extract was determined with improved titrimetric method as described by (AOAC, 2000). Methanol and isopropanol (chloroform: methanol: Isopropanol = 2:1:2) were added to chloroform extract. The extract was titrated with NaOH using metacresol purple as an indicator. The percentage of free fatty acid was calculated as oleic acid percentage.

Total volatile base nitrogen (TVB-N) and trimethyl amine nitrogen (TMA-N) were determined using a micro diffusion method (Conway, 1950). A 1 ml of sulphuric acid was added to inner chamber of the diffusion unit and 1 ml of muscle solution which was extracted with TCA was added followed by addition of 1 ml of saturated potassium carbonate. The unit was sealed and kept undisturbed overnight. The amount of unreacted acid in the inner chamber was determined by titrating against standard sodium hydroxide with Tashiro's indicator. TVB-N was calculated and expressed in  $\text{mg } 100\text{g}^{-1}$  of the sample. The TMA was determined following the method used for TVB-N estimation, except the addition of neutral formaldehyde to the outer chamber for TMA-N estimation. TMA-N was calculated as explained in TVB-N and expressed in  $\text{mg } 100\text{g}^{-1}$  of the sample.

Determination of  $\alpha$ -amino nitrogen (AAN) content was done according to the method given by Benjakul and Morrissey, 1997. The sample was extracted with TCA and then added with few drops of thymolphthalein and NaOH. Then 10 ml of this was added to 30 ml mixed solution of  $\text{CuCl}_2$ , tri-sodium phosphate and borate buffer (1: 2: 2). A 10 ml of the filtrate is added with 0.5ml of glacial acetic acid and 0.5 g of potassium iodide. The liberated iodine was titrated against sodium thiosulphate using starch as an indicator, when yellow solution of Iodine became faint yellow, few drops of starch solution was added and titration was continued till blue color disappeared. AAN was calculated as  $\text{mg } 100\text{g}^{-1}$  of muscle.

The total plate count, *S. aureus*, psychrotrophic count and anaerobic count were carried out for all the lots throughout storage period by using the method given

by Tomlinson (1995). For estimating total plate count, appropriate dilutions were prepared from the homogenate using physiological saline and placed on plate count agar by pour plate method. The petri dishes were incubated at 37°C for 24 - 48 h. The colonies developed on agar plates were counted and expressed as cfu g<sup>-1</sup>. In case of psychrotrophic count estimation, the petri dishes were pour plated and incubated at 7±1°C for 7-8 days. The colonies developed on agar plates were counted and expressed as cfu g<sup>-1</sup>.

For determining staphylococcal count, the petri dishes were plated by spread plate method on Baird parker agar. The plates were incubated at 37°C for 48 h. the colonies of *S. aureus* were expressed as cfu g<sup>-1</sup>. Total anaerobes were determined by 3 tube MPN technique. The sample was inoculated with 10 ml each into 3 tubes of 10 ml thioglycollate broth, 1 ml each to 3 tubes of thioglycollate broth and 0.1 ml of each to 3 tubes of thioglycollate broth. The test tubes were overlaid with sterile paraffin oil and incubated at 35°C for 48 hours. The probable number of anaerobic count were calculated according to the various combinations of +ve and -ve reactions by using Mac crady table.

Sensory evaluation was done by a 10 point scoring system using a modified version of one developed by Peryam and Pligirms (1957) based on characterization and differentiation of the various sensory characters, such as appearance, texture, odour and flavor. Scores were given by five trained panelists, based on a ten-point hedonic scale. Score 5 was taken as the borderline of acceptability. Evaluation for Melanosis was done by a 4 point scoring system, a modified version used by Gomez-Guillen et al. (2005) for *Parapenaeus longirostris*. Low score indicated less melanosis, good quality and vice versa.

## RESULTS AND DISCUSSION

### Raw material characteristics

The proximate composition of fresh farmed tiger shrimp is given in figure 1. The proximate composition analysed was 75.16% moisture, 1.98% crude fat, 21.07% crude protein and 1.79% crude ash. There were no significant differences found in the proximate composition with increased storage period. Initial sensory, chemical and bacteriological analysis of fresh tiger shrimp indicated that the quality of the raw material was very fresh. The fresh tiger shrimp used in the present study had pH, TMA-N, TVB-N, AAN and indole values of 6.58, 8.92 mg%, 0.34 mg%, 18.46 mg% and 4 µg/100g respectively. It was reported that the raw tiger shrimp used in their study had TMA-N, TVB-N and indole content of 3.78-7.28 mg%, 15.96 - 28.28 mg%, and 5.85 - 6.85 µg/100 g respectively (Antony et al., 2002). The fresh tiger shrimp used in another study had pH and TVB-N values of 6.63 and 5.88 mg% (Rahaman et al., 2001).

The fresh tiger shrimp used in present study had TPC and psychrotrophic counts of 4.08 log cfu/g and 4.02 log cfu/g. Anaerobes and Staphylococcus were not detected in fresh shrimp.

### **Changes in bio-chemical characteristics during ice storage**

The TMA-N content, which indicates the extent of protein breakdown, usually caused by bacteria or enzymes, increased gradually in all the samples with storage time, as observed by earlier authors (Figure 2). In the present work, the initial TMA-N value was 0.34 mg%, which increased to 3.26 mg%, and 2.44 mg% for the TAP and TVP samples respectively on the 8<sup>th</sup> of day storage. The TMA-N further gradually increased up to 17<sup>th</sup> day of storage. On 17<sup>th</sup> day, the TMA-N was recorded to be 5.58 mg% and 3.66 mg% for TAP and TVP samples respectively. These findings are in agreement with the earlier reports on usage of vacuum packaging and treatment with various chemicals to preserve seafoods (Rajesh et al., 2002; Sawant, 2008 and Shakila et al., 2005).

The TVB-N was 8.92 mg % at the beginning of the storage in TAP and TVP samples, which gradually increased to 26.32 mg% and 21.77 mg% for TAP and TVP respectively on 17<sup>th</sup> day of storage (Figure 3). Further on the 26<sup>th</sup> day of storage the values rapidly increased and reached 30.01 mg% for TVP samples. The lower TVB-N values were recorded in TVP sample during storage. The similar trend was observed by Rajesh et al. (2002) and Sawant (2008) during their studies on seafood quality.

A change in pH of fish muscle is usually a good index for quality assessment. In the present study, all the samples exhibited increase in pH during storage, from initial value of 7.11 for both TAP and TVP samples on 0 day, reaching to a value of 7.81 and 7.78 in TAP and TVP samples on 17<sup>th</sup> and 27<sup>th</sup> day respectively (Figure 4.). The study showed similar changes to those observed by Sawant (2008) and Shalini et al. (2000).

The initial AAN content was 18.46 mg% for both TAP and TVP samples (Figure 5). In case of TAP, it increased up to 38.14 mg% which in TVP increased up to 29.40 mg% on 27<sup>th</sup> day. In the present study, the AAN showed a steady increase during the storage period of 26 days and reached 37.62 mg% on 26<sup>th</sup> day. Decrease of AAN during ice storage is attributed to leaching out of soluble nitrogenous constituents (Govindan, 1972). In the present study, the TAP and TVP shrimp samples were not in indirect contact with ice or water. Because of impermeable flexible pouches and hermetic sealing, there was no leaching and AAN values showed only slow increase throughout the storage period. The changes observed were in agreement with the observations made by Rajesh et al. (2002) and Sawant (2008).

In the present study FFA increased from the initial level of 0.16% oleic acid in both TAP and TVP shrimp to 1.89 and 1.26% oleic acid on 17<sup>th</sup> day of storage (Figure 6). The FFA value of TVP samples further increased and reached 1.74% on 26<sup>th</sup> day of storage. In the present study, the FFA value increased significantly with storage period and the increase in TAP was found to be more than that in TVP samples. Similar results have also been reported previously by various researchers

(Haung et al., 1991; Juvekar, 2007; Rajesh et al., 2002 and Shalini et al., 2000) during their work on seafood quality assessment.

### **Changes in microbiological characteristics during ice storage**

During the present study, it was noticed that total plate count (TPC) showed a steady rise after a fall in count during experimental period (Figure 7). The count increased from an initial value of 4.08 log cfu g<sup>-1</sup> to 4.91 and 4.32 log cfu g<sup>-1</sup> for TAP and TVP samples respectively on 8<sup>th</sup> day of storage. The gradual increase still continued and the total plate count recorded were 6.74 and 5.04 log cfu g<sup>-1</sup> for TAP and TVP samples respectively on the 17<sup>th</sup> day. Further it was found that on 26<sup>th</sup> day of storage, the value increased to 7.65 log cfu g<sup>-1</sup> in case of TVP samples. This increase in TPC with storage period was also observed previously in different seafoods during storage (Juvekar, 2007; Sawant, 2008 and Shalini et al., 2000).

Psychrotrophs are the major spoilage bacteria in foods stored at low temperatures. During the present study the psychrotrophic count increased from an initial value of 4.02 log cfu g<sup>-1</sup> to 4.95 and 4.47 log cfu g<sup>-1</sup> on 8<sup>th</sup> day for TAP and TVP samples (Figure 8). The count further showed an increase to 7.04 and 5.76 log cfu g<sup>-1</sup> in case of TAP and TVP samples on 17<sup>th</sup> day of storage. The psychrotrophic count of TVP samples further increased and reached 7.87 log cfu g<sup>-1</sup> on 26<sup>th</sup> day of storage. A similar observations were also made by others (Juvekar, 2007; Sawant, 2008 and Shalini et al., 2000) during their studies.

In the present study, anaerobic counts could not be detected in fresh shrimp, but the counts increased up to 7.2 and 9.4 MPN/g in case of TAP and TVP samples respectively at the end of 8<sup>th</sup> day of storage (Figure 9). Significant increase in anaerobe counts were observed i.e. 21 and 29 MPN/g in case of TAP and TVP samples respectively at the end of 17<sup>th</sup> day of storage. In case of TVP samples the count further increased and reached 43 MPN/g on 26<sup>th</sup> day of storage. The present observation is supported by observations made by others (Juvekar, 2007; Sawant, 2008 and Lyon and Reddmann, 2000).

The presence of *Staphylococcus* in fish is an indication of post harvest contamination. *S. aureus* was not detected in both TAP and TVP shrimp during entire storage period. The absence of *S. aureus* throughout the storage period was also reported by earlier researchers (Baug, 2008; Joseph et al., 1998 and Sanchez et al., 1994).

### **Changes in sensory characteristics during ice storage**

In the present study melanosis was absent in both TAP and TVP samples during the entire storage in ice as observed for 27 days (Figure 10). Whereas samples treated with same concentration of sulfite and ice stored, showed melanosis on 11<sup>th</sup> day. This extension in occurrence of melanosis in TAP and TVP samples might be due to the presence of sodium metabisulfite for whole storage period. However, in our previous studies it was observed that the amount of sulphite residues in treated

and ice stored shrimps was reduced with storage period, which might be due to the effect of washing with ice melted water (Reddy and Patange, 2011).

It was reported that the combined effect of controlled atmosphere and melanosis inhibitors was used to delay blackspot development as compared to the shrimps stored in ice alone (Lyon and Reddmann, 2000; Reddy and Patange, 2011).

There was a significant decrease in sensory score in all the TAP and TVP samples during storage period (Figure 11). From the initial score of 9.75, the scores declined to 4.50 and 4.05 in case of TAP and TVP samples respectively on the day of rejection of 17<sup>th</sup> and 25<sup>th</sup> day of storage. The sensory score of 5 was observed on 16<sup>th</sup> and 24<sup>th</sup> day for TAP and TVP samples respectively. In the present study the shelf life showed significant difference between TAP and TVP samples which was 16 days and 24 days respectively. An extension of 8 days of shelf life was noticed due to packaging under inert condition. The present result is supported by identical observation recorded by earlier workers (Rajesh et al., 2002 and Shakila et al., 2005).

Form the study it was observed that a combination of anitmelanotic treatment and vacuum packaging extends the melanosis and increases the shelf life of tiger shrimp significantly. The samples were acceptable up to 16 days in TAP and 24 days in TVP samples. Extension of storage life in TAP and TVP samples were observed compared to treated shrimps without packaging, which had a shelf life of 14 days in our previous experiment (Reddy and Patange, 2011). This might be due to the presence of sulfites in packaged shrimps, and leaching out of sulfites with melting ice in shrimps without packaging. Extension of storage life in ice in vacuum-packed samples and samples treated with antimelanotic substances has been reported previously (Dalgaard et al., 1993; Mendonka et al., 1989 and Zhuang et al., 1996). In fact there is a very little literature available on combined effect of vacuum packaging and antimelanotic treatment on shrimp quality. It was also observed that the sensory scores correlated well with all other parameters estimated. The sensory scores decreased with the storage period and were acceptable up to 16 days and 24 days for TAP and TVP shrimp.

## CONCLUSION

The use of combination of antimelanotic treatment and vacuum packaging was found to prolong the shelf life of tiger shrimp. The tiger shrimps were acceptable up to 16 days in TAP and 24 days in TVP samples. An extension of 8 days of shelf life was noticed due to packing under vacuum, compared to air-packed samples. An extension of 2 days of shelf life and complete absence of melanosis was noticed due to packing, compared to treated shrimps without packing. This indicates that even packing without vacuum after treatment with sulfite-based treatment ( $30 \text{ g kg}^{-1}$ ) gives a good shelf life and preserves the quality in ice. It is concluded that a good quality and a shelf life up to 24 days in ice for fresh tiger shrimp can be obtained by packing under vacuum and a shelf life of up to 16 days by packing without vacuum after treating with  $30 \text{ g kg}^{-1}$  sulfites.

### ACKNOWLEDGEMENTS

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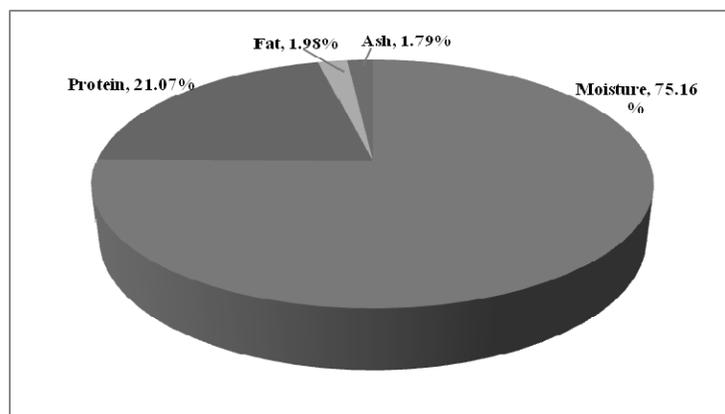
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**Table 1: Physical properties of the packaging material**

<b>*Tensile strength</b>	
Machine direction (MD)	363 Kg/cm <sup>2</sup>
Cross-direction (CD)	349 Kg/cm <sup>2</sup>
<b>*Elongation at break (MD)</b>	
Elongation at break (CD)	80%
<b>*Heat seal strength (MD)</b>	
Heat seal strength (CD)	194 Kg/cm <sup>2</sup>
Water vapour transmission rate	3.62 g/m <sup>2</sup> /24 h at 37°C at 90 + 2%
(as per ISI 060, Part II, 1960)	RH
Oxygen transmission rate (OTR) (as per ASTM 1975)	65 ml/m <sup>2</sup> /atmosphere/24 h/ room temperature (28- 32°C)
*As per IS2508, 1984.	

Figure 1: Proximate composition of fresh tiger shrimp (*Penaeus monodon*)

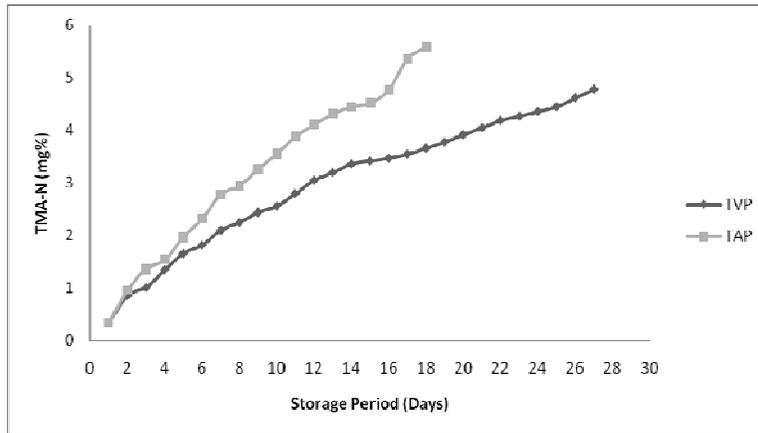


Figure 2: Changes in TMA-N (mg%) content in tiger shrimp during ice storage

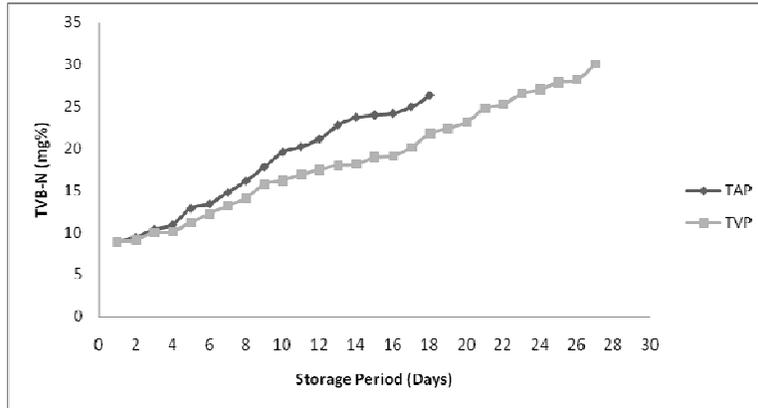


Figure 3: Changes in TVB-N (mg%) content in tiger shrimp during ice storage

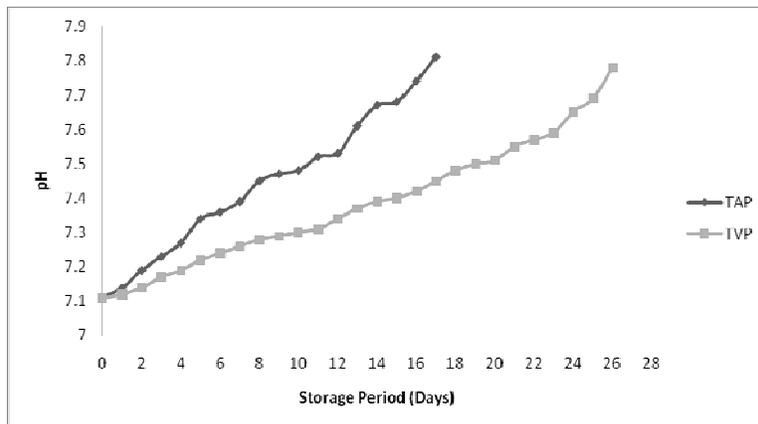


Figure 4: Changes in pH in tiger shrimp during ice storage

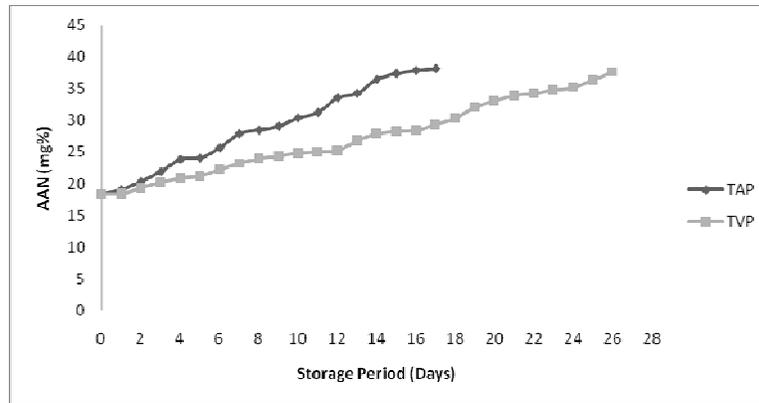


Figure 5: Changes in AAN (mg%) content in tiger shrimp during ice storage

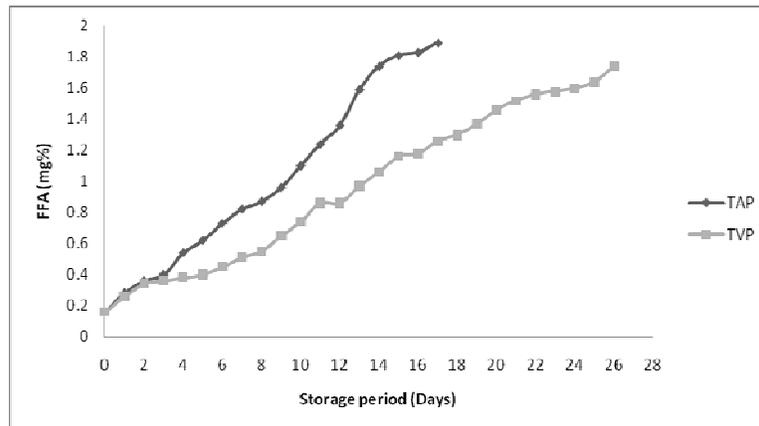


Figure 6: Changes in FFA (Oleic acid %) content in tiger shrimp during ice storage

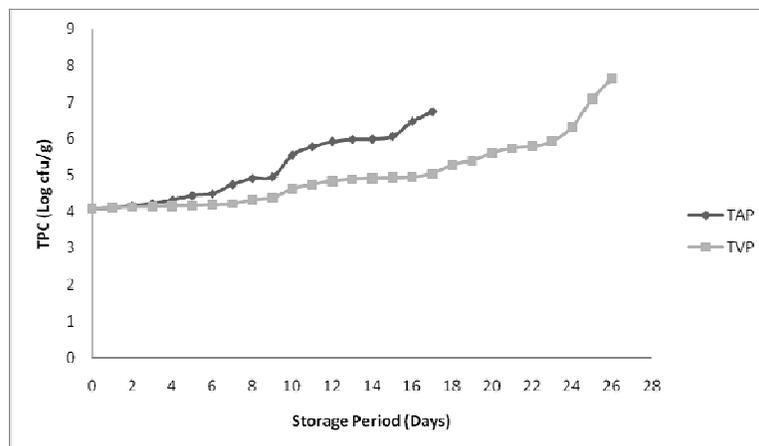


Figure 7: Changes in TPC (Log cfu g<sup>-1</sup>) in tiger shrimp during ice storage

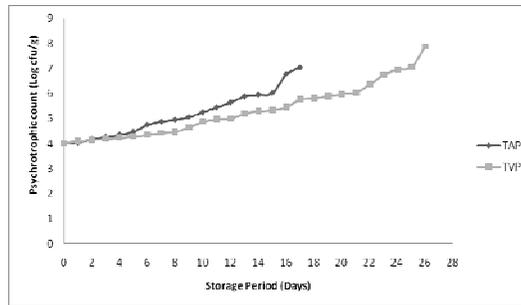


Figure 8: Changes in psychrotrophic count (Log cfu g<sup>-1</sup>) in tiger shrimp during ice storage

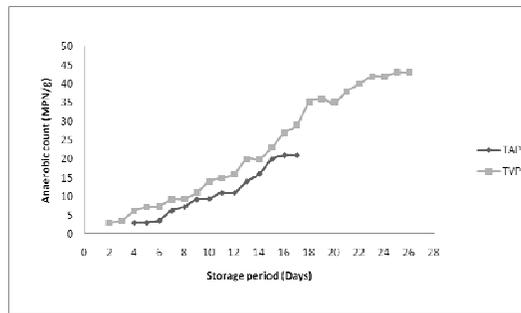


Figure 9: Changes in anaerobic count (MPN g<sup>-1</sup>) in tiger shrimp during ice storage

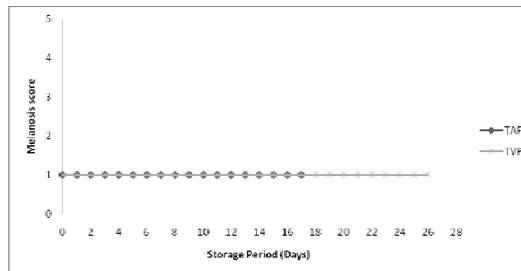


Figure 10: Changes in melanosis score in tiger shrimp during ice storage

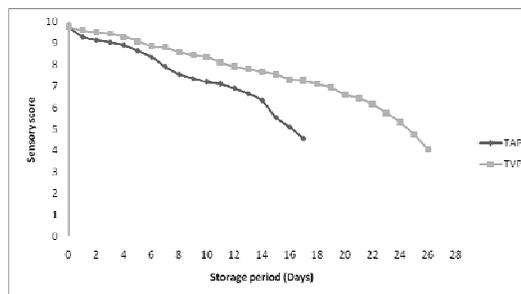


Figure 11: Changes in sensory score in tiger shrimp during ice storage

## EFFECT OF INTEGRATED DISEASE MANAGEMENT PACKAGES ON DISEASES INCIDENCE AND BULB YIELD OF ONION (*Allium cepa* L.)

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### ABSTRACT

A field experiment was conducted to find out the effect of different integrated disease management (IDM) packages on severity of basal rot (*Fusarium oxysporum*), white rot (*Sclerotium rolfsii*) and stemphylium blight (*S. vesicarium*) diseases, and bulb yield of onion (*Allium cepa* L.) variety Agrifound Light Red. Nine IDM packages consisting of NPK fertilizers, farm yard manure, biocontrol agents and spray of fungicides starting from 30 days of transplanting at 15 days interval were applied. The incidence of basal rot and white rot of onion bulbs ranged from 0.98-4.31% and 0.00-0.96%, respectively. The highest incidence of basal rot (4.31%) and white rot (0.96) was recorded in bulbs harvested from untreated control. The lowest incidence of basal rot (0.98%) was found in bulbs harvested from the IDM package consisting of NPK @ 100:50:50 kg ha<sup>-1</sup> + FYM @ 10 t ha<sup>-1</sup> + vermicompost @ 1 t ha<sup>-1</sup> + *Ps. fluorescens* @ 5 kg ha<sup>-1</sup> + copper oxychloride @ 0.3%). White rot did not appear under this package. The lowest Stemphylium blight intensity (3.87%) was achieved with the package having four foliar sprays of propiconazole @ 0.1% followed by mancozeb @ 0.25% and copper oxychloride @ 0.3%. The IDM package increased the bulbs yield over standard check by 25.54% and over untreated control by 109.42%. The average highest bulb diameter (54.15 mm) and bulb size index (23.27 cm<sup>2</sup>) and lowest incidence (0.98%) of basal rot disease in onion bulbs was also obtained with IDM package consisting of NPK @ 100:50:50 kg ha<sup>-1</sup> + FYM @ 10 t ha<sup>-1</sup> + vermicompost @ 1 t ha<sup>-1</sup> + *Ps. fluorescens* @ 5 kg ha<sup>-1</sup> + copper oxychloride @ 0.3%.

**Key words:** *Allium cepa*, integrated disease management.

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## INTRODUCTION

Onion (*Allium cepa* L.) is an exportable bulb crop among the cultivated *Allium* species in India. The country is a traditional exporter of fresh onion with 67% share of the total export of fresh vegetables. Onion is exported to the Iran, UAE, Malaysia, Bangladesh, Sri Lanka, Indonesia, Vietnam, Reunion France and Singapore (Anonymous 2008). Onion is cultivated in summer (Rabi), rainy (Kharif) and dry periods (late Kharif) seasons in the country and maximum area under cultivation is being covered in Rabi season. India ranks second in cultivated area as well as onion production in the world. It has been estimated that more than 25% yield losses occur due to foliar diseases, especially Stemphylium blight (*S. vesicarium*) and purple blotch (*Alternaria porri*). Onion crop also suffers from basal rot (*Fusarium oxysporum*) and white rot (*Sclerotium rolfsii*) diseases causing various extents of losses (Entwistle, 1990). About 30-40% storage losses in onion occur due to incidence of *Fusarium* basal rot (Gupta et al., 2008; Barnoczkine, 1986). The diseases are also responsible for deterioration of export quality of onion bulbs (Gupta et al., 2009).

*Trichoderma viride* and *Pseudomonas fluorescens* have antagonistic capacity against plant pathogens which enhance systemic acquired resistance and plant growth promoting character. *Pseudomonas fluorescens* plays an important role in phosphate solubilization which improves the soil and plant health. Scanty information is available on quality production of onion by approaching integrated crop health and disease management (Gupta et al., 2009).

Gupta et al., (2008) conducted *in-vitro* study for the management of soil borne fungal pathogens like *Fusarium oxysporum* and *Sclerotium cepivorum* using bio agents and oil cakes and findings indicated that *T. viride* inhibited 50.74% mycelia growth as well as 93.59% inhibition in sclerotia production of *S. cepivorum* while 70.40% inhibition in mycelia growth of *F. oxysporum*. Gupta et al. (2011) reported that basal rot and white rot in onion can be managed by the application of *Trichoderma viride*, *T. harzianum* and *Pseudomonas fluorescens*. In view of the above facts, the present study was undertaken to find out the effect of different IDM packages on severity of soil borne and foliar diseases, and yield of onion.

## MATERIALS AND METHODS

The integrated disease management packages were: T<sub>1</sub>=N-P-K (100:50:50) + FYM @ 20t ha<sup>-1</sup> + *T. viride* @ 5kg ha<sup>-1</sup> + Spray of propiconazole @ 0.1%, T<sub>2</sub>= N-P-K (100:50:50) + Vermicompost @ 2 t ha<sup>-1</sup> + *T. viride* @ 5 kg ha<sup>-1</sup> + Spray of propineb @ 0.25%, T<sub>3</sub>= N-P-K (100:50:50) + FYM @ 10 t ha<sup>-1</sup> + Vermicompost @ 1t ha<sup>-1</sup> + *T. viride* @ 5 kg ha<sup>-1</sup> + Spray of copper oxychloride @ 0.3%, T<sub>4</sub>= N-P-K (100:50:50) + FYM @ 10 t ha<sup>-1</sup> + Vermicompost @ 1t ha<sup>-1</sup> + *Ps. fluorescens* @ 5 kg ha<sup>-1</sup> + Spray of copper oxychloride @ 0.25%, T<sub>5</sub>= N-P-K (100:50:50) + FYM @ 20 t ha<sup>-1</sup> + *T. viride* @ 5 kg ha<sup>-1</sup> + Seedling root dip (10g *T.viride*+100g FYM + 1lit.

water) + Spray of propiconazole @ 0.1%, T<sub>6</sub>= N-P-K (100:50:50) + FYM @ 20 t ha<sup>-1</sup> + *Ps. fluorescens* @ 5 kg ha<sup>-1</sup> + Seedling root dip (10g *Ps. fluorescens* +100g FYM + 1lit. water) + Spray of propineb @ 0.25%, T<sub>7</sub>= N-P-K (100:50:50) + FYM @ 10 t ha<sup>-1</sup> + Vermicompost @ 1t ha<sup>-1</sup> + *T. viride* @ 5 kg ha<sup>-1</sup> + *Ps. fluorescens* @ 5 kg ha<sup>-1</sup> + Spray of copper oxychloride @ 0.3%, T<sub>8</sub>= N-P=K (100:50:50) + foliar spray of polyfeed and multi-k @ 1.0% with foliar spray of Mancozeb @ 0.25% (standard check) and T<sub>9</sub>= Untreated Control

The experiment was conducted during Rabi, 2008-09 and 2009-10 in the experimental farm of Regional Research Station, National Horticultural Research and Development Foundation, Nashik, India. The soil of the experimental field was black soil in texture, having pH-7.2, EC-0.155, organic carbon 0.57%, available phosphorus 47.09 kg ha<sup>-1</sup> and potash 414.80 kg ha<sup>-1</sup>. The field was prepared following standard methods for good tilth (Singh et al., 2001) and divided into 3.0m x 1.5m unit plots maintaining 0.50 m space between plots. Drain was made around the unit plots by digging soil from the space between plots. The soil of the drain was used to raise the unit plot. The experiment was laid out in a randomized complete block design with 3 replications. Eight weeks old onion seedlings of variety Agrifound Light Red were transplanted maintaining 10.0 cm plant to plant and 15.0 cm row to row distances. The dosage of manure, fertilizers, biopesticides and fungicides were the same according to IDM packages selected for the experiment. Manure and fertilizers were applied at the time of final land preparation. *Trichoderma viride* (2x10<sup>6</sup> cfu g<sup>-1</sup>) and *Pseudomonas fluorescens* (2x10<sup>8</sup> cfu g<sup>-1</sup>) were obtained from M/s. Sun Agro Biosystem Pvt. Ltd., Chennai. The biopesticides were applied to the soil before transplanting or by dipping roots of seedlings in their suspensions before transplanting. Spraying of the fungicides for the management of foliar diseases was started after 30 days of transplanting. A total of 4 sprays were given at 15 days intervals started after 30 days of transplanting. The insecticide deltamethrin @ 0.1% was sprayed uniformly to control insect pest.

All the recommended intercultural practices were followed to grow the crop. Irrigation was applied at regular intervals to maintain the optimum moisture level in the field. The crop was harvested at bulbs maturity. The incidence and intensity of basal rot and white rot of bulbs were recorded after harvesting based on the covering area of infected bulb by the fungal pathogen following 0 to 5 scales. The severity of Stemphylium blight was indexed at 30, 45, 60 and 75 days after transplanting on a 0-5 scale and Percent Disease Index (PDI) was computed (Sharma, 1995). Data on incidence and intensity of basal rot and white rot disease of onion bulbs, exportable bulb yield, gross yield and bulb size were recorded after harvest. The data recorded in two consecutive years were pooled and analyzed statistically by Randomized Block Design. The cost benefit ratio was also computed.

## RESULTS AND DISCUSSION

### Incidence of basal rot and white rot diseases

The incidence of basal rot and white rot diseases in onion bulbs ranged from 0.98-4.31% and 0.00-0.96%, respectively. The highest incidence of basal rot as well as white rot was recorded in bulbs harvested from untreated control. The lowest incidence of basal rot (0.98%) was found in bulbs harvested from the treatment T<sub>4</sub> (NPK @ 100:50:50 + FYM @ 10 t ha<sup>-1</sup> + vermicompost @ 1 t ha<sup>-1</sup> + *Ps. fluorescens* @ 5 kg ha<sup>-1</sup> + copper oxychloride @ 0.3%). White rot did not appear under this treatment (T<sub>4</sub>). However, the lowest incidence of the disease (0.18%) was recorded from T<sub>5</sub> (NPK @ 100:50:50 kg ha<sup>-1</sup> + FYM @ 20 t ha<sup>-1</sup> + *T. viride* @ 5 kg ha<sup>-1</sup> + seedling root dip (10g *T. viride*+100g FYM + 1 liter water + spray of propiconazole at 0.1%). It was found that soil application of *Ps. fluorescens* and *Trichoderma viride* supplemented with root dip method effectively control the soil borne fungal disease of onion. The data presented in figure 4 revealed that 100% control of white rot disease and maximum control of basal rot disease were obtained with the application of *Ps. fluorescens* @ 5 kg ha<sup>-1</sup> in the soil in treatment T<sub>4</sub> as compared to the untreated control. The present study revealed that *Ps. fluorescens* was most effective against soil borne disease of onion followed by *T. viride* (Table 1).

Data presented in figure 4 revealed that 100% control of white rot and basal rot of onion bulb was obtained with soil application of *Ps. fluorescens* @ 5kg ha<sup>-1</sup> followed by *T. viride* @ 5 kg ha<sup>-1</sup> as soil application as well as seedling root dip (10g *T. viride*+100g FYM + 1liter water) compared to the results of the present study reveal that *Ps. fluorescens* and *T. viride* is most effective against soil borne diseases of onion.

Rajendran and Rangnathan (1996) reported that soil application of *T. viride* and other species were effective against basal rot (*Fusarium oxysporum*) pathogen. Other workers also reported that biocontrol agents such as *T. viride*, *Gliocladium zeae* and *Coniothyrium minitans* gave effective control of white rot disease of onion caused by *Sclerotium rolfsii* (Gupta et al., 2011; Ahmad and Tribe, 1977).

### Incidence and intensity of Stemphylium blight disease

All the tested fungicides significantly reduced incidence and intensity of Stemphylium blight (*Stemphylium vesicarium*) over control. The lowest Stemphylium blight incidence (25.0%) was recorded before 4<sup>th</sup> spray of propiconazole @ 0.1% (T<sub>5</sub>) and mancozeb @ 0.25% (T<sub>8</sub>). However, the lowest Stemphylium blight intensity (3.87%) was achieved with sprays of propiconazole @ 0.1% followed by intensity (4.20 %) in mancozeb @ 0.25%. Significantly the highest Stemphylium blight incidence (43.0%) and intensity (9%) was recorded from untreated control (Table 2).

Propiconazole @ 0.1% was most effective against foliar diseases as well as increasing the onion bulb yield and could be recommended as curative measure against Stemphylium blight as an ad-hoc decision making component of integrated disease management strategy (Table 2).

The data presented in figure 3 revealed that the highest control of *Stemphylium* blight intensity (62.21%) of onion was achieved with foliar sprays of propiconazole @ 0.1% followed by mancozeb @ 0.25% (55.63 and copper oxychloride @ 0.3% (54.78%). The results of the present study reveal that propiconazole was the most effective against *Stemphylium* blight followed by mancozeb and copper oxychloride.

Gupta et al. (1996) reported that spray of contact fungicide mancozeb @ 0.25% reduced the incidence and intensity of *Stemphyllium* blight in onion. Foliar spray of mancozeb @ 0.25 % was also recommended by some other workers for the control of foliar diseases of onion like *Stemphyllium* blight and purple blotch (*Alternaria porri*) of onion (Borkar and Patil, 1995; Srivastava et al., 1999; Mathur and Sharma, 2006; Gupta et al., 2011a & b). Mathur and Sharma (2006) reported that spraying of mancozeb @ 0.2% and copper oxychloride @ 0.3% thrice at 15 days intervals significantly superior in reducing purple blotch and *Stemphylium* blight as well as increasing the yield of onion bulbs. Gupta and Pandey (2011) conducted a field study on management of foliar diseases of onion and reported that propiconazole and mancozeb sprays at 15 days intervals effectively controlled the foliar diseases of onion.

#### **Exportable and gross yield of onion**

Significantly highest exportable onion bulb yield (289.63 t ha<sup>-1</sup>) as well as gross yield (311.81 t ha<sup>-1</sup>) was recorded from the IDM package T<sub>4</sub> (NPK @ 150:50:50 + FYM @ 10 t ha<sup>-1</sup> + vermicompost @ 1 t ha<sup>-1</sup> + *Ps. fluorescens* @ 5 kg ha<sup>-1</sup> + copper oxychloride @ 0.3%) which was followed by yield the treatment T<sub>6</sub> (N.P.K. @100:50:50 + FYM @ 20 t ha<sup>-1</sup> + *Ps. fluorescens* @ 5 kg ha<sup>-1</sup> + seedling root dip (10g *Ps. fluorescens* +100g FYM + 1 lit. water) + Spray of propineb @ 0.25% and treatment T<sub>5</sub> (Table 1). Significantly the highest bulb diameter (54.15 mm) and bulb size index (23.27 cm<sup>2</sup>) was recorded in the package T<sub>4</sub>. The highest cost benefit ratio (1: 4.24) was also recorded in this package (T<sub>4</sub>). However, the lowest exportable yield of onion bulbs (230.89 t ha<sup>-1</sup>) was recorded in standard check T<sub>8</sub> (NPK (100:50:50) + foliar spray of polyfeed and multi-k @ 1.0% with foliar spray of Mancozeb @ 0.25%) and the lowest exportable yield (13.830 t ha<sup>-1</sup>) was observed under control. The highest gross yield (31.181t ha<sup>-1</sup>) of onion bulbs was also recorded from treatment T<sub>4</sub> which was at par with other treatments except gross yield noticed in treatment T<sub>3</sub> (N.P.K. (100:50:50) + FYM @ 10 t ha<sup>-1</sup> + vermicompost @ 1t ha<sup>-1</sup> + *T. viride* @ 5 kg ha<sup>-1</sup> + Spray of copper oxychloride @ 0.3%) and untreated check (Table 1).

Figures 1 and 2 show that, remarkably higher exportable yield of onion bulbs was observed in integrated nutrient applying chemical fertilizers along with organic manures (T<sub>4</sub>) which caused an increase of 25.44% over recommended doses of NPK fertilizers (standard check) and 109.42% over untreated control.

The findings of the present study revealed that judicious use of fertilizers, organic manures, and chemical pesticides supplemented with biopesticide are effective to

increase yield as well as quality of the onion bulb. The improvement in bulb diameter, size index and yield attributes due to phosphate solubilizers may be due to the ability of *Ps. fluorescens* to solubilize and increase availability from insoluble or fixed phosphorus to soluble or readily available phosphorus (P). Similar findings have been reported by other workers (Gupta, 2009; Verma and Mathur, 1989).

The favorable nutritional environment in the root zone created by the addition of organic manures and biofertilizers resulted in increased absorption of the nutrients from soil which was responsible for increasing the yield of onion bulbs (Gupta et al., 2009). Similar findings have been reported in carrot (Luzzati et al., 1980) and in potato (Mandal and Roy, 2001). The higher yield was obtained in potato based cropping systems by the use of fertilizers in combination with organic manures than the use of inorganic fertilizers alone (Sharma and Dua, 1995). Studies conducted on integrated nutrient management (INM) in onion bulb crop improved the quality and yield of onion bulbs through integration of chemical fertilizers along with organic manures (Singh et al., 2001).

### CONCLUSION

The findings of the present study revealed that application of NPK @ 100:50:50 + FYM @ 10 t ha<sup>-1</sup> + vermicompost @ 1 t ha<sup>-1</sup> + *Ps. fluorescens* @ 5 kg ha<sup>-1</sup> with foliar spray of copper oxychloride @ 0.3% was adjudged better in increasing the yield of export quality onion bulbs. Four sprays of mancozeb @ 0.25% and propiconazole @ 0.1% starting from 30 days after transplanting with an interval of 15 days are effective to reduce foliar diseases of onion.

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**Table 1: Yield of onion bulbs, yield attributing characters, soil borne diseases and cost benefit ratio (pooled data of Rabi, 2008-09 and 2009-10)**

Treatments	Exportable yield (t/ha)	Gross yield (t/ha)	Bulb diameter (mm)	Bulb size index (cm <sup>2</sup> )	Basal rot (Inc. %)	White rot (Inc. %)	C:B ratio
T <sub>1</sub>	23.874	26.496	49.0	19.49	1.95	0.45	1:2.09
T <sub>2</sub>	23.463	27.315	48.5	18.67	2.04	0.26	1:2.60
T <sub>3</sub>	21.566	25.378	49.5	18.89	1.95	0.41	1:2.19
T <sub>4</sub>	28.963	31.181	54.2	23.27	0.98	0.00	1:4.24
T <sub>5</sub>	24.785	28.015	48.9	19.08	1.12	0.18	1:2.27
T <sub>6</sub>	25.074	28.109	51.0	20.56	1.55	0.51	1:2.19
T <sub>7</sub>	22.257	26.785	48.5	18.79	2.76	0.46	1:2.32
T <sub>8</sub>	23.089	27.837	49.1	19.30	2.95	0.56	1:1.69
T <sub>9</sub>	13.830	18.500	48.2	18.48	4.31	0.96	
S. Em	20.47	21.81	-	-	0.73	0.42	-
CV %	8.90	8.19			33.78	10.55	
C.D. at 5%	46.30	49.33	-	-	1.65	2.32	-

T<sub>1</sub>=N-P-K (100:50:50) + FYM @ 20t ha<sup>-1</sup> + *T. viride* @ 5kg ha<sup>-1</sup> + Spray of propiconazole @ 0.1%, T<sub>2</sub>= N-P-K (100:50:50) + Vermicompost @ 2 t ha<sup>-1</sup> + *T. viride* @ 5 kg ha<sup>-1</sup> + Spray of propineb @ 0.25%, T<sub>3</sub>= N-P-K (100:50:50) + FYM @ 10 t ha<sup>-1</sup> + Vermicompost @ 1t ha<sup>-1</sup> + *T. viride* @ 5 kg ha<sup>-1</sup> + Spray of copper oxychloride @ 0.3%, T<sub>4</sub>= N-P-K (100:50:50) + FYM @ 10 t ha<sup>-1</sup> + Vermicompost @ 1t ha<sup>-1</sup> + *Ps. fluorescens* @ 5 kg ha<sup>-1</sup> + Spray of copper oxychloride @ 0.25%, T<sub>5</sub>= N-P-K (100:50:50) + FYM @ 20 t ha<sup>-1</sup> + *T. viride* @ 5 kg ha<sup>-1</sup> + Seedling root dip (10g *T.viride*+100g FYM + 1lit. water) +Spray of propiconazole @ 0.1%, T<sub>6</sub>= N-P-K (100:50:50) + FYM @ 20 t ha<sup>-1</sup> + *Ps. fluorescens* @ 5 kg ha<sup>-1</sup> + Seedling root dip (10g *Ps. fluorescens* +100g FYM + 1lit. water) +Spray of propineb @ 0.25%, T<sub>7</sub>= N-P-K (100:50:50) + FYM @ 10 t ha<sup>-1</sup> + Vermicompost @ 1t ha<sup>-1</sup> + *T. viride* @ 5 kg ha<sup>-1</sup> + *Ps. fluorescens* @ 5 kg ha<sup>-1</sup> + Spray of copper oxychloride @ 0.3%, T<sub>8</sub>= N-P-K (100:50:50) + foliar spray of polyfeed and multi-k @ 1.0% with foliar spray of Mancozeb @ 0.25% (standard check) and T<sub>9</sub>= Untreated Control.

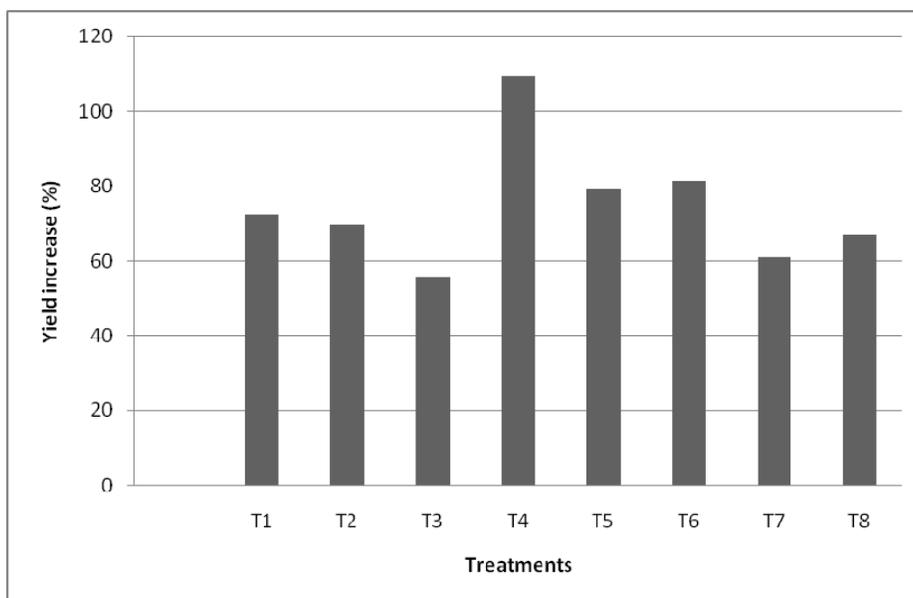


Figure 1: Exportable yield of onion bulbs increased (%) in different treatments over Control CD (P=0.05) 21.32 under field condition

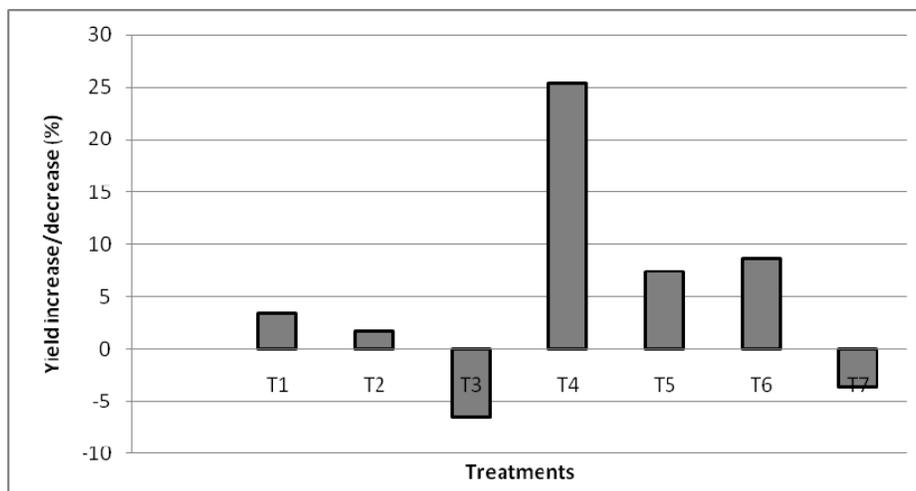


Figure 2: Exportable yield of onion bulbs increased or decreased (%) in different treatments over standard check CD (P=0.05) 23.30 under field condition

**Table 2: Incidence and intensity of Stemphylium blight disease (*Stemphylium vesicarium*) of onion (pooled data of Rabi, 2008-09 and 2009-10)**

Treatment s	Percent incidence and intensity of <i>Stemphylium</i> blight disease							
	Before 1 <sup>st</sup> spray		Before 2 <sup>nd</sup> spray		Before 3 <sup>rd</sup> spray		Before 4 <sup>th</sup> spray	
	Inc.	Int.	Inc.	Int.	Inc.	Int.	Inc.	Int.
T <sub>1</sub>	11.67	0.60	20.00	1.33	28.34	3.74	30.00	5.07
T <sub>2</sub>	15.00	0.94	20.00	1.54	28.34	4.42	31.67	5.60
T <sub>3</sub>	15.00	0.87	21.67	1.74	30.00	4.87	30.00	6.07
T <sub>4</sub>	15.00	0.93	20.00	1.54	26.67	3.60	30.00	5.40
T <sub>5</sub>	11.67	0.74	18.33	1.27	25.00	3.00	25.00	3.87
T <sub>6</sub>	18.34	0.94	20	1.54	30.00	4.14	30.00	5.34
T <sub>7</sub>	16.67	1.20	23.34	2.20	30.00	4.80	35.00	6.47
T <sub>8</sub>	13.34	0.67	18.34	1.53	21.67	2.94	25.00	4.20
T <sub>9</sub>	28.33	2.00	33.33	3.13	36.67	6.14	43.34	9.27
S.E.	3.91	0.35	3.57	0.42	2.93	0.35	2.55	0.55
CV %	24.26	35.13	16.47	24.17	10.26	8.33	8.18	9.61
C.D. at 5%	8.84	NS	8.08	0.95	6.63	0.79	5.77	1.24

**Inc.** : Incidence, **Int.**: Intensity

T<sub>1</sub>=N-P-K (100:50:50) + FYM @ 20 t ha<sup>-1</sup> + *T. viride* @ 5kg ha<sup>-1</sup> + Spray of propiconazole @ 0.1%, T<sub>2</sub>= N-P-K (100:50:50) + Vermicompost @ 2 t ha<sup>-1</sup> + *T. viride* @ 5 kg ha<sup>-1</sup> + Spray of propineb @ 0.25%, T<sub>3</sub>= N-P-K (100:50:50) + FYM @ 10 t ha<sup>-1</sup> + Vermicompost @ 1t ha<sup>-1</sup> + *T. viride* @ 5 kg ha<sup>-1</sup> + Spray of copper oxychloride @ 0.3%, T<sub>4</sub>= N-P-K (100:50:50) + FYM @ 10 t ha<sup>-1</sup> + Vermicompost @ 1t ha<sup>-1</sup> + *Ps. fluorescens* @ 5kg ha<sup>-1</sup> + Spray of copper oxychloride @ 0.25%, T<sub>5</sub>= N-P-K (100:50:50) + FYM @ 20 t ha<sup>-1</sup> + *T. viride* @ 5 kg ha<sup>-1</sup> + Seedling root dip (10g *T.viride*+100g FYM + 1lit. water) +Spray of propiconazole @ 0.1%, T<sub>6</sub>= N-P-K (100:50:50) + FYM @ 20 t ha<sup>-1</sup> + *Ps. fluorescens* @ 5 kg ha<sup>-1</sup> + Seedling root dip (10g *Ps. fluorescens* +100g FYM + 1 lit. water) +Spray of propineb @ 0.25%, T<sub>7</sub>= N-P-K (100:50:50) + FYM @ 10 t ha<sup>-1</sup> + Vermicompost @ 1t ha<sup>-1</sup> + *T. viride* @ 5 kg ha<sup>-1</sup> + *Ps. fluorescens* @ 5 kg ha<sup>-1</sup> + Spray of copper oxychloride @ 0.3%, T<sub>8</sub>= N-P-K (100:50:50) + foliar spray of polyfeed and multi-k @ 1.0% with foliar spray of Mancozeb @ 0.25% (standard check) and T<sub>9</sub>= Untreated Control.

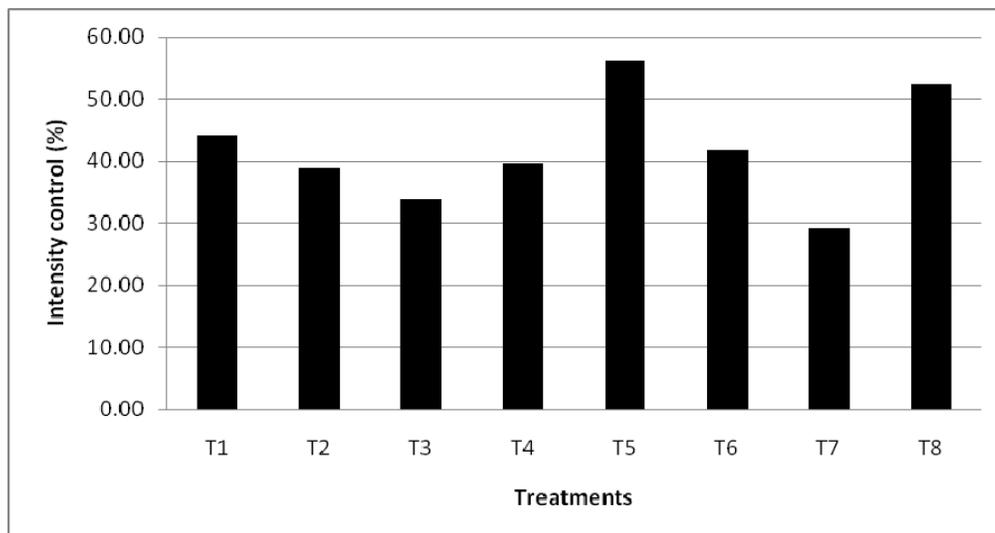


Figure 3: Percent disease control (PDC) of stemphylium blight in different treatments over control CD (P=0.05) 13.82 under field condition

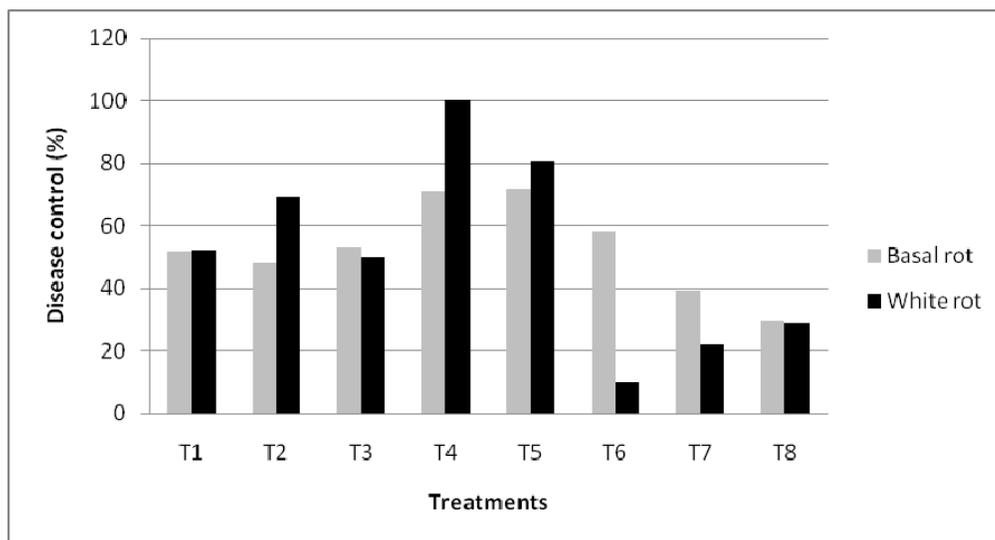


Figure 4: Percent disease control (PDC) of basal rot and white rot in different treatments over control CD (P=0.05) 33.95 under field condition

## INTRA AND INTER CLUSTER STUDIES FOR QUANTITATIVE TRAITS IN GARLIC (*Allium sativum* L)

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### ABSTRACT

Garlic is propagated vegetatively, the clonal selection is the important breeding method and very scanty work has been done on the association between different traits in garlic. Analysis of variance revealed significant divergence in fifteen garlic clone with checks i. e. Yamuna Safed (G-1), Agrifound White (G-41), Yamuna Safed-2 (G-50) and Yamuna Safed-3 (G-282) for different traits, indicating sufficient genetic diversity among the cultivars. Genotypes belonging to the cluster with maximum inter cluster distance were genetically more divergent. Therefore, it is suggested that based upon large cluster distances to select genotypes from all the clusters, which may lead to broad spectrum of favorable genetic variability for bulb yield improvement. Cluster-III had highest value of plant height (93.05 cm), bulb diameter (4.71 cm), bulb size index (16.08 cm<sup>2</sup>), 20 bulb weight (700 g), clove diameter (1.75 cm), clove size index (4.43 cm<sup>2</sup>), weight of 50 cloves (97.50 g) and gross yield (159.63 t ha<sup>-1</sup>) and minimum neck thickness (1.45 cm) number of cloves per bulbs (17) days for bulb initiation (61.66 days) and days to harvesting (149.83). The traits total soluble solids contributed maximum (20.46%) toward genetic divergence followed by gross yield (16.37%), bolters (12.86%), marketable yield (11.11%), number of cloves per bulbs (10.52%), weight of 50 cloves (10.52%), days for bulb initiation (10.52%) and days for harvesting (4.09%). These traits were considered to be most important for genetic divergence, they contributed (96.45%) towards genetic divergence and selection of genotypes based on these traits will contribute to wider genetic diversity in the existing gene pool of garlic genotypes.

**Key words:** Garlic, *Allium sativum*, genetic divergence and D<sup>2</sup> analysis

### INTRODUCTION

Among the spices grown in India, garlic (*Allium sativum* L.) is undoubtedly one of the important crops, propagated vegetatively and hence selection and use of

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diverse parents for its improvement through hybridization is difficult. Garlic consumed by almost all the sections of societies and highly placed for its flavor enhancing capacity and good export potential as fresh bulb as well as in the form of dehydrated product (Gupta and Singh, 1998). Lack of high yielding varieties is one of the main constraints in limiting the production and productivity. Garlic exhibits greater susceptibility to agro-techniques and environmental conditions and possesses a wide range of variability for bulb traits and yield attributes as well as the storability in spite of being vegetatively propagated crop. As this crop is propagated vegetatively, the clonal selection is the important breeding method and very scanty work has been done on the association between different traits which are prerequisite for executing a selection programme. Ministry of Agriculture, India estimate; the annual area under garlic during the year 2012-13 was 2.42 lakh ha and production was 12.28 lakh MT with average yield 5.07 t ha<sup>-1</sup>. Among different states in India, Madhya Pradesh is the leading state accounting for more than 27% of area and 21% of production with average yield of 4.47 t ha<sup>-1</sup>. The other major garlic growing states are Gujarat, Rajasthan, Orissa, Uttar Pradesh, Maharashtra and Tamilnadu. In India per hectare yield are highest in Kerala (19.0 t ha<sup>-1</sup>) followed by Manipur (11.91 t ha<sup>-1</sup>), Punjab (10.96 t ha<sup>-1</sup>), Andhra Pradesh (10.38 t ha<sup>-1</sup>) and West Bengal (9.79 t ha<sup>-1</sup>) (Bhonde et al, 2012).

Diversity arises either due to geographical separation or due to genetic barriers to cross ability and it plays an important role in plant breeding. The knowledge of genetic diversity, its nature and degree of variability would be helpful for selecting desirable genotypes and cultivars for a successful breeding programme. Application of this method helps us for maintaining large number of germplasm and avoiding the duplicates. To meet the domestic requirement and fulfill the export demand, selection of suitable variety for growing under different agro-climatic condition is necessary. In this regards National Horticultural Research and Development Foundation (NHRDF), collected more than 400 germplasm from different garlic growing areas. Among of those 19 germplasm were selected including four checks on the basis of yield and yield contributing traits. The present study was therefore, conducted to determine the genetic divergence among 15 selected genotypes along with four checks i. e. Yamuna Safed (G-1), Agrifound White (G-41), Yamuna Safed-2 (G-50) and Yamuna Safed-3 (G-282) and factors influencing genetic diversity and variability of economic traits to identify suitable genotypes of garlic.

## MATERIALS AND METHODS

The experiment was carried out at National Horticultural Research and Development Foundation, (NHRDF) Salaru, Karnal during 2005-06 and 2006-07. Nineteen diverse genotypes including four checks (released varieties) namely Yamuna Safed (G-1), Agrifound White (G-41), Yamuna Safed-2 (G-50) and Yamuna Safed-3 (G-282) (table-1) were evaluated in randomized complete block design with three replications. Uniform size cloves were selected and planted during the first

fortnight of October in bed of 3.0 m x 1.5 m with a spacing of 10 cm x 7.5 cm. The climate of Karnal is subtropical with minimum and maximum temperature ranging between 2<sup>o</sup> to 45<sup>o</sup>C and favorable for garlic cultivation during Rabi season. Recommended dose of chemical fertilizer such as 100 kg N, 50 kg P<sub>2</sub>O<sub>5</sub> and 50 kg K per hectare) were adopted to ensure a healthy crop growth and development. Harvesting was done as per maturity of different genotypes. Observations were recorded on 10 randomly selected plants in each replications for all the characters viz.-plant height (cm), leaves per plant, neck thickness (cm), bulb diameter (cm), bulb size index (cm<sup>2</sup>), weight of 20 bulbs (g), clove diameter (cm), clove size index (cm<sup>2</sup>), cloves per bulbs, weight of 50 cloves (g), bolters (%), total soluble solid (%), dry matter (%), days for bulb initiation, days for harvesting, gross yield (t ha<sup>-1</sup>) and marketable yield (t ha<sup>-1</sup>).

Pooled data of 2006-2008 were analyzed statistically as suggested by Mahalanobis, (1936). “D<sup>2</sup>” statistics was used to find out generalized distance between the genotypes as per Rao, (1952). The D<sup>2</sup> values were determined to have clustering which was done following Tocher’s method. The clusters were grouped in to four divergence classes (DC) on the basis of mean (M) and standard deviation (S).

## RESULTS AND DISCUSSION

On the basis of D<sup>2</sup> values, all the 19 genotypes were grouped in three clusters (Table-2). Cluster I was largest consisting of 17 genotypes. Cluster II and III had one genotype each. The genotypes belonging to same status or origin were grouped in to different cluster and the genotypes belonging to different origin were grouped in same cluster. The grouping pattern of the genotypes suggested no parallelism between genetic divergence and geographical distribution of genotypes. Singh et al. (2011), Singh et al, (2012), Lokhande et al. (1987), Mohanty (2001) and Mohanty and Prusti (2002) also reported that genotype diversity was independent of geographical region.

Intra and inter cluster D<sup>2</sup> values and corresponding genetic distance are presented in table-3. The highest inter cluster value (D<sup>2</sup>) and genetic distance was noted for cluster-I (852.88), (29.20) while other clusters II and III had zero intra cluster value. Clusters with single genotype (II and III) indicated their independent identity and importance due to various unique characters possessed by them. The intra cluster values were less than inter cluster values indicating the homogenous and heterogeneous nature of the genotypes within and between the clusters. Inter cluster distance (D<sup>2</sup>) is the main criterion for selection of genotype. The genotypes belonging to the cluster with maximum inter cluster distance are genetically more divergent. Therefore, it is suggested that selection of genotypes should be based upon large cluster distances, which may lead to broad spectrum of favorable genetic variability for bulb yield improvement.

Estimates of cluster means for different traits are the measures of inter cluster divergence and degree of homogeneity in these clusters. Cluster mean was worked out and presented in table-4, indicating that the different clusters were superior in respect of various traits.

Cluster-III had highest value of plant height (93.05 cm), bulb diameter (4.71 cm), bulb size index (16.08 cm<sup>2</sup>), 20 bulb weight (700 g), clove diameter (1.75 cm), clove size index (4.43 cm<sup>2</sup>), weight of 50 cloves (97.50 g) and gross yield (159.63 t ha<sup>-1</sup>) and minimum neck thickness (1.45 cm) cloves per bulbs (17) days for bulb initiation (61.66 days) and days to harvesting (149.83) (Table-3). Cluster-II was promising for highest number of leaves per plant (7.60) and minimum bolters (11.38%). Cluster I showed highest cloves per bulb (38.57), total soluble solids (38.11%), dry matter content (40.22%) and marketable yield (134.89 t ha<sup>-1</sup>). The largest cluster-I had more and less average values for most of the traits like yield and yield contributory traits. Any attempt to strengthen the existing gene pool with introduction of garlic materials and their evaluation for desirable traits like resistance to insect pest and disease, adoptability, yield and quality is relevant for improving the potentiality of garlic crop.

The characters contributing maximum to the D<sup>2</sup> values are to be given greater emphasis for deciding the cluster for the purpose of further selection. The study revealed that total soluble solids contributed maximum (20.46%) toward genetic divergence followed by gross yield (16.37%), bolters (12.86%), marketable yield (11.11%), cloves per bulbs (10.52%), weight of 50 cloves (10.52%), days for bulb initiation (10.52%) and days for harvesting (4.09%). All above mentioned traits were considered to be most important for genetic divergence and they contributed (96.45%) towards genetic divergence in this present investigation. Similar findings for contribution towards marketable yield and gross yield were also reported by Khar et al. (2006) in garlic and Mehta et al. (2005) in onion. The result of the present study will help to avoiding duplicates and minimizes the cost of maintenance of garlic germplasm. This also indicates that there is good scope for selection of varieties for desirable traits and cultivation in different part of India for higher productivity.

Therefore it is suggested that based upon large cluster distances to select genotypes from all the clusters, which may lead to broad spectrum of favorable genetic variability for bulb yield improvement.

#### **ACKNOWLEDGEMENT**

The authors are grateful to the Director, National Horticultural Research and Development Foundation, Nashik, for providing necessary facilities during the course of investigation.

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**Table 1: Details of genotypes used in experiment and their source of collection**

SN	Genotype	Source of collection
1	G-4	Calcutta, West Bengal
2	G-176	Rajkot, Gujarat
3	G-189	NBPGR, New Delhi
4	G-192	NBPGR, New Delhi
5	G-200	NBPGR, New Delhi
6	G-222	NBPGR, New Delhi
7	G-255	NBPGR, New Delhi
8	G-284	Mandsaur, Madhya Pradesh
9	G-302	Rajkot, Gujarat
10	G-304	IARI, New Delhi
11	G-305	Karnal, Haryana
12	G-324	NBPGR, New Delhi
13	G-366	Indore, Madhya Pradesh
14	G-368	Dindigul, Tamil Nadu
15	G-369	Kota, Rajasthan
16	Yamuna Safed G-1(C)	Azadpur, New Delhi
17	Agrifound White G-41(C)	Nalanda, Bihar
18	Yamuna Safed-2 G-50 (C)	Karnal, Haryana
19	Yamuna Safed-3 G-282(C)	Dindigul, Tamil Nadu

**Table 2: Distribution of 19 garlic genotypes in different clusters as obtained by multivariate analysis**

Cluster	Genotypes	Name of genotypes
I	17	G-4, G-222, G-176, G-189, G-1, G-284, G-255, G-200, G-304, G-50, G-192, G-41, G-324, G-305, G-368, G-366, and G-369
II	01	G-302
III	01	G-282

**Table 3: Intra and inter cluster  $D^2$  value and distance ( $\sqrt{D^2}$ ) in garlic genotypes**

Clusters	I	II	III
I	852.88 <b>(29.20)</b>	2285.42 <b>(47.81)</b>	5053.38 <b>(71.09)</b>
II		0.00 <b>(0.00)</b>	2070.12 <b>(45.50)</b>
III			0.00 <b>(0.00)</b>

**Table 4: Cluster mean for different traits in garlic genotypes**

Characters	Clusters			% Contribution
	I	II	III	
Plant height (cm)	92.20	87.33	93.05	0.00
Number of leaves/plant	7.14	7.60	7.13	0.00
Neck thickness (cm)	1.48	1.50	1.45	0.00
Bulb diameter (cm)	4.57	4.70	4.71	0.00
Bulb size index (cm <sup>2</sup> )	14.62	13.86	16.08	1.17
Weight of 20 bulbs (g)	601.47	620.00	700.00	0.00
Clove diameter (cm)	1.08	1.18	1.75	0.00
Clove size index (cm <sup>2</sup> )	2.70	3.06	4.43	2.33
No. of cloves/bulbs	38.57	25.66	17.00	10.52
Weight of 50 cloves (g)	45.80	59.66	97.50	10.52
Bolters (%)	17.92	11.38	15.25	12.86
Total soluble solids (%)	38.11	34.96	35.65	20.46
Dry matter (%)	40.22	36.82	38.06	0.00
Days for bulb initiation	65.88	67.00	61.66	10.52
Days for harvesting	158.86	151.66	149.83	4.09
Gross yield (t ha <sup>-1</sup> )	165.57	117.28	159.63	16.37
Marketable yield (t ha <sup>-1</sup> )	134.89	81.23	122.42	11.11

## EVALUATION OF PHYSIOLOGICAL AND ORGANOLEPTIC PROPERTIES OF MANGO CV. KESAR AS INFLUENCED BY IONIZING RADIATION AND STORAGE TEMPERATURE

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### ABSTRACT

Radiation processing of fruits involves exposure to short wave energy to achieve a specific purpose to maintain the physiological changes and sensory quality of the product. The effect of gamma irradiation and storage temperature on physiological changes and organoleptic properties of mango fruit var. Kesar was studied. The fruits were exposed to gamma radiation at different doses i.e. 0.00kGy, 0.20kGy, 0.40kGy and 0.60kGy from the radio isotope <sup>60</sup>Co and stored at different storage environments i.e. at ambient storage (27±2°C with 60-70% RH); at 9°C with 90% RH; at 12°C with 90% RH and control atmosphere storage (at 12°C, O<sub>2</sub> 2%, CO<sub>2</sub> 3% and RH 90%). The fruits irradiated with 0.40kGy gamma rays and stored at 9°C storage temperature with 90% RH recorded maximum reduction in physiological loss in weight and reduced ripening. The minimum physiological loss in weight and ripening and highest marketability of fruits was recorded from fruits irradiated with 0.40kGy gamma rays and stored at 12°C storage temperature with 90 RH including maximum scores on skin colour, pulp colour, texture, taste and overall acceptability at the end of shelf life (41.43 days).

**Key words:** Colour, gamma irradiation, Kesar mango, marketability, ripening, storage temperature, taste

### INTRODUCTION

Mango (*Mangifera indica* L.) is a tropical to sub-tropical fruit; its rapid ripening process and infection caused by microorganism are the major causes of postharvest losses and limit the transport of fresh fruit from the site of harvest to market (Mitra, 1997). Mangoes are growing in over 90 countries worldwide and Asia accounts

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for 77% of global mango production, whereas the America and Africa account for 13 and 19%, respectively (Pereira et al., 2010). Kesar, one of the leading mango cultivars of India is recently emerged out for establishing commercial mango orchard. In export market it becomes more favorite to Alphonso. The ripe fruit has attractive colour, shape and size, rich flavour, pleasant aroma, excellent taste, sweet and fibreless flesh. It has excellent sugar: acid blend. Mangoes are extremely perishable in nature and to extend their ripening during long distance shipment, fruits are generally harvested at physiologically mature stage, stored at low temperatures and ripened at destination under favorable conditions (Yimyong et al., 2011). At the present stage of development, however, sea shipment does not guarantee good quality fruit on arrival for successful marketing. Therefore, if freshly harvested fruit is allowed to ripen at normal ambient conditions (this can vary between 22°C- 32°C and 40-65% RH), ripening processes increase rapidly within the week; the ripe fruit may stay edible for a few days thereafter (Udipi and Ghurge, 2010). The most commonly encountered postharvest problems for mango are shriveling due to water loss, fast ripening and ultimately less marketability. Moreover, chilling injury may occur under low temperature. On the other hand, as the temperature increases, the higher will be the fruits water loss. Mango is susceptible to chilling injury and an optimum temperature of 12-13°C is generally recommended (Gomez – Lim, 1993; Yimyong et al., 2011). Irradiation of fruits has been successfully shown to delay ripening (Pimentel and Walder, 2004) and is the most recent commercial phytosanitary treatment for fresh commodities. Irradiation is a physical process for the treatment of foods compared to conventional process like heating or freezing. It prevents food poisoning, reduces wastage and at the same time preserves quality (Mahindru, 2009). Therefore, the new knowledge is critical because it is important to maintain a balance between the optimum doses required to achieve safety and the minimum change in the sensory quality of the fruit. Therefore, a balance between the required effective dose and tolerance of fruit to irradiation has to be investigated under various storage temperatures. So, irradiation can be used in combination with low temperature for assessing the effects of different doses of gamma irradiation and storage temperature on physiological and organoleptic quality of the fruit. The loss in appearance, taste and texture of fruits is likely to reduce the consumer's acceptability drastically. The objectives of the study were to evaluate the effect of irradiation and storage temperature on physiological and sensory changes during storage. Kesar mangoes from Gujarat have captured sizeable Indian market and have good export potential, but the protocol for their irradiation and post harvest storage needed to be standardized. In this paper the results of studies for standardization protocol of irradiation and storage are presented and discussed.

## MATERIALS AND METHODS

### Material and preparation

The present experiment was carried out at Navsari Agricultural University, Navsari (Gujarat) India during the year 2008-2010. The export grade mangoes of cv. Kesar were harvested from university orchard and were sorted for uniformity in size, maturity and free from defects and washed with chlorinated water. After drying, fruits were packed in corrugated fibre board (CFB) boxes (370 X 275 X 90mm) cushioned with tissue paper. The packed boxes were kept in cold storage at 12°C for 8 h for pre-cooling.

### Radiation processing

After pre-cooling; fruits were irradiated at ISOMED plant (Board of Radiation and Isotope Technology, Sir Bhabha Atomic Research Centre, Mumbai, India). The fruits were exposed to gamma radiation for different doses from the source radio isotope <sup>60</sup>Co with energy 1.33MeV. There were four irradiation doses i.e. I<sub>1</sub> -0.00kGy (Non-irradiated), I<sub>2</sub> -0.20kGy (1.15hrs), I<sub>3</sub> -0.40kGy (2.30hrs) and I<sub>4</sub> -0.60kGy (3.45 hrs).

### Storage conditions

The CFB boxes were stored in cold storage as per storage temperature treatments. The treatments were S<sub>1</sub>-Ambient (27±2°C, 60-70% RH), S<sub>2</sub>-9°C, 90% RH, S<sub>3</sub>-12°C, 90% RH and S<sub>4</sub>- Control atmosphere (CA) storage (12°C, O<sub>2</sub> 2%, CO<sub>2</sub> 3% and RH 90%).The relative humidity maintained through automated RH sensors.

### Parameters and evaluation protocols

The selected parameters and protocols were performed at Post Harvest Technology Unit of American Sprayers and Pressing Equipments (ASPEE) College of Horticulture and Forestry, Navsari Agricultural University, Navsari, India. The parameters and protocols are discussed below.

### Physiological loss in weight (%)

Four fruits from each treatment were weighed on first day of treatment and subsequently their weight was recorded at six day interval up to the end of shelf life. The physiological loss in weight (PLW) was expressed in percentage and calculated as follows.

$$PLW \% = \frac{W_1 - W_2}{W_1} \times 100$$

Where, W<sub>1</sub> = initial weight and W<sub>2</sub> =final weight (Shankar et al., 2009)

**Ripening (%)**

Ripening was measured by the number of fruits having change in colour from greenish to yellow and soft in texture were counted from day 2 day of storage to at six day interval up to the eating ripeness and expressed in percentage over total number of fruits taken for the study.

**Marketable fruits (%)**

The number of good quality and visibly sound fruits that could be marketed were counted and expressed as percentage over the total number of fruits at prescribed interval up to 90 per cent fruits had marketability.

**For senso organoleptic evaluation**

The various treatments were evaluated by a panel consisted of twenty trained panelists and evaluated the sample on the basis of colour, taste, aroma and by pressing the fruit and points were given as per hedonic scale procedure (Rangana, 1986). Higher scoring was treated as more acceptable from the attraction point of view.

**Statistical analysis**

The two years data obtained from experiment was analyzed using ANOVA for completely randomized design with factorial concept in three repetitions. Significance of differences among treatments means were compared using the Fisher's analysis of variance at 5% probability level, technique as followed by Panse and Sukhatme (1967). The data were subjected to appropriate transformation (arcsine) to meet the assumptions of normality.

**RESULTS AND DISCUSSION****Physiological loss in weight (%)**

It is evident from the data (Figure1) that the shelf life was extended up to 38 days and on this day the significantly minimum PLW (5.850%) was recorded in the fruits exposed to 0.40kGy gamma rays ( $I_3S_2$ ) followed by 0.20kGy ( $I_2S_2$ ) gamma rays (7.470%) fruits stored at 9°C temperature. In general, gamma rays @ 0.40kGy ( $I_3$ ) was the most effective treatment in reducing the physiological weight loss in Kesar mango at 9°C ( $S_2$ ) and 12°C ( $S_3$ ) storage temperature as compared to all other treatments. The physiological loss in weight was possibly on account of loss of moisture through transpiration and utilization of some reserve food materials in the process of respiration (Mayer et al., 1960). The irradiation significantly reduced physiological loss in weight during storage period over control which might be attributed to reduction in utilization of reserve food material in the process of respiration (Purohit et al., 2004). The delay in respiration rate as a result of irradiation is also reported by Singh and Pal (2009) in guava. Similar findings were also observed by Prasadini et al. (2008) and El-Salhy et al. (2006) in mango. Similarly, in the different storage conditions, the highest physiological loss in weight was observed in fruits subjected to am-

bient temperature. Minimum physiological loss in weight was noted in cold storage temperature which might be due to lesser water vapour deficit compared to ambient condition and the low temperature which had slowed down the metabolic activities like respiration and transpiration (Mane, 2009). The results are at par with the results of Roy and Joshi (1989) and Waskar and Masalkar (1997) in mango; Nagaraju and Reddy (1995) in banana and Gutierrez et al. (2002) in guava. The significantly minimum reduction in physiological loss in weight of mango fruits subjected to irradiation and stored at various temperatures i.e. at 9<sup>o</sup>C, 12<sup>o</sup>C and in CA (12<sup>o</sup>C) might be due to the mutual complementary effect of irradiation and low temperature.

#### **Fruit ripening (%)**

Figure 2 indicated that the irradiated fruits did not fully ripe up to 38<sup>th</sup> day of storage when fruits were exposed to 0.40kGy gamma rays and kept at 12<sup>o</sup>C (I<sub>3</sub>S<sub>3</sub>) and 0.60kGy kept at 9<sup>o</sup>C (I<sub>4</sub>S<sub>2</sub>) storage temperature were discarded due to the end of their shelf life. The fruits exposed to 0.20kGy and 0.40kGy gamma rays showed 97.93 and 97.30 per cent ripening, respectively at 9<sup>o</sup>C storage (I<sub>2</sub>S<sub>2</sub> and I<sub>3</sub>S<sub>2</sub>). Mangoes are classified as climacteric fruits and ripe rapidly after harvest. The un-irradiated mangoes had early ripeness whereas; gamma ray exposed mangoes had a significantly delayed ripening. The possible mechanisms that have been postulated include: a) irradiations results in decreased sensitivity to ripening action of ethylene and b) alteration in carbohydrates metabolism by regulating certain key enzymes, which interfere with production of ATP which is required for various synthetic processes during ripening (Udipi and Ghugre, 2010). Same findings were noted by Farzana (2005) in mango and Aina et al. (1999) in banana. The lower and delayed ripening was noted at 9<sup>o</sup>C and 12<sup>o</sup>C and in CA (12<sup>o</sup>C) storage as compared to ambient temperature. The decrease of ripening per cent and increase in days to ripening at low temperature may be due to desirable inhibition of enzymatic activities leading to reduction in the respiration and ethylene production (Mane, 2009). These results are supported by Mann and Singh (1975) in mango and Deka et al. (2006) in banana.

#### **Marketable fruits (%)**

The data on per cent marketable fruit are presented in figure 3. The maximum marketable fruit (97.69%) was noted in the fruits irradiated with 0.40kGy gamma rays and stored at 12<sup>o</sup>C (I<sub>3</sub>S<sub>3</sub>), which was followed by I<sub>2</sub>S<sub>3</sub> and I<sub>4</sub>S<sub>3</sub>. The fruits stored at either at 9<sup>o</sup>C (S<sub>2</sub>) or at 12<sup>o</sup>C (S<sub>3</sub>) temperature with (I<sub>2</sub>, I<sub>3</sub> and I<sub>4</sub>) or without (I<sub>1</sub>) gamma rays treatment also showed considerable marketable fruits percentage for long time. The regulation of ripening is an extremely important factor in supplying the consumers with fruit of acceptable eating quality. Innovation in irradiation and cold storage are the new tools for the enhancement of physiological changes and health promoting components in climacteric fruits so, the fruits provide long time marketability. The per cent marketable fruit was significantly influenced by various doses of gamma irradiation and storage temperatures possibly due to the fact that irradiation maintains water content in the fruit and low temperature coupled with high

humidity in cold storage maintains the health of the fruits. Control ripening also a possible reason for long time and higher marketability. These results are in conformity with the findings of El-Salhy et al. (2006) with respect to irradiation and Mane (2009) with respect to low temperature in mango.

### **Skin and pulp colour**

The data pertaining to skin colour showed (Table 1) significant effect on skin and pulp colour. Due to irradiation the maximum score (7.8) for skin colour was obtained in the fruits treated with treatment  $I_3$  (0.40kGy). The maximum score (8.0) for skin colour was obtained in the fruits stored under treatment  $S_3$  ( $12^{\circ}\text{C}$ ). Combined effect of irradiation and storage temperature recorded maximum skin colour score (8.5) when fruits were exposed to 0.40kGy gamma rays stored at  $12^{\circ}\text{C}$  storage temperature ( $I_3S_3$ ). Significantly the maximum pulp colour score (7.5) of fruits was recorded (Table 1) when fruits were exposed to 0.40kGy gamma rays ( $I_3$ ). The maximum score (7.7) for pulp colour obtained from fruits stored under treatment  $S_3$  ( $12^{\circ}\text{C}$ ). Combinedly the maximum pulp colour score of fruits (7.8) was recorded in fruits exposed to 0.40kGy gamma rays and stored at  $12^{\circ}\text{C}$  storage temperature ( $I_3S_3$ ). The visual changes in the colour of the fruits were noticeable throughout the storage. It was noticed that outer skin and inner pulp of mango fruits irradiated at medium (0.40kGy) and lower dose (0.20kGy) of gamma rays had greener skin and yellow to pink pulp than the skin and pulp of the higher dose (0.60kGy) exposed fruits at the end of storage. The large changes in skin and tissue colour during ripening may have overwhelmed any irradiation responses (Zhao et al., 1996). Similar results were also found by El-Salhy et al., (2006) in mango and by Paull (1996) in papaya. The storage temperature had significant effect on skin and pulp colour. The fruit stored at  $12^{\circ}\text{C}$ , ambient temperature and in CA ( $12^{\circ}\text{C}$ ) storage recorded higher colour score whereas, minimum colour score was recorded in fruits stored at  $9^{\circ}\text{C}$ . It may be due to the slow degradation of chlorophyll due to slow rate of respiration and ethylene production by desired temperature and it is also associated with chloroplast to chromoplast transition. The present results support the results of Medlicott et al. (1990), Waskar and Masalkar (1997) and El-Salhy et al. (2006) in mango and Gutierrez et al. (2002) in guava.

### **Texture of fruit**

The panelists found significant differences in the texture of the un-irradiated and irradiated fruits. It is evident from the data, presented in Table 2 that significantly maximum texture score (7.6) was recorded in fruits exposed to treatment  $I_3$  (0.40kGy). Maximum score (7.8) for texture was obtained from fruits stored at treatment  $S_3$  ( $12^{\circ}\text{C}$ ). Collectively the maximum texture value (8.0) was recorded in fruits exposed to 0.40kGy gamma rays and stored at  $12^{\circ}\text{C}$  storage ( $I_3S_3$ ). The panelists found a significant difference between the dose levels of gamma ray with respect to texture score. The possible reasons may be that medium to lower dose of irradiation can break chemical bonds, increase membrane permeability and metabolic activity,

which will lead to more water vapor movement to intercellular space and maintain the texture of the fruit and another reason may be that the changes in pectin by irradiation are possible cause of the radiation induced softening (Zhao et al. 1996). These results are in agreement with findings of Moreno et al. (2006), El-Salhy et al. (2006) and Lacroix et al. (1990) in mango; Pimentel and Walder (2004) in banana and Singh and Pal (2007) and Singh and Pal (2009) in guava. Significantly better texture score was recorded at the full ripening stage in fruits stored at 12<sup>0</sup>C, ambient temperature and in CA (12<sup>0</sup>C) storage compared to fruits stored at 9<sup>0</sup>C temperature, which may be due to retardation of the biochemical changes and ripening process at desired temperature (Mane, 2009) and significantly decrease in texture during storage of fruits was due to changes in nature of pectin substances which cementing the cell wall and hydrolysis of starch, hemicelluloses and cellulose during ripening of fruit (Leopold, 1964). This finding is in conformity with the findings of Waskar and Masalkar (1997) in mango; Purvoko (2002) in banana.

#### **Taste of fruits**

Significantly the maximum (Table 2) taste score (7.7) was recorded in fruits exposed to 0.40kGy gamma rays (I<sub>3</sub>). Similarly, maximum score (7.9) for taste obtained in fruits stored under treatment S<sub>3</sub> (12<sup>0</sup>C) followed by treatment ambient storage (S<sub>1</sub>). In combination, the maximum taste score (8.2) was recorded in fruits exposed to 0.40kGy gamma rays stored at 12<sup>0</sup>C (I<sub>3</sub>S<sub>3</sub>) Significantly enhanced taste of fruits might be due to optimum moisture and uniform ripening at desired temperature and radiation dose. The 9<sup>0</sup>C stored fruits lost their taste and flavour due to prolonged storage of mango fruits at lower temperature and high humidity (Dhemre and Waskar, 2004).

#### **Overall acceptability**

Significantly the maximum overall acceptability score (7.8) was recorded (Table 2) in fruits exposed to 0.40kGy gamma rays (I<sub>3</sub>). The maximum score (7.9) for overall acceptability obtained when fruits were stored under treatment S<sub>3</sub> (12<sup>0</sup>C). Combined effect of irradiation and storage temperature showed the maximum overall acceptability score (8.4) in fruits exposed to 0.40kGy gamma rays and stored at 12<sup>0</sup>C storage (I<sub>3</sub>S<sub>3</sub>). The higher score of acceptability of fruits in these treatments may be due to good appearance, better texture and best in taste. Another reason is that the combined advantage of irradiation and desired temperature which reduces undesirable sensory changes, since free radicals are less mobile at low temperature and hence their ability to interact with other constituents reduces (Udipi and Ghurge, 2010). The same results were obtained by Rajput et al. (2004) in apple with respect to irradiation and Mane (2009) in mango with respect to storage temperature.

### CONCLUSION

Exposure of mango fruits (var. Kesar) to 0.40kGy gamma rays subsequently stored at 9°C temperature delayed the ripening process which maintained lower percentage of physiological loss in weight and ripening per cent and higher percentage of marketable fruits. This was followed by the fruits treated with 0.40kGy gamma irradiation subsequently stored at 12°C temperature. The highest score for skin and pulp colour; texture and taste of fruit and overall acceptability was observed in fruits exposed to 0.40kGy gamma rays and stored under 12°C storage temperature compared to minimum in fruits stored at 9°C temperature irradiated with 0.60kGy gamma radiation. Overall conclusions are that the irradiation increases the fruit health and overall acceptability for consumers point of view. It fulfills the quarantine criteria for export purpose.

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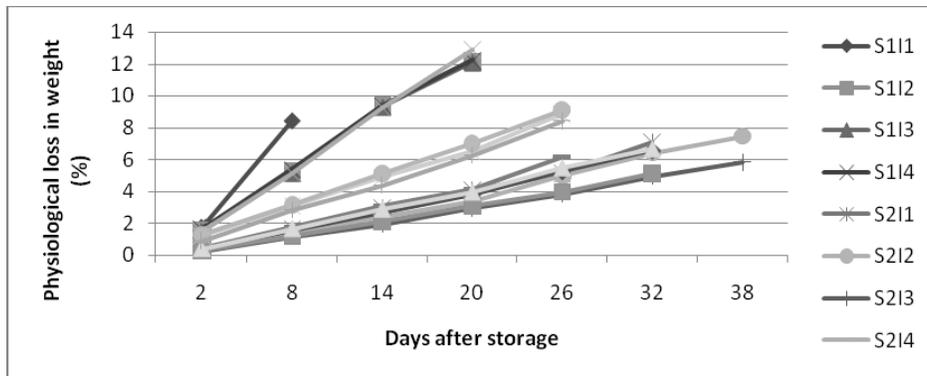


Figure 1: Weight loss by irradiation and storage temperature during storage of mango cv. Kesar

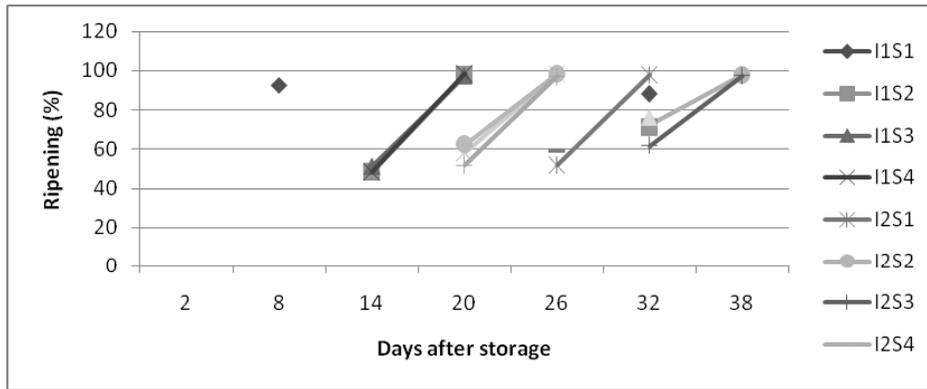


Figure 2: Ripening changes by irradiation and storage temperature of mango cv. Kesar during storage

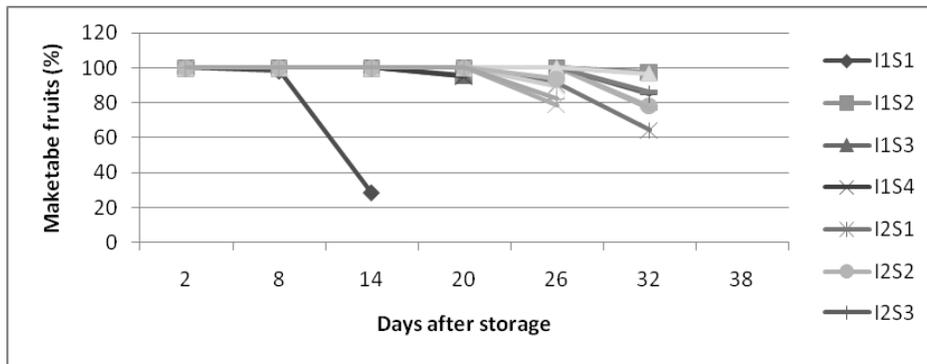


Figure 3: Response to irradiation and storage temperature on marketable fruits of mango cv. Kesar

**Table 1: Effect of irradiation and storage temperature on skin and pulp colour score of mango cv.Kesar**

Factors	Skin colour					Pulp colour				
	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>4</sub>	Mean	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>4</sub>	Mean
S <sub>1</sub>	7.5	7.8	7.8	7.0	7.5	7.4	7.5	7.7	7.3	7.5
S <sub>2</sub>	6.8	7.0	7.0	6.7	6.9	6.2	6.8	7.1	6.2	6.5
S <sub>3</sub>	7.9	8.0	8.5	7.7	8.0	7.5	7.7	7.8	7.5	7.7
S <sub>4</sub>	7.5	7.7	7.8	7.2	7.5	7.2	7.2	7.5	7.2	7.3
Mean	7.5	7.6	7.8	7.1		7.1	7.3	7.5	7.0	
Factors	I	S	I X S			I	S	I X S		
S. Em ±	0.004	0.004	0.008			0.009	0.005	0.009		
CD (P≤0.05)	0.012	0.012	0.023			0.026	0.014	0.026		

Where, I=irradiation, S-storage temperature

**Table 2: Effect of irradiation and storage temperature on texture, taste and overall acceptability score mango CV Kesar**

Factors	Texture					Taste					Overall acceptability				
	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>4</sub>	Mean	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>4</sub>	Mean	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>4</sub>	Mean
S <sub>1</sub>	7.4	7.5	7.8	7.0	7.5	7.6	7.7	7.9	7.5	7.7	7.4	7.6	7.7	7.1	7.5
S <sub>2</sub>	6.6	6.8	7.1	6.4	6.7	6.7	6.8	7.2	6.5	6.8	6.6	6.8	7.5	6.5	6.8
S <sub>3</sub>	7.8	8.0	8.0	7.5	7.8	7.8	7.9	8.2	7.7	7.9	7.6	8.1	8.4	7.5	7.9
S <sub>4</sub>	7.2	7.3	7.4	7.0	7.2	7.5	7.6	7.7	7.4	7.6	7.3	7.4	7.6	7.2	7.4
Mean	7.2	7.4	7.6	7.0		7.4	7.5	7.7	7.2		7.2	7.5	7.8	7.1	
Factors	I	S	I X S			I	S	I X S			I	S	I X S		
S. Em ±	0.005	0.005	0.010			0.005	0.005	0.010			0.005	0.005	0.010		
CD (P≤0.05)	0.015	0.015	0.028			0.015	0.015	0.030			0.016	0.016	0.030		

Where, I=irradiation, S-storage temperature

## CHARACTERIZATION AND FIELD PERFORMANCE OF 15 STRAWBERRY GERMPLASM UNDER BANGLADESH CONDITIONS

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### ABSTRACT

Fifteen strawberry germplasm collected from local and exotic sources were evaluated for different qualitative and quantitative characters. The germplasm showed variation for majority of the studied traits. Plant height during peak harvest ranged from 14.67 to 25.00 cm and FA 007 exhibited the tallest plants. The maximum number of leaves plant<sup>-1</sup> was produced by FA 006 (46.67). The maximum number of runners plant<sup>-1</sup> was obtained in FA 003 (68.67) followed by FA 004 (63.00), while the highest number of crown plant<sup>-1</sup> was found in FA 007 (15.33) followed by FA 006 (14.67). Days to flowering varied from 48.33 to 102.30 among the germplasm and FA 008 required minimum (48.33) days for flowering. The germplasm FA 004 produced the highest number of flower trusses (35.50 plant<sup>-1</sup>) while it was the lowest in FA 010 (5.00 plant<sup>-1</sup>). The number of flowers plant<sup>-1</sup> was found maximum in FA 003 (168.00 plant<sup>-1</sup>) and minimum in FA 010 (40.00 plant<sup>-1</sup>). Among the germplasm pollen viability varied significantly and maximum viable pollen was recorded in FA 010 (84 %), while it was lowest in FA 004 (12.00 %). The highest per cent fruit set was recorded in BARI Strawberry-1 (86 %) while, FA 013 (39 %) showed the lowest. Among the germplasm the highest yield plant<sup>-1</sup> was recorded from FA 005 (737.70 g) followed by FA 006 (702.30 g) and was significantly higher than others while the lowest yield plant<sup>-1</sup> was recorded from FA 013 (52.00 g), FA 014 (69.00 g), FA 009 (81.33 g) and FA 010 (121.30 g).

**Keywords:** Strawberry, qualitative growth characters, quantitative growth characters, reproductive characters

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## INTRODUCTION

Strawberries (*Fragaria x ananassa* Duch.) are widely grown fruit crop in the world, adapted in geographically diverse area (Biswas et al., 2009). Strawberry is now grown successfully in Bangladesh. There are more than 20 *Fragaria* species and numerous cultivars commercially cultivated in several countries (Gaafar and Saker, 2006). The cultivars vary remarkably in morphological characters (Biswas et al., 2009). The genotypic and phenotypic interaction of strawberry plant is highly cultivar dependent. Therefore, the choice of a cultivar is of paramount importance for successful strawberry cultivation (Asrey and Singh, 2004). On the other hand, as a new crop it is necessary to evaluate the morphological characters under Bangladesh condition. A germplasm collection with good variability for the desirable characters is the basic requirement of any crop improvement program (Singhania et al., 2006; AVRDC, 1990). In addition, crop improvement is primarily based on extensive evaluation of germplasm. Morphological characters of a plant is most important for nature as well as yield of a crop. Hence, studies about these important traits are necessary for successful cultivation of this crop in a new area like Bangladesh.

Bangladesh Agricultural Research Institute has released one variety of strawberry named BARI strawberry-1, which is not enough for increasing demand of strawberry cultivation. So, it is necessary to develop more variety of this promising crop. None of work based on field performance and morphological characterization of strawberry under Bangladesh condition was done. However, a good number of investigations were done in India, Pakistan and elsewhere in the world. Therefore, the present study was undertaken to characterize the collected strawberry germplasm for morphological characters and to select superior one on the basis of morphological characters.

## MATERIALS AND METHODS

**Experiment site:** This study was conducted at the Fruit Research Farm of Horticulture Research Centre of Bangladesh Agricultural Research Institute, (Latitude 23°59' N, Longitude 90°24' E, Altitude 14.33 m), Gazipur, Bangladesh during August 2008 to May 2010. This region falls in sub-tropical zone having hot summers (May–August) and mild winter (December–February). Cumulative rainfall of about 104 – 112 mm during cultivation period with average 79.4 % relative humidity. The mean maximum and minimum temperature during cropping period were 25.89 and 17.05°C, respectively. Soil of the experimental farm was clay loam, having pH 6.2, which was low in organic carbon (0.95 %), very low in available phosphorus (9 ppm) and low in potash (0.17 meq/100 g soil).

**Treatments:** Healthy and disease free daughter plants of 15 strawberry germplasm collected from local and exotic sources were considered as treatments and planted in the experimental field. The released strawberry variety in Bangladesh namely BARI Strawberry-1 was used as check (Appendix 1).

**Experimental design and layout:** The experiment was laid out in a randomized complete block (RCB) design with three replications. The unit plot size was 100 x 280 cm and the plants were spaced 50 x 40 cm on open beds. Beds were raised 30 cm above main field with 50 cm drain in-between 2 beds. Each plot contained double row accommodating 14 plants. Daughter plants of strawberry germplasm were planted in September 15, 2008 and 2009. Data were collected from inner plants from each row to avoid border effect. In each unit plot ten inner plants were selected for recording data.

**Intercultural operations:** Runners were removed at every 3 to 4 days intervals in order to make the crown capable to initiate flowers. Straw mulch was applied around the plants as a normal practice in order to conserve soil moisture, decreasing weed and to avoid contact of the fruits with soil for reduction of rot caused by soil microbes. Weeds were removed whenever necessary to keep the crop weed free. Irrigation was given whenever necessary to keep soil moisture available in the field for better plant growth. All other necessary cultural practices and plant protection measures were followed uniformly for all the plots and treatments during the entire period of experimentation.

**Observations recorded:** Following vegetative characters viz. growth habit, plant vigor, foliage density, foliage color of strawberry plants and position of inflorescences were recorded by close observation according to the descriptor developed by IBPGR (1986). On the other hand, plant height (cm), plant spread (cm), leaves plant<sup>-1</sup>, runners plant<sup>-1</sup>, crowns plant<sup>-1</sup>, days to flowering, flower trusses plant<sup>-1</sup>, flowers plant<sup>-1</sup>, size of flower (cm), disk size of flower (receptacle) (cm), pollen viability (%) and percent fruit set were recorded for quantitative growth characters of strawberry plants and yield plant<sup>-1</sup> (g).

**Pollen viability (%):** Freshly opened flowers were collected to assess viability of the pollen grains. The pollen grains from the anther cone were dusted on a glass slide. Carmino Acetic Acid (CAA) solution (single drop) was used to stain the specimen and was covered with a cover slip. Pollen grains were viewed under a light microscope. The normal and properly stained pollen grains were considered as viable.

**Data analysis:** Two year's data of different quantitative parameters were pooled and analyzed, following RCB design using MSTAT-C (Nisen, 1983). The mean comparison was done following the Duncan's Multiple Range Test (DMRT).

## RESULTS AND DISCUSSION

### Qualitative growth characters

**Growth habit:** Regarding growth habit of plants, the germplasm were categorized into 3 types viz. erect, intermediate and prostrate (Table 1). Six germplasm (FA 001, BARI Strawberry-1, FA 005, FA 008, FA 013 and FA 014)

exhibited intermediate type of growth habit, five germplasm (FA 004, FA 006, FA 007, FA 016 and FA 017) exhibited erect type of growth and rest of the germplasm exhibited prostrate type of growth habit. Meulenbroek et al. (2002) found that the plants of 'Pavana' variety exhibited open (prostrate) growth habit.

**Plant vigor:** Based on plant vigor, the germplasm were classified into 3 viz., weak, intermediate and strong (Table 1). Among 15 germplasm studied, strong vigorous plants were observed in 6 germplasm (FA 003, FA 004, FA 006, FA 007, FA 016 and FA 017); 5 germplasm (FA 001, BARI Strawberry-1, FA 005, FA 008 and FA 011) were intermediate and the rest 4 germplasm (FA 009, FA 010, FA 013 and FA 014) were weak in plant vigor. Ludvikova and Paprstein (2003) found that plants of 'Pegasus' and 'Honeoye' varieties were vigorous in nature. Their findings corroborated with the present observation.

**Foliage density:** Foliage densities of strawberry plants were classified into sparse, intermediate and dense (Table 1). Among the germplasm plants of 7 germplasm (FA 001, BARI Strawberry-1, FA 005, FA 010, FA 011, FA 013 and FA 014) showed intermediate, 5 germplasm (FA 004, FA 006, FA 007, FA 016 and FA 017) showed dense and remaining 3 (FA 003, FA 008 and FA 009) sparse type of foliage. Ludvikova and Paprstein (2003) found that the variety 'Pegasus' was dense and 'Honeoye' was intermediate in nature. Results of their findings are similar to the present observation.

**Foliage color:** Foliage coloration of plants was grouped into light, intermediate and dark (Table 1). Among the germplasm, 8 (FA 001, BARI Strawberry-1, FA 003, FA 008, FA 009, FA 011, FA 013 and FA 014) were dark in foliage color; 2 (FA 004 and FA 010) germplasm were intermediate and plants of rest germplasm (FA 005, FA 006, FA 007, FA 016 and FA 017) were light in color.

**Position of inflorescences:** Positions of inflorescences in different germplasm were categorized into beneath, level with and above the foliage (Table 1). Among the studied germplasm, it was found that the plants of 9 germplasm (FA 003, FA 004, FA 005, FA 006, FA 007, FA 010, FA 011, FA 016 and FA 017) produced inflorescences, which were beneath the foliage. Inflorescences of 4 germplasm (FA 001, BARI Strawberry-1, FA 008 and FA 013) were above the foliage and rest two germplasm (FA 009 and FA 014) produced inflorescences level with the foliage (Table 1). The germplasm having inflorescences above the foliage was suitable for harvesting. Meulenbroek et al. (2002) found that cultivar 'Pavana' had long flower clusters that make the fruit easy to pick.

#### **Quantitative growth characters**

**Plant height:** Significant variation among the germplasm was observed for plant height. The tallest plant (25.00 cm) was recorded in the genotype FA 007 and the shortest (14.67 cm) in the genotype FA 013 (Table 2). The variation in plant height might be due to the genetic makeup. Results of present investigation regarding plant height were partial

agreement with the findings of Rahman (2011), and Asrey and Singh (2004). Riyaphan et al. (2005) obtained significant variation in plant height of strawberry plants in Thailand, which ranged from 10 to 20 cm at mid-harvesting time.

**Plant spread:** Significant variation among the germplasm was observed for plant spread. The highest plant spread (24.67 cm) was recorded in the genotype FA 005 and FA 006, and the lowest (15.00 cm) in the genotype FA 013 (Table 2). The variation in plant size might be due to the genetic makeup of strawberry plants. Asrey and Singh (2004) and Rahman (2011) found a significant variation in plant spread. Results of present investigation regarding plant size was in agreement with the findings of Asrey and Singh (2004) and Rahman (2011).

**Leaves plant<sup>-1</sup>:** Leaf number is important for photosynthesis and it differed significantly ranging from 46.67 (FA 006) to 20.33 (FA 014). Among the germplasm, FA 006 produced the maximum number of leaves and FA 014 produced the minimum (Table 2.). The number of leaf plant<sup>-1</sup> in different germplasm varied mainly due to inherent characters of the germplasm. Rahman (2011) obtained the number of leaves plant<sup>-1</sup> varying from 60.52 to 49.00 in different germplasm which was more or less closer to the findings of the present study. But the finding about leaves plant<sup>-1</sup> reported by Asrey and Singh (2004) was lower than the present study. They found that number of leaves ranged from 3.33 to 6.33 plant<sup>-1</sup> under irrigation in semi-arid region of Punjab, which might be due to the genetic as well as the environmental effect. Perez-de-Camacaro et al. (2002) obtained a variation in leaf number among cultivars during the later part of the season and found that cv. 'Bolero' and 'Everest' produced the highest and 'Elsanta' the least number of leaves plant<sup>-1</sup>. In another study revealed a significant variation in leaves plant<sup>-1</sup> among strawberry germplasm and recorded 4.6 to 9.8 leaves plant<sup>-1</sup> at 86 days after planting in Cambridge.

**Number of runner and crown plant<sup>-1</sup>:** Significant variation among the germplasm was observed in respect of numbers of runner and crown plant<sup>-1</sup>. Among the germplasm, FA 003 produced the highest number of runner (68.67) followed by FA 004 (63.00) while FA 007 produced the lowest (6.33). FA 007 produced the maximum number of crown (15.33) followed by FA 006 (14.67) and FA 005 (11.33), while FA 009 and FA 011 produced the lowest (6.67) (Table 2.). The variation in different germplasm as recorded might be due to genetic makeup. This result is in confirmatory with the findings of Pérez-de-Camacaro et al. (2002). They found a significant difference in number of runner and crown plant<sup>-1</sup> among the strawberry cultivars in UK. 'Elsanta' produced more runners than 'Bolero' and 'Everest'. While cv. 'Bolero' produced more crowns than 'Everest' and 'Elsanta'. In the present study, FA 005, FA 006 and FA 007 produced the minimum number of runners but the maximum number of crowns. This result is at par with that of Riyaphan et al. (2005). They found significant variation in numbers of runners and crowns plant<sup>-1</sup> and reported that cv. 'Tioga' produced the maximum number of crowns and the minimum number of runners in Thailand.

**Relationship between runner and crown per plant:** A negative linear relationship was observed between number of crown per plant and number of runner per plant (Figure 1). The equation was  $y = -0.0952x + 12.612$  and the value of the coefficient of determination ( $R^2 = 0.4801$ ) gave a good fit and the fitted regression line had a significant regression coefficient, indicating number of crown plant<sup>-1</sup> will be increased with a significant manner with the decrease of number of runner plant<sup>-1</sup>. So, there is a clear indication that higher number of runners inhibited the crown producing ability of strawberry plants. This result is in strong agreement with that of Riyaphan et al. (2005).

### **Reproductive characters**

**Days to flowering:** Significant variation among the germplasm in relation to days to flowering was observed. The maximum number of days (102.3) was required for flowering in FA 004 and the minimum (48.33 days) in FA 008 (Table 3). This is because FA 008 might be early genotype. It is in agreement with the findings of Riyaphan et al. (2005) who recorded that number of days from transplanting to first blooming ranged from 67.50 to 78.90 in cv. ‘Tioga’ and from 40.80 to 47.23 in cv. ‘Tochiotome’. Asrey and Singh (2004) found a significant variation in days to flowering, which is also in consonant with the present findings. They found that strawberry cultivar ‘Seascape’ required maximum number of days for flowering followed by ‘Chandler’, while minimum in Fern. Rahman (2011) found that days to flowering varied significantly among the strawberry germplasm and it ranged from 72.42 to 81.87 days. Macit et al. (2007) reported that ‘Sweet Charlie’ and ‘Camarosa’ were earlier in first flowering than other varieties, while Kabarla was late under conventional system in Samsun, Turkey.

**Flower trusses and flowers plant<sup>-1</sup>:** Marked variation was observed among the germplasm for number of flower trusses (inflorescences) and flowers plant<sup>-1</sup>. Among the germplasm, FA 004 produced maximum number of flower trusses plant<sup>-1</sup> (35.5), while FA 010 produced only 5 flower trusses plant<sup>-1</sup>. During the study period, the highest number of flowers plant<sup>-1</sup> (168) was observed in FA 003 followed by FA 016 (145), FA 017 (144) and FA 004 (142), which were statistically similar; while the lowest was in FA 010 (40) shown in table 1.3.3. The number of flower trusses and flowers plant<sup>-1</sup> in different germplasm varied mainly due to inherent mechanism of the germplasm. This result was partially similar with that of Rahman et al. (2011), who found that number of flower trusses and number of flowers plant<sup>-1</sup> differed significantly among the germplasm, ranged from 6.58 to 8.10 and from 37.11 to 53.42, respectively. Michel et al. (2006) found that the total number of flowers plant<sup>-1</sup> was linearly related to its number of inflorescences (trusses). They found on an average 25.90 flowers inflorescence<sup>-1</sup>. But Sønsteby and Heide, (2008) recorded less number of inflorescences with a more number of flowers plant<sup>-1</sup> in cv. ‘Korona’. Riyaphan et al. (2005) and Pérez-de-Camacaro et al. (2002) found that the number of inflorescences plant<sup>-1</sup> significantly varied in strawberry cultivars, which are in close conformity with the present findings. They also observed that the early flowering germplasm produced less number of trusses as well as flowers

plant<sup>-1</sup> compared to late flowering germplasm. This the fact might be due to that the early flowering germplasm produced inflorescences before optimum vegetative growth, which might have suppressed inflorescences and flowers number plant<sup>-1</sup>. On the other hand, late flowering germplasm reach reproductive phase after attaining optimum vegetative growth, which might enhance flowering potentiality. In addition, the early flowering germplasm produced flower under higher temperature which have negative impact on inflorescences and flower production. These observations are supported by the findings of Patterson (1995) and Chercuitte et al. (1991), while Maurer and Umeda (1999) found that irrespective of flowering time, flowers number plant<sup>-1</sup> varied significantly among the germplasm.

**Relationship between days to flowering and flower trusses plant<sup>-1</sup>:** A positive quadratic relationship was observed between days to flowering and flower trusses plant<sup>-1</sup> (Figure 2). The equation was  $y = -138.92 + 4.1467x - 0.0241x^2$  and the value of the coefficient of determination ( $R^2 = 0.7786$ ) was significant at 5% level of probability, indicating flower trusses plant<sup>-1</sup> increased with a significant manner with the increase of days to flowering up to 85 days. So, there is a clear indication that early flowering germplasm produced less number of flower trusses plant<sup>-1</sup> compared to late flowering germplasm. This result is in line with the findings of Patterson (1995) and Chercuitte et al. (1991). But flower trusses decreased in those plants that required more than 85 days for earlier flowering. This might be due to extreme delayed plants flowered at higher temperature which inhibited flowering in strawberry. Ledesma et al. (2008) found comparatively high temperature significantly reduced the number of inflorescences in strawberry.

**Size of flower:** Size of flower in respect of diameter and thickness varied significantly. This might be due to genetic variation among the germplasm. Among the germplasm FA 007 produced the largest flower in respect of diameter and thickness (3.43 cm and 2.07 cm, respectively) followed by FA 006 (3.30 cm and 2.03 cm, respectively), while FA 004 produced the smallest flower having the lowest diameter (2.07 cm) and thickness (0.87 cm) and shown in table 4. This result is in agreement with the findings of Rahman (2011) and Verma et al. (2002), who found that diameter and thickness of flower ranged from 1.74 to 2.44 cm and 1.40 to 2.36 cm, respectively.

**Disk size of flower (receptacle):** Size of receptacle (disk) of flower in respect of diameter and thickness varied significantly. Among the genotype FA 006 produced flowers having the biggest receptacle in respect of diameter (2.60 cm) while, FA 004 produced flowers with the smallest receptacle diameter (1.33 cm). The thickest receptacle was found in the flowers of FA 007 (1.60 cm) and the thinnest receptacle was found in the flowers of FA 004 and FA 003 (Table 4). Rahman (2011) and Verma et al. (2002) found that flower disk diameter and thickness differed significantly among the germplasm ranging from 0.66 to 0.90 cm. and 0.31 cm to 0.65 cm, respectively which were lower compared to present observation. This might be due to difference of cultivation or the growing environment.

**Pollen viability:** Pollen viability is one of the most important attributes for proper fruit setting. The germplasm varied significantly in respect of pollen viability. The maximum viable pollens were found in FA 010 (84.00 %) followed by FA 008 (81.67 %), FA 001 (81.33 %) and BARI Strawberry-1 (76.00 %) and those were statistically similar, but the lowest viable pollen grains were found in FA 004 (12.00 %) and FA 003 (12.33 %) (Table 4). Such variation in pollen variability might be due to the influence of ploidy level and effect of high temperature which prevailed during the time of flowering in these two germplasm. Ledesma and Sugiyama (2005) stated that pollen viability of strawberry varied significantly among the germplasm, which was supported by the present observation. Such variation might be due to the effect of germplasm, temperature and their interaction. In addition, pollen quality was also influenced by high temperature, but the effect of higher temperature on pollen viability is cultivar specific. According to Kronenberg (1959), pollen viability differs among strawberry cultivars which affect the fruit set and yield of strawberry.

**Per cent fruit set:** The number of fruit set is directly dependent on the abundance of flowers and their existence in the plant which was controlled by environment and genetic makeup of the germplasm. The highest per cent fruit set was recorded in BARI Strawberry-1 (86 %) followed by FA 008 (81 %) and FA 010 (73 %) and they were statistically similar, and significantly higher than rest of the germplasm while FA 013 (39 %) showed the lowest (Figure 3). Among the germplasm FA 003 and FA 004 failed to produce any fruit due to very poor pollen viability (Table 4). The variation in different germplasm is due to variation of genetic makeup and differential response of the germplasm to climatic conditions of the locality. Ledesma et al. (2008) found that per cent fruit set of strawberry varied significantly among the cultivars ranging from 60.20 to 92.40 they also observed a significant interaction between temperature and cultivars regarding fruit set (%) which was similar to the results of the present experiment.

**Relationship between pollen viability and percent fruit set:** A positive linear relationship was observed between pollen viability (%) and per cent fruit set (Figure 4). The equation was  $y = 1.1062x - 17.108$  and the value of the coefficient of determination ( $R^2 = 0.8139$ ) gave a fitted regression line and indicating a significant regression coefficient, representing fruit set will be increased with a significant manner with the increase of pollen viability (%). So, there is a clear indication that increase in percent pollen viability has significant positive effect on percent fruit set. Abdul-baki and Stommel, (1995) and Cross et al. (2003) stated that fruit set was directly hampered by poor pollen performance. Their results are strongly supported by the present findings.

**Yield plant<sup>-1</sup>:** Among the germplasm the highest yield plant<sup>-1</sup> was recorded from FA 005 (737.70 g) followed by FA 006 (702.30 g) and was significantly higher than others (Figure 5). The lowest yield plant<sup>-1</sup> was recorded from FA 013, FA 014, FA 009 and FA 010 which produced only 52.00 g, 69.00 g, 81.33 g and 121.30 g fruits plant<sup>-1</sup>,

respectively. The variation in yield plant<sup>-1</sup> was due to the inherent character of the germplasm. Rahman (2011) found that yield of strawberry varied significantly and ranged from 442.50 to 129.85 g. Present findings was consonant with the results of Rahman (2011). From different experiments Pires et al. (2006) and Crespo (2010) stated that fruit yield plant<sup>-1</sup> in strawberry varied significantly among the cultivars studied which were strongly supported by the present result.

### CONCLUSION

The germplasm of strawberries showed variation in morphological as well as quantitative traits and also in yield plant<sup>-1</sup>. On the basis of physico-morphological characters, it was concluded that genotype FA 006 and FA 007 were identical with FA 016 and FA 017, respectively. On the basis of different physico-morphological characters along with the higher yield plant<sup>-1</sup> the germplasm FA 005, FA 006 and FA 007 were selected for further investigation. On the other hand, based on poor pollen viability, poor fruit set as well as low yield plant<sup>-1</sup> the germplasm FA 003 and FA 004 were discarded.

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**Table 1: Qualitative growth characters of strawberry germplasm**

Germplasm	Growth habit	plant vigor	Foliage density	Foliage color	Position of inflorescences
FA 001	Intermediate	Intermediate	Intermediate	Dark	Above the foliage
FA 003	Prostrate	Strong	Sparse	Dark	Beneath the foliage
FA 004	Erect	Strong	Dense	Intermediate	Beneath the foliage
FA 005	Intermediate	Intermediate	Intermediate	Light	Beneath the foliage
FA 006	Erect	Strong	Dense	Light	Beneath the foliage
FA 007	Erect	Strong	Dense	Light	Beneath the foliage
FA 008	Intermediate	Intermediate	Sparse	Dark	Above the foliage
FA 009	Prostrate	Weak	Sparse	Dark	Level with the foliage
FA 010	Prostrate	Weak	Intermediate	Intermediate	Beneath the foliage
FA 011	Prostrate	Intermediate	Intermediate	Dark	Beneath the foliage
FA 013	Intermediate	Weak	Intermediate	Dark	Above the foliage
FA 014	Intermediate	Weak	Intermediate	Dark	Level with the foliage
FA 016	Erect	Strong	Dense	Light	Beneath the foliage
FA 017	Erect	Strong	Dense	Light	Beneath the foliage
BARI Sb.-1	Intermediate	Intermediate	Intermediate	Dark	Above the foliage

**Table 2: Quantitative growth characters of strawberry germplasm**

Germplasm	Plant height (cm)	Plant spreading (cm)	Leaves plant <sup>-1</sup>	Runners plant <sup>-1</sup>	Crowns plant <sup>-1</sup>
FA 001	15.33 de	22.00 ab	31.00 e	44.67 c	7.33 cd
FA 003	16.33 de	23.33 a	34.00 de	68.67 a	9.00 b-d
FA 004	17.00 d	21.67 ab	36.67 cd	63.00 a	9.67 bc
FA 005	21.67 bc	24.67 a	42.67 b	11.67 f	11.33 b
FA 006	22.67 b	24.67 a	46.67 a	8.33 f	14.67 a
FA 007	25.00 a	24.33 a	40.00 bc	6.33 f	15.33 a
FA 008	16.67 de	23.00 a	20.67 f	31.33 d	8.67 b-d
FA 009	15.67 de	20.00 a-c	21.33 f	44.33 c	6.67 d
FA 010	15.00 de	18.00 b-d	21.00 f	43.00 c	7.00 cd
FA 011	15.00 de	18.00 b-d	23.67 f	53.00 b	6.67 d
FA 013	14.67 e	15.00 d	21.33 f	53.67 b	7.00 cd
FA 014	17.00 d	16.00 cd	20.33 f	43.00 c	7.33 cd
FA 016	20.00 c	21.67 ab	40.00 bc	19.33 e	9.00 b-d
FA 017	21.33 bc	23.33 a	40.67 bc	20.67 e	9.00 b-d
BARI Strawberry-1	17.00 d	23.00 a	31.33 e	33.33 d	8.67 b-d
<b>Level of significance</b>	<b>**</b>	<b>**</b>	<b>**</b>	<b>**</b>	<b>**</b>
<b>CV (%)</b>	<b>4.82</b>	<b>8.69</b>	<b>5.55</b>	<b>7.49</b>	<b>12.24</b>

Figures having the same letter(s) in a column do not differ significantly by DMRT.

Level of significance \*\* means significant at 1 % level

**Table 3: Days to flowering, flower trusses per plant and flowers per plant 16 in strawberry germplasm**

Germplasm	Days to flowering	Flower trusses plant <sup>-1</sup>	Flowers plant <sup>-1</sup>
FA 001	55.67 e-g	5.70 gh	74.10 c-e
FA 003	95.67 b	33.60 a-c	168.00 a
FA 004	102.3 a	35.50 a	142.00ab
FA 005	61.33 de	34.00 ab	136.00 b
FA 006	66.33 cd	31.50 b-d	135.40 b
FA 007	69.33 c	31.00 cd	124.00 b
FA 008	48.33 h	6.70 gh	80.40 cd
FA 009	56.67 ef	6.00 gh	42.00 f
FA 010	50.33 f-h	5.00 h	40.00 f
FA 011	52.00 f-h	14.50 e	65.25 c-f
FA 013	49.67 gh	11.50 f	46.00 ef
FA 014	50.00 gh	11.60 f	52.20 d-f
FA 016	61.67 de	29.00 d	145.00 ab
FA 017	64.00 cd	32.00 bc	144.00 ab
BARI Strawberry-1	49.67 gh	8.40 g	84.00 c
<b>Level of sig.</b>	<b>**</b>	<b>**</b>	<b>**</b>
<b>CV (%)</b>	<b>4.29</b>	<b>6.08</b>	<b>12.58</b>

Figures having the same letter(s) in a column do not differ significantly by DMRT.  
Level of significance \*\* means significant at 1 % level.

**Table 4: Size of flower, size of flower disk and per cent pollen viability in strawberry**

Germplasm	Size of flower (cm)		Size of flower disk (cm)		Pollen viability (%)
	Diameter	Thickness	Diameter	Thickness	
FA 001	2.97 d	1.37 a-d	2.13 cd	1.10 cd	81.33 ab
FA 003	2.13 fg	0.87 d	1.43 gh	0.60 e	12.33 g
FA 004	2.07 g	0.87 d	1.33 h	0.60 e	12.00 g
FA 005	3.20 bc	1.90 a-c	2.50 ab	1.47 ab	74.67 b-d
FA 006	3.30 ab	2.03 ab	2.60 a	1.47 ab	66.33 d-f
FA 007	3.43 a	2.07 a	2.30 bc	1.60 a	69.67 c-e
FA 008	2.97 d	1.43 a-d	1.97 de	1.10 cd	81.67 ab
FA 009	2.37 e	1.27 cd	1.80 ef	0.83 de	64.33 ef
FA 010	2.33 ef	1.17 cd	1.67 fg	0.83 de	84.00 a
FA 011	2.30 ef	1.27 cd	1.63 fg	0.83 de	74.33 b-d
FA 013	2.23 e-g	1.30 b-d	1.60 f-h	0.87 de	69.33 c-e
FA 014	2.33 ef	1.27 cd	1.80 ef	0.87 de	60.00 f
FA 016	3.03 cd	1.60 a-d	2.23 b-d	1.37 a-c	72.33 c-e
FA 017	3.17 b-d	1.67 a-c	2.13 cd	1.43 ab	73.33 b-d
BARI Strawberry-1	3.03 cd	1.53 a-d	2.23 b-d	1.23 bc	76.00 a-c
<b>Level of significance</b>	<b>**</b>	<b>**</b>	<b>**</b>	<b>**</b>	<b>**</b>
<b>CV (%)</b>	<b>3.14</b>	<b>8.61</b>	<b>5.82</b>	<b>11.09</b>	<b>5.53</b>

Figures having the same letter(s) in a column do not differ significantly by DMRT.  
Level of significance \*\* means significant at 1 % level.

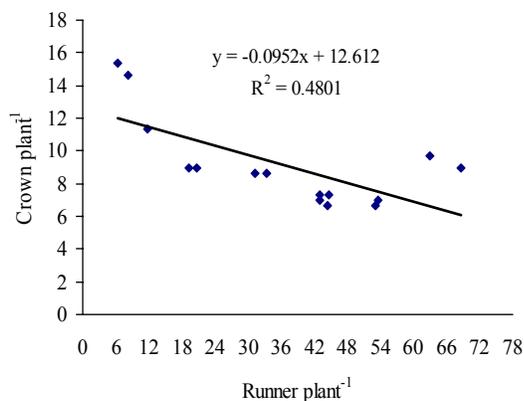


Figure 1: Relationship between number of runner plant<sup>-1</sup> and number of crowns plant<sup>-1</sup>

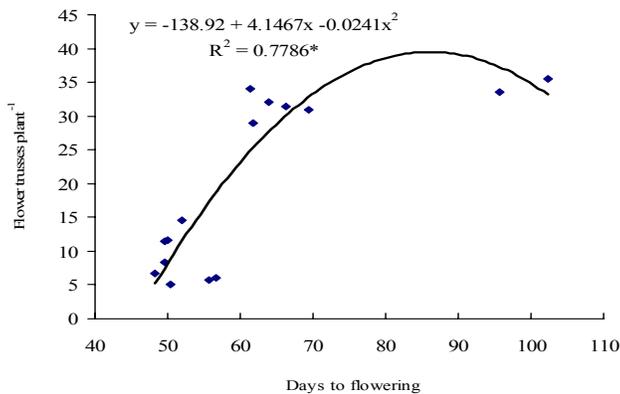


Figure 2: Relationship between days to flowering and number of flower trusses plant<sup>-1</sup>.

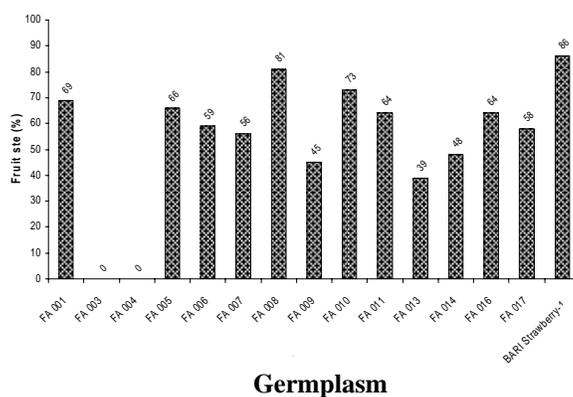


Figure 3: Fruit set (%) in different strawberry germplasm.  
Note. Plant of FA 003 and FA 004 failed to produce any fruits.

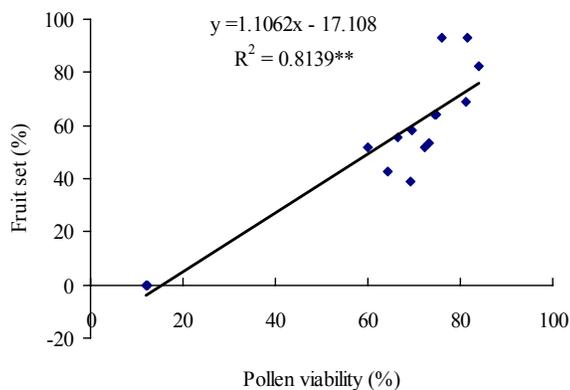


Figure 4. Relationship between pollen viability and percent fruit set in strawberry

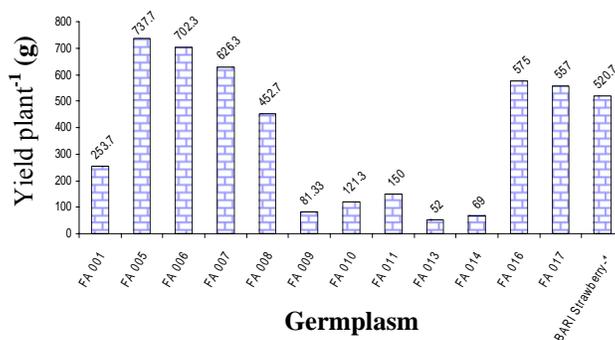


Figure 5: Yield plant<sup>-1</sup> (g) in different strawberry germplasm.

#### Appendix 1: Accession number, source of collection and country of origin of collected strawberry germplasm/varieties

Accession No.	Source of collection	Country of origin
FA 001	BARI	Japan
FA 003	Parvin Nursery	Unknown
FA 004	Parvin Nursery	Unknown
FA 005	Personal contact	Florida, USA
FA 006	Personal contact	Florida, USA
FA 007	Personal contact	Florida, USA
FA 008	Rajshahi University	Japan
FA 009	Rajshahi University	Japan
FA 010	Rajshahi University	Japan
FA 011	Kashban Nursery	India
FA 013	Kashban Nursery	India
FA 014	Parvin Nursery	India
FA 016	Krishibid Nursery	Unknown
FA 017	Krishibid Nursery	Unknown
BARI Strawberry-1	BARI	Japan

## MOLECULAR DIVERSITY ANALYSIS IN POTATO (*Solanum tuberosum* L.) THROUGH RAPD MARKERS

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### ABSTRACT

Random Amplified Polymorphic DNA (RAPD) markers were used to study the molecular diversity of 12 popular potato varieties in Bangladesh. DNA was extracted from tender leaf sample for PCR amplification. The PCR amplified DNA profile was visualized on 2% agarose gel, staining with ethidium bromide. Eight RAPD primers were used to evaluate the genetic diversity of potato varieties. Some total of 36 DNA fragments were amplified and out of them 24 were polymorphic. Those primers generated 61.53% of polymorphic DNA band. The primer OPX 04 produced highest (9) number of DNA band and out of 9 amplicon 6 were polymorphic. Lowest number of amplification was observed in the primer OPA-17 and it was only 3. The highest Nei's genetic distance (0.9701) was noticed between the variety Granola and Provinto. The highest (0.8205) number of genetic identity/similarity was observed between the varieties Cardinal and Diamant. The unweighted pair group method of arithmetic mean (UPGMA) dendrogram based on Nei's genetic distance revealed that the 12 varieties followed into two main clusters. The present finding showed that there was high level of genetic diversity among the varieties which can be used for parental selection in potato breeding program.

**Key words:** Molecular diversity, RAPD, potato.

### INTRODUCTION

Potato (*Solanum tuberosum* L.) is a highly heterogeneous and vegetatively propagated crop. It is one of the important food crops of Bangladesh as well as in many other countries of the world. It produces more calories and protein per unit of land with minimum time than any other field crops (Upadhyaya, 1995). Because of its high yield potential and food value as compared to rice and wheat, it is considered as a promising candidate crop for feeding the hungry people of the world (Pushkarnath,

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1976). The yield level of potato in Bangladesh is lower than other potato growing countries of the world (BBS, 2010). The use of local seed and traditional varieties are the major constrains of low yield in potato. Development of high yielding varieties having good keeping quality is one of the challenges for potato breeders. Genetic variability has been considered as is prerequisite for crop improvement program. The quantification of genetic diversity made it possible to select diverse parents for successful hybridization program. In recent years, several molecular markers had been used to identify and assess the genetic diversity and phylogenies relationship in plant. The traditional methods based on morphological traits require more time, cost expensive and has large effect on environment. By the development of a wide range of molecular technique, marker assisted breeding is now used to enhance conventional breeding program for crop improvement. Among the different molecular markers RAPD technique (Williams et al. 1990) is reliable, faster and easier for exploiting molecular diversity analysis within and among species. RAPD markers have been widely used for identification of genetic relationship among cultivars (Tosti and Nejri, 2002). Hence, the present investigation was undertaken for molecular diversity analysis of some released potato varieties through RAPD markers and to identify the divergence genotypes for potato improvement program.

## MATERIALS AND METHODS

The experiments were carried out at the Department of Biotechnology, Sher-e-Bangla Agricultural University, Sher-e-Bangla Nagar, Dhaka. Twelve popular released potato varieties were collected from Tuber Crops Research Centre (TCRC), Bangladesh Agricultural Research Institute (BARI), Gazipur and used as experimental materials. The potato varieties were Cardinal, Diamant, Granola, Asterix, Sagita, Courage, Lady Rosetta, Felsina, Multa, Provinto, Petronige and TPS-1.

**Seedling raising:** Good quality, disease free, healthy tuber (breeder stock) were sown in plastic pots and kept in nets house. All management practices were done for raising quality seedlings from those materials. Fresh leaves were collected at 3-4 leaf stage of plant for isolation of DNA.

**Extraction and quantification of DNA:** Total genomic DNA from each variety was isolated by CTAB method with slight modification according to Frey et al., (2004). The extracted DNA was purified by propanol and treated with 10 $\mu$ g/ml RNase A for 20-25 min at 37°C to remove the RNA. The purified DNA was dissolved in TE buffer and quantification of DNA was done through electrophoresis on 1% agarose gel staining by ethidium bromide. The sample DNA was stored at -20°C freezer for further use.

**Primer selection and PCR amplification:** Seventeen RAPD primers were selected on the basis of previous works to evaluate the molecular polymorphism among the potato varieties. PCR reaction was performed using BIONEER KIT

(korea). The PCR reaction having 20.0 µl mixture containing 3.0 µl sterile deionized water, 10X PCR buffer 4.0 µl, enzyme dilution buffer 4.0 µl, 20 mM MgCl<sub>2</sub>, 3.0 µl dNTPs (10mM) 1.0 µl top DNA polymerase 0.5 µl, primer 2.5 µl and sample DNA (approx. 40-50 ng) 2.0 µl. The reaction mixture was subjected to the following thermal profile for amplification in a thermocycler : 5 min at 95°C for initial denaturation, followed by 33 cycle of 1.10 min denaturation at 94°C, 1.0 min at annealing and 1.30 min at 72°C for extension. A final extension step was done at 72°C for 7 min. Electrophoresis was done to visualize the PCR amplified product. It was carried out on 2.0% agarose gel and amplified fragments were visualized by staining with ethidium bromide. The amplified bands were scored as present (1) and absent (0) for each primer. The score of bands were pooled to create a single data matrix. These were used to estimate polymorphic loci. Genetic distance and identity were calculated based on Nei's (1972). Phylogenetic tree and dendrogram were established based on an unpaired group method of arithmetic means (UPGMA) using the software POPGENE (Version 1.31) (Yeh et al., 1999).

## RESULTS AND DISCUSSION

Molecular diversity and polymorphism studies in 12 potato varieties of Bangladesh was done through RAPD primer. Seventeen RAPD Primers (10-mer) were initially screened on 12 popular potato varieties for their ability to amplify polymorphic fragment of DNA. Out of them only eight primers viz. OPA-17, OPG-17, OPJ-13, OPP-12, OPX-01, OPX-04 and OPX-07 showed distinct polymeric DNA profiles. Some total of 39 bands were obtained from these primers with an average of 4.87 bands per primer. Among the amplified product 24 polymorphic DNA bands were observed. The polymorphic DNA fragments ranged from 2-6 in different RAPD oligomer. It was observed that the primer OPX-04 product had highest (9) number of polymorphic DNA band and it was lowest (2) in OPA-17 and OPX-07 primers. The percent of polymorphic DNA fragment was 61.53 under this present investigation (Table 2). The maximum DNA fragment was generated by the primer OPX-04 and it was minimum (3) in OPA-17. The DNA profile of 12 potato varieties using OPX-04, OPX-07 and OPX-17 primers are shown in figure 1 and 2, respectively. The number of polymorphic bands was considered appropriate to assess the genetic divergence of potato genotypes. It might be due to more amount of GC content (60-70%) of the primers used in this study. Fukuoka et al. (1992) observed an increased number of bands with increasing GC content of the primer. The explanation for the correlation between GC content and the number of bands may be the stability of base complementation of A with T. The amplified DNA profiling was scored according to the presence and absence of bands. Absence of bands might be failure of primers to anneal at a binding site in some genotypes due to nucleotide sequence differences or may be insertion or deletions of primer binding site. Rocha et al. (2010) reported on genetic diversity in potato cultivar by RAPD and SSR markers. They notice that, genomic DNA of 16 potato cultivars was amplified with 25 RAPD primers that

generated 92 polymorphic bands. The cultivar identification using RAPD markers is well documented in studies of molecular characterization (Bianchi et al, 2003). Fingerprinting based on RAPD marker type was used for identification and characterization of potato cultivars in North America (Sosinski and Donches, 1996). The genetic identity and genetic distance among the 12 potato varieties are presented in table 2. The Nei's genetic identity was the highest (0.82050) in the varietal pair Cardinal and Diamant and it was the lowest (0.333) in Provinto and Granola. The highest Nei's genetic distance (0.970) was noticed between Granola and Provinto. It was the lowest (0.137) in two different varietal pair viz, (a) Petronige and Provinto (b) Courage and Provinto, respectively. However, high levels of genetic distance were also noticed in the varietal pairs: Lady Rossetta and Cordage (0.955), Provinto and Lady Rossetta (0.893), Courage and Granola (0.8910). A dendrogram based on Nei's (1972) genetic distance using unmeasured pair group method of arithmetic mean (UPGMA) was established with 12 popular potato varieties (Figure 3). These varieties segregated into two main clusters. The variety Granola and Sagita were into one cluster and rest of the materials were in cluster-II. The cluster -II was sub-divided into two sub groups. Seven varieties were clustered in one sub-group and three varieties were clustered in second sub-group. Those sub-groups were further segregated in different sub-sub cluster group on the basis of their identity. The results indicated that, low and high level genetic distance exists between the varieties. The variety Granola, Provinto, Lady Rossetta and Courage showed highest level of genetic diversity which can be used for further potato breeding program. Sawy et al. (2007) reported that, RAPD technique can be successfully applied to determine the genetic fidelity of potato plant. A limited study has been made on genetic divergence in potato either at tetraploid (Gaur et al. 1978 and Sidhu et al., 1981 or at diploid level (Grag 1988). Mondal (2007) reported that an understanding of the nature and magnitude of variability among the genetic stock is of prime importance to the breeders. Hence, it is important to analyze the genetic variability of parental materials. Molecular based analysis of present finding can provide information on actual genetic diversity among the potato cultivars.

### CONCLUSION

Information on the genetic diversity allows to assist the parent selection and paving the way to genetic gains. The results of present study revealed the existence of high level of genetic diversity among the studied 12 popular widely grown potato varieties in Bangladesh. These varieties can further be used as parental material for fixation of heterosis in potato improvement program.

### ACKNOWLEDGEMENT

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#### PCR Amplification with RAPD Primer OPX-07

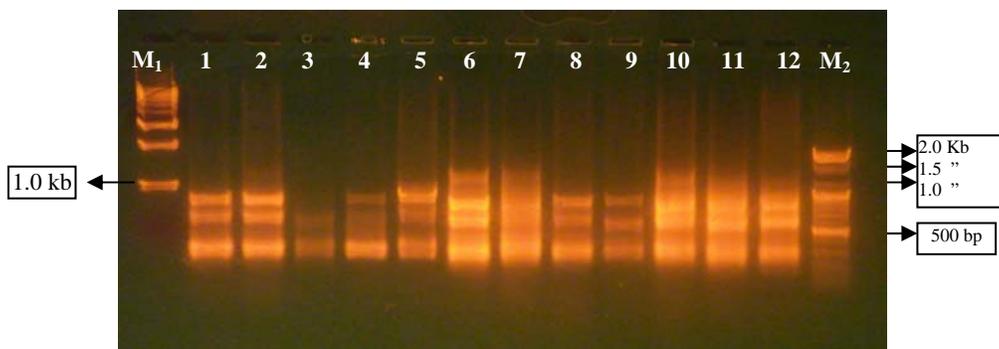


Figure 1: RAPD profile of 12 potato varieties using primer OPX-07. Lane: 1. Cardinal; 2. Diamant; 3. Granola; 4. Lady Rossetta; 5. Sagita; 6. Courage; 7. Asterix; 8. Felsina; 9. Multa; 10. Provinto; 11. Petronige; 12. T.P.S. 1; M<sub>1</sub>= Molecular marker 1kb (B.G. Nei, India) and M<sub>2</sub>=100bp (Bioneer, Korea)

#### PCR Amplification with RAPD Primer OPX-04

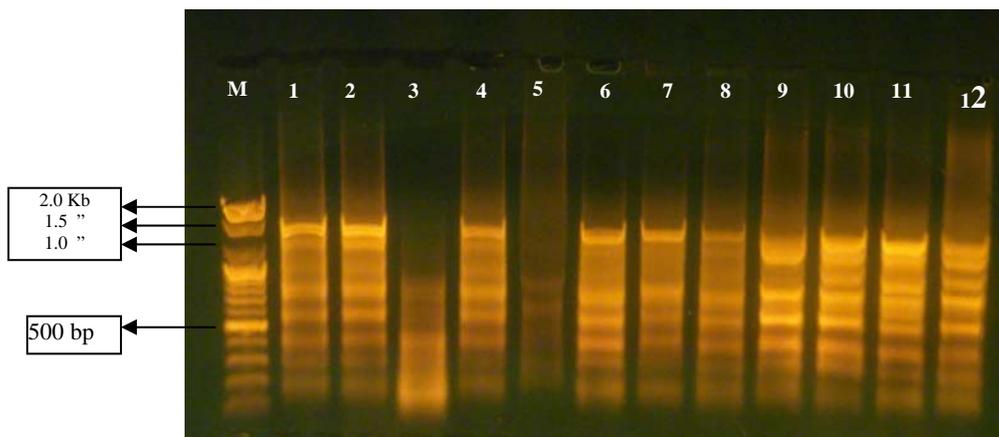


Figure 2: RAPD profile of 12 potato varieties using primer OPX-04. Lane: 1. Cardinal; 2. Diamant; 3. Granola; 4. Lady Rossetta; 5. Sagita; 6. Courage; 7. Asterix; 8. Felsina; 9. Multa; 10. Provinto; 11. Petronige; 12. T.P.S. 1; M= Molecular marker 100bp (Bioneer, Korea)

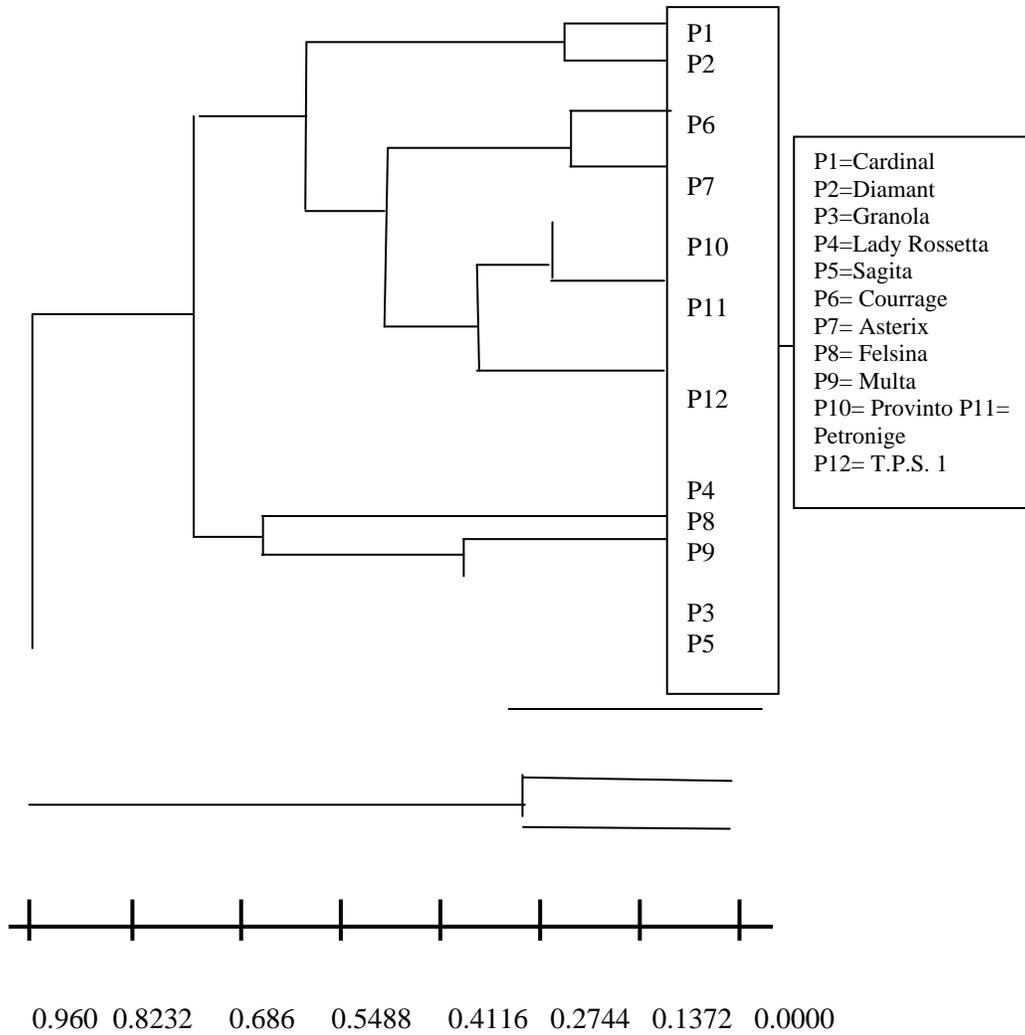


Figure 3: UPGMA dendrogram based on Nei's (1972) genetic distance, between 12 potato varieties according to RAPD analysis.

**Table 1: Number and percentage of polymorphic loci obtained in 12 potato varieties**

Name of RAPD primer	Sequence of the primer	GC content (%)	No. of bands scored	Size ranges (bp) observed	No. of polymorphic bands	Percentage of polymorphic loci
OPA17	GACCGCTTGT	60	3	278-735	2	66.66
OPG17	ACGACCGACA	60	6	249-1225	3	50.0
OPJ13	CCACACTACC	60	5	295-1491	4	80.0
OPP12	AAGGGCGAGT	60	5	400-1264	3	60.0
OPX01	CTGGGCACGA	70	7	144-934	4	57.14
OPX04	CCGCTACCGA	70	9	198-1579	6	66.66
OPX07	GAGCGAGGCT	70	4	301-1272	2	50.00
Total	-	-	39	-	24	61.53

**Table 2: Genetic identity (above diagonal) and genetic distance (below diagonal) values among the twelve potato varieties**

	Cardinal	Diamant	Granula	Asterix	Lady Rossetta	Courage	Sagita	Felsina	Multa	Provinto	Petronige	TPS1
Cardinal	-	0.8205	0.5897	0.6154	0.5641	0.7692	0.6923	0.6410	0.5641	0.7436	0.6667	0.7692
Diamant	0.1978	-	0.6667	0.7436	0.5897	0.6410	0.5641	0.4615	0.5385	0.6667	0.7436	0.6923
Granula	0.5281	0.4055	-	0.6154	0.7692	0.4103	0.4359	0.5385	0.5641	0.3333	0.4615	0.4615
Asterix	0.4855	0.2963	0.4855	-	0.5897	0.5897	0.5641	0.6154	0.6923	0.6154	0.7436	0.6410
L.Rossetta	0.5725	0.5281	0.2624	0.5281	-	0.3846	0.4615	0.6667	0.6923	0.3590	0.4872	0.5385
Courage	0.2624	0.4447	0.8910	0.5281	0.9555	-	0.8718	0.6154	0.5897	0.8718	0.7436	0.6923
Sagita	0.3677	0.5725	0.8303	0.5725	0.7732	0.1372	-	0.7436	0.6154	0.8462	0.7179	0.6667
Felsina	0.4447	0.7732	0.6190	0.4855	0.4055	0.4855	0.2963	-	0.7692	0.6410	0.5641	0.6667
Multa	0.5725	0.6190	0.5725	0.3677	0.3677	0.5281	0.4855	0.2624	-	0.6667	0.7436	0.6923
Provinto	0.2963	0.4055	0.9701	0.4855	0.8931	0.1372	0.1671	0.4447	0.4055	-	0.8718	0.8205
Petronige	0.4055	0.2963	0.7732	0.2963	0.7191	0.2963	0.3314	0.5725	0.2963	0.1372	-	0.7949
TPS1	0.2624	0.3677	0.7732	0.4447	0.6190	0.3677	0.4055	0.4055	0.3677	0.1978	0.2296	-

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## IDENTIFICATION AND DISTRIBUTION OF SUGARCANE STEM BORER IN BANGLADESH

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### ABSTRACT

Field surveys were conducted during the cropping season of 2010-2011 to assess the distribution of Sugarcane stem borer species in 12 AEZs of Bangladesh. These surveys documented abundance and their distribution and results clearly showed the existence of the stem borer at all locations surveyed, but with a higher incidence in the Atwary (36%) and initiation of infestation was observed on 20 May. Stem borer incidence and distribution varied significantly among the different locations. Second highest rate of infestation (32%) was recorded in Bashudebpur followed by Dinajpur (31%), Pabna and Akandabaria farm (30%). The lower infestation was recorded in Kaliganj. The percentage of stems attacked at the Kaliganj has never exceeded 23% followed by Rajshahi (28%), Thakurgaon and Faridpur (29%). The rate of infestation of stem borer (*Chilo tumidicostalis*) in different locations varied from 23-36%. While morphological characteristics of stem borer species were identified with standard keys and species composition was only predominated by *Chilo tumidicostalis* Hampson, though previous workers found other borers in addition to this species. The sex ratio of adult moth *Chilo tumidicostalis* was 1:1.42 after emergence from the reared collected pupae from different locations.

**Key words:** Alternate host, Caterpillar growth stage, Moths, Pupae, Lepidopteran pest, Sugarcane.

### INTRODUCTION

Sugarcane (*Saccharum officinarum* L.), a perennial tropical crop with a high self-tolerant nature, is grown for sugar stored in its stem and propagated through stem cuttings. Sugarcane is grown around the world between tropical and sub-tropical

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climate (North latitude = 35°C and South latitude = 35°C). Sugarcane is cultivated in more than 100 countries of the tropical and sub-tropical regions of the world (Humbert, 1968). Around 70% of the world's sugar is produced from this crop (Chowdhury and Vasil, 1993). Sugarcane is cultivated on an area of about 0.16 million hectare of land which almost 50% area is located in the sugar mills zone, and the remaining 50% is grown in the non-mills zone, where sugarcane is mostly diverted for jaggary and juice production. Presently, 15 sugar mills are in operation under Bangladesh Sugar and Food Industries Corporation (BSFIC) with a capacity of 0.21 million tons of sugar production per year (BSFIC, 2008).

Stem borer, *Chilo tumidicostalis* Hampson is a serious insect pest of sugarcane in Bangladesh (Karim and Islam, 1977), Thailand (Suasa-ard et al., 2000), Nepal, Burma (Williams et al., 1969) and Australia (Sallam, 2006). Incidence of stem borer infestation starts from the end of April and continues up to November. However, it remains most abundant from June to September. Hundred percent crop losses by stem borer infestation were recorded in Setabgonj Sugar Mills in Bangladesh during 1973-1974 cropping season when attack reached as high as 100% due to the stem borer infestation (Karim and Islam, 1977). In India, it caused 8.2-12.6% yield loss sometimes reaching up to 70% with 10.75-48.55 % sugar recovery in endemic areas (Khanna et al., 1957; Butani, 1961). Gupta and Avasthey (1960) also reported 25-70% and 12-60% in cane yield due to primary and secondary infestation, respectively in different sugarcane varieties, along with 0.5-3.5 unit loss of sugar recovery in West Bengal, India. Stem borer is the most destructive pest of sugarcane in growing areas of Bangladesh. Maximum weight losses of 28.73, 18.64 and 18.01% and sucrose losses of 9.74, 11.21 and 15.93% were estimated in Isd 16, Isd 21 and Isd 30 respectively having more than three bores in infested cane (BSFIC, 2008).

The stem borer genus *Chilo* contains 41 species of which *Chilo tumidicostalis* had been so far identified worldwide attacking many crops particularly Graminae/Poaceae family. This species and *Chilo auricilius* are very common and destructive in Bangladesh and India (Khanna et al., 1957; Avasthy, 1983). In addition, internode borers, *Chilo sacchariphagus indicus* Kapur is also a major pest of sugarcane in Peninsular India (Gupta and Avasthy, 1957). Climatic and soil conditions of both Bangladesh and India are almost similar. Besides, availability of alternate hosts during off season facilitates over-wintering of stem borer in Bangladesh. Therefore, there is a high possibility of invasion of the pest in Bangladesh. In such situation, distribution and identification of this pest at certain intervals are essential to know whether any new pest species has invaded in this country. In fact, no comprehensive work has so far been undertaken in Bangladesh to identify the different species of stem borer attacking sugarcane cultivated throughout the country since 1962. In this context, the present study was undertaken to identify and to assess the distribution of sugarcane stem borers in Bangladesh.

## MATERIALS AND METHODS

### Study site

The survey was conducted at nine locations covering 12 Agro-Ecological Zones (AEZ) of Bangladesh with date of sample collection, date of placing sample in rearing box in the laboratory (Table 1). Stem borer infested sugarcane were collected from the selected locations from August to September 2011. Nine Sugar Mills zones of Bangladesh namely Atwary, Panchagar Sugar Mills (PSM); Patuadangi farm, Thakurgaon Sugar Mills (TSM); Sultanpur farm, Setabganj Sugar Mills (STSM); Puthia, Rajshahi Sugar Mills (RJSM); Bashudebpur, Natore Sugar Mills (NTSM); BSRI farm, Pabna Sugar Mills (PBSM); Akandabaria farm, Carew & Co. Sugar Mills; Modhukhali, Faridpur Sugar Mills (FSM) and Kaliganj, Mobarakganj Sugar Mills (MKSM) were selected for this study (Figure 1).

### Collection of sample

A comprehensive survey was conducted through questionnaire by extensive visit throughout the selected locations during the cropping season of 2010-2011 and stem borer infested sugarcane fields were observed in the morning and afternoon. Adult emergence of *Chilo tumidicostalis* from pupae was recorded by rearing the collected specimen of 9 locations during the period from August to October 2011 and periods of stem borer infestation were recorded for nine locations of Bangladesh during the period from May to June 2011 (Figure 2 & 3). A total of 100 infested sugarcane (Plate 6) were checked for recording the incidence of target pest in each field of cane grower. Hundred plants were examined randomly at each sugarcane field to obtain the percent infestation of stem borer (Figure 4). At each location, three sugarcane fields were randomly selected from which specimens were collected at random from 3 different cane growers. The percent of stem borer infested sugarcane per location was calculated based on the number of total plants observed and the number of infested plants. Accordingly fifty (50) plant specimens from each location were cut and tied into bundle. The bundle of cane containing borers was brought to Entomology Laboratory of Regional Sugarcane Research Station (RSRS), Thakurgaon as well as Entomology Laboratory of Bangladesh Sugarcane Research Institute (BSRI), Ishurdi, Pabna for morphometric study. Rearing boxes were kept ready before the specimen were carried to the laboratory.

### Rearing of sugarcane stem borer

Infested sugarcane collected from different locations were placed separately in netted rearing cage (90 cm × 60 cm) and placed on laboratory desk. In order to culture the specimen larvae were collected by splitting borer infested stem of sugarcane. They were reared in the plastic boxes (23.0 cm in dia. and 10.5 cm high) with pieces of sugarcane stalk as food until pupation. The pupae were kept in a Petri dish (11.5 cm × 1.5 cm) with adequate moisture provided with water-soaked filter paper in the bottom of the Petri dish until the adult emergence. Pupa were then transferred to the

insect rearing cage (60 cm × 60 cm × 90 cm). Stock culture of stem borer was maintained at room temperature ranging from 20-30°C. Adults emerged from full grown pupae at 7-15 days after pupation. Emerged adults were placed separately in a killing jar and their number was counted. Adult moths were then pinned, stretched and preserved for identification.

#### **Identification of sugarcane stem borer (*Chilo tumidicostalis* Hampson)**

The major characteristics used to identify the adult male and female of sugarcane stem borer at different stages are given in appendix I. Specimens were primarily identified with the help of taxonomic keys described by Butani (1956), Alam (1967) and Butani and Jotwani (1984) (Appendix II). The laboratory reared freshly emerged adults (moths) collected from nine locations were compared with adult characteristics (Plate 1-6). For this purpose specific characters of each specimen were studied thoroughly and checked with the characters of the keys and all the characters were studied under binocular microscope. Larval (caterpillar) characteristics were also considered to confirm the identification.

**Design and Statistical analysis:** The experiment was laid out in Completely Randomized Design (CRD) with three replications. The treatment means were compared using Duncan's Multiple Range Test i.e. DMRT (Gomez and Gomez, 1984).

## **RESULTS AND DISCUSSION**

Infested samples with larvae collected from different locations, number of pupae and number of total adult individuals emerged from the laboratory-reared sugarcane are presented in Table 2. The recently emerged adults collected from nine locations of Bangladesh belong to the genus *Chilo* under the family Pyralidae. In the present study through identifying characteristics keys as described by Butani (1956), it might be assumed that only *Chilo tumidicostalis* would be the major pest of sugarcane in all the nine locations of Bangladesh. The present results are supported by Isaac and Rao (1941) and Isaac and Venkataraman (1941) who had shown the same larval and pupal characters of stem borer.

Morphological variations were not observed among the emerged adults of both male and female. The results indicated that the morphological characteristics of larva, pupa and adults were completely comparable to that of *Chilo tumidicostalis* only. Thus, only one species *tumidicostalis* belonging to the genus *Chilo* under the family Pyralidae was identified. Avasthy (1983) reported that two species of *Chilo* (i.e., *Chilo tumidicostalis* and *Chilo auriciliusi*) had bored into sugarcane, corn and paddy. He observed that these species also had attacked other plants under the family Graminae/Poaceae. Kapur (1950) reported that *Argyria tumidicostalis* later renamed as *Chilo tumidicostalis* Hampson was the major borer pest of sugarcane which prevails in Bangladesh.

**Number of pupae emerged**

The highest number of pupae (73) was produced from larvae of infested sugarcane collected from Patuadangi farm, Thakurgaon (Table 2). The second highest number of pupae (70) was obtained from the samples with larvae collected from Sultanpur farm, Setabganj. On the other hand, the lowest number of pupae (56) was found from the collected samples with larvae of Akandabaria farm, Carew & Co., Dorshona. Almost similar number of pupae viz., 60, 62, 65, 67, 68 and 69 were recorded from the sugarcane samples (with larvae) of Kaliganj, BSRI farm, Pabna, Puthia, Modhukhali, Bashudebpur and Atwary, respectively.

**Number of Adults emerged**

From the Figure 2 the highest number of adults (50) emerged from the pupae were recorded from Patuadangi farm followed by Atwary, Sultanpur farm and Modhukhali (49). The lowest number of adults (45) emerged from the pupae of Akandabaria (Carew & Co.) followed by BSRI farm (46). The number of adults emerged from the pupae of Puthia, Kaliganj and Bashudebpur were 47, 47 and 48, respectively.

**Male and female Sex Ratio**

The numbers of male and female stem borer moths emerged from the pupae of different locations are presented in Table 2. The highest number (24) of male moths emerged from the pupae of Patuadangi farm and the second highest number (22) of male moths emerged from Sultanpur farm. Number of male moths of 21 and 20 emerged from the pupae of the samples was collected from Atwary and Bashudebpur, respectively. Male moth emergence was poor from the pupae of samples collected from Modhukhali (19), Puthia (18), BSRI farm (18), Kaliganj (18) and Akandabaria farm (17). The highest number (30) of female moths emerged from the pupae of the samples from Modhukhali and the second highest number (29) from the samples of Puthia and Kaliganj followed by Atwary (28), Bashudebpur (28), BSRI farm (28) and Akandabaria farm (28). Twenty seven and twenty six female moths emerged individually from the pupae of samples collected from Sultanpur farm and Patuadangi farm. The number of female moth emergence ranged from 26-30. The lowest number (26) of female moth emerged from the pupae of samples collected from Patuadangi farm. A total of 430 moths emerged from the collected specimens of 9 different locations. Of them, 177 were males and 253 were females with a sex ratio of 1:1.42.

**Periods and rate of infestation by *Chilo tumidicostalis* in different locations**

From the figure 3 the initiation of infestation by stem borer was observed on 20 May in Atwary, 22 May in Patuadangi farm and 25 May in Sultanpur farm. On the other hand, stem borer infestation was found on 10 June in Rajshahi, 15 June in Natore, 18 June in Faridpur, 16 June in Pabna, 14 June in Akandabaria farm and 20 June in Mobarakganj. Panchagor, Thakurgaon and Dinajpur mills zones were the most *Chilo tumidicostalis* Hampson prone areas in Bangladesh. These zones have medium high

land, sandy and sandy loam soil texture, rain fed in March, April and May but were rainy in June, July and August because these areas are closer to Dargilling and Himaloy. Sugarcane is a major crop in these areas covering 14-18 months in the field. Maize and Rice is the alternate/sequential crop which is alternate host of sugarcane stem borer. In these areas, sugarcane is planted in the early season. For this reason, stem borer infestation reached as high as 100% in Sultanpur farm and Dinajpur (Karim and Islam, 1977).

#### **Survey record on the infestation rate of stem borer in different locations**

From the figure 4 survey report indicates that the highest rate of infestation (36%) was observed in Atwary. Second highest rate of infestation (32%) was recorded in Bashudebpur followed by Dinajpur (31%), Pabna and Akandabaria farm (30%). The lower infestation was recorded in Kaliganj (23%), followed by Rajshahi (28%), Thakurgaon and Faridpur (29%). The rate of infestation of stem borer (*Chilo tumidicostalis*) in different locations varied from 23-36%.

#### **Distribution of stem borer on host and locality**

Stem borer, *Chilo tumidicostalis* (Hampson) was the most abundant species which recorded 430 individuals that emerged from 590 pupae (72.88%) reared in infested sugarcane collected from 9 different locations situated at North, South, West and Central regions of Bangladesh. These species predominantly emerged from the specimens collected from Thakurgaon but became gradually infrequent in the infested sugarcane collected from Dinajpur, Panchagor, Natore, Faridpur, Rajshahi, Pabna, Kaliganj and Dorshona (Figure 3). The present study indicated that *Chilo tumidicostalis* is the most damaging insect pest that bored into the sugarcane, though previous workers found other borers in addition to *Chilo auricilius*. Other borers which attack maize and rice choose sugarcane as alternate host. A research report was found in the web site that there were many species of *Chilo* so far identified worldwide which attack many crops particularly under Graminae/Poaceae family. The genus *Chilo* contains 41 species and of them, 8 species (i.e., *Chilo agamemnon*, *C. auricilius*, *C. infuscatellus*, *C. orichalocociliellus*, *C. partellus*, *C. sacchariphagus*, *C. terrenellus* and *C. tumidicostalis*) have damage potential in Australian cane if they invade the mainland. Additionally, two *Chilo* species of Australia (i.e., *Chilo polychrysus* and *C. suppressalis*) are major pest of rice and minor pest of sugarcane in some countries in Asia (David and Easwaramoorthy, 1990). However, *Chilo suppressalis* appears to be strictly a pest of rice, but there is no evidence in the literature that it can survive on sugarcane. Two other species (*Chilo diffusilineus* and *C. zacconius*) were found in Africa but might have negligible impact. The remaining 29 *Chilo* genera are not known to be the pests of sugarcane.

Stem borer, *Chilo tumidicostalis* (Hampson) is originally an Asian species. Populations in Madagascar, Malaysia, Mauritius and Reunion have probably been introduced by human in the mid 1800s (Williams, 1983). *Chilo tumidicostalis* is reported to feed exclusively on sugarcane, found in Bangladesh, India, Myanmar, Nepal and Thailand (David and Easwaramoorthy, 1990; Suasa-ard et al., 2000). It is known as the Bengal borer in India and used to be considered a major pest of

sugarcane in Purnea and adjoining parts of Bhagalpur, Munger and Darbhanga districts of Bihar, but its importance seems to have declined during the 1980s (Kumar and Kalra, 1987). However, in Thailand, *C. tumidicostalis* used to be considered a major pest of sugarcane until the late nineties, when it unexpectedly became the most important pest of sugarcane. Severe outbreaks were reported in the provinces of Sa Kaew and Buri Rum in Thailand (Suasa-ard et al., 2000) and Dinajpur, Setabganj in Bangladesh (Karim and Islam, 1977) where infestation reached up to 100%. However, this insect pest should have a high spread and colonization potential in sugarcane growing areas, especially in North Queensland, Taiwan and Pakistan (Sallam and Allsopp, 2002; Cheng et al., 1997; Ullah et al., 2006).

### CONCLUSION

The present study indicated that *Chilo tumidicostalis* Hampson is the only insect pest that bored into sugarcane, though previous workers found other borers in addition to *Chilo tumidicostalis*. Other borers which attack sugarcane are the rice, maize borer and they choose sugarcane as alternate host. However, it is now essential to undertake more research work to find out other borers which may have been attacking sugarcane as alternate host. Another study could be conducted to identify some of the borers invading Graminae/Poaceae crops of Bangladesh.

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**Table 1. Name of locations, date of collection and placement of sugarcane samples in rearing boxes in the Entomology Laboratories**

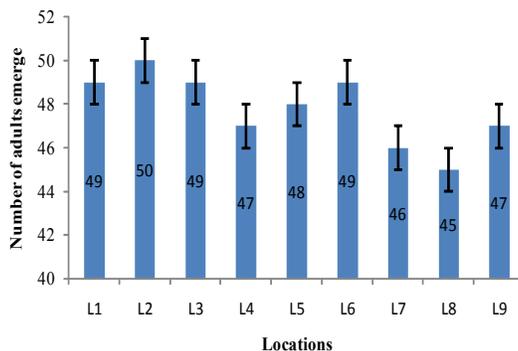
Selected locations	Date of sample collection	Date of sample placement in rearing box	Laboratory
Atwary, Panchagor Sugar Mills (PSM)	24 August 2011	25 August 2011	RSRS, Thakurgaon
Patuadangi farm, Thakurgaon Sugar Mills (TSM)	26 August 2011	27 August 2011	RSRS, Thakurgaon
Sultanpur farm, Setabganj Sugar Mills (STSM)	25 August 2011	26 August 2011	RSRS, Thakurgaon
Puthia, Rajshahi Sugar Mills (RJSM)	10 September 2011	11 September 2011	BSRI, Ishurdi, Pabna
Bashudebpur, Natore Sugar Mills (NTSM)	11 September 2011	12 September 2011	BSRI, Ishurdi, Pabna
Modhukhali, Faridpur Sugar Mills (FSM)	12 September 2011	13 September 2011	BSRI, Ishurdi, Pabna
BSRI farm, Pabna Sugar Mills (PBSM)	17 September 2011	18 September 2011	BSRI, Ishurdi, Pabna
Akandabaria, Carew & Co. Sugar Mills (Carew & Co.)	18 September 2011	20 September 2011	BSRI, Ishurdi, Pabna
Kaliganj, Mobarakganj Sugar Mills (MKSM)	19 September 2011	21 September 2011	BSRI, Ishurdi, Pabna

**Table 2. The number of stem borer moth emerged from infested sugarcane samples from Nine sugar Mills of Bangladesh**

Selected locations	Number of observed samples	Number of pupae produced	Number of adult emerged	♂ and ♀ ratio of <i>Chilo tumidicostalis</i>	
				♂	♀
Atwary, Panchagor Sugar Mills (PSM)	100	69	49	21 (30.43)	28 (40.58)
Patuadangi farm, Thakurgaon Sugar Mills (TSM)	100	73	50	24 (32.88)	26 (35.62)
Sultanpur farm, Setabganj Sugar Mills (STSM)	100	70	49	22 (31.43)	27 (38.57)
Puthia, Rajshahi Sugar Mills (RJSM)	100	65	47	18 (27.69)	29 (44.62)
Bashudebpur, Natore Sugar Mills (NTSM)	100	68	48	20 (29.41)	28 (41.18)
Modhukhali, Faridpur Sugar Mills (FSM)	100	67	49	19 (28.36)	30 (44.78)
BSRI farm, Pabna Sugar Mills (PBSM)	100	62	46	18 (30.00)	28 (46.67)
Akandabaria, Carew & Co. Sugar Mills (Carew & Co.)	100	56	45	17 (30.36)	28 (50.00)
Kaliganj, Mobarakganj Sugar Mills (MKSM)	100	60	47	18 (29.03)	29 (46.77)
Total	900	590	430	177	253

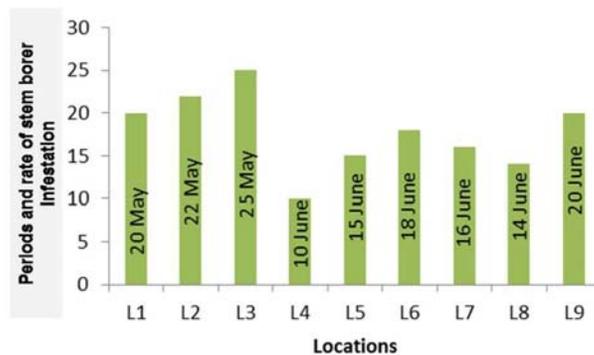
Male : Female = 1 : 1.42





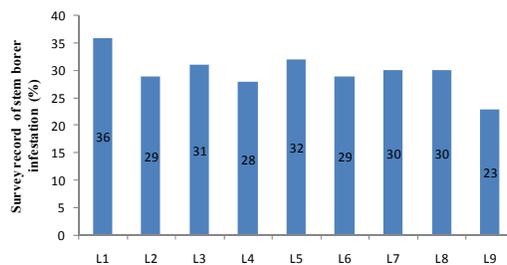
L<sub>1</sub> = PSM, L<sub>2</sub>= TSM, L<sub>3</sub>= STSM, L<sub>4</sub>= RJSM, L<sub>5</sub>= NTSM, L<sub>6</sub>= FSM, L<sub>7</sub>= PBSM,  
 L<sub>8</sub>= Carew & Co. and L<sub>9</sub>= MKSM, Standard bar indicates standard error.

Figure 2. Adult emergence from the pupae of *Chilo tumidicostalis* reared from the collected specimens of nine locations of Bangladesh during August-October, 2011



L<sub>1</sub> = PSM, L<sub>2</sub>= TSM, L<sub>3</sub>= STSM, L<sub>4</sub>= RJSM, L<sub>5</sub>= NTSM, L<sub>6</sub>= FSM, L<sub>7</sub>= PBSM,  
 L<sub>8</sub>= Carew & Co. and L<sub>9</sub>= MKSM

Figure 3. Periods and rate of infestation by stem borer in nine locations of Bangladesh during May-June, 2011 ( $P < 0.01$ ).



L<sub>1</sub> = PSM, L<sub>2</sub>= TSM, L<sub>3</sub>= STSM, L<sub>4</sub>= RJSM, L<sub>5</sub>= NTSM, L<sub>6</sub>= FSM, L<sub>7</sub>= PBSM,  
 L<sub>8</sub>= Carew & Co. and L<sub>9</sub>= MKSM

Figure 4. Mean percent infestation by stem borer in nine locations of Bangladesh during the cropping season of 2010-2011 ( $P < 0.01$ ).



Plate 1. Seven spots present in terminal side of adult *Chilo tumidicostalis*



Plate 2. Two – five tier eggs of *Chilo tumidicostalis*

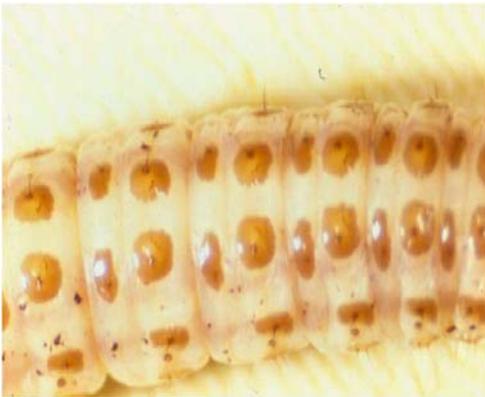


Plate 3. Four pinkish brown stripes (mature larva)

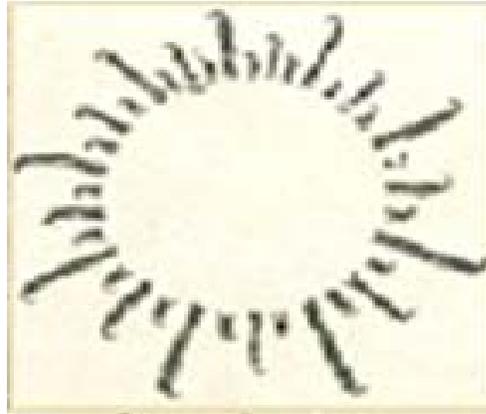


Plate 4. Complete crochets on planta of prolegs in triple series



Plate 5. The 5<sup>th</sup>, 6<sup>th</sup>, and 7<sup>th</sup> abdominal segments chitinized and branched spines



Plate 6. Entry wholes by stem borer in sugarcane

### Appendix I. Characteristics of egg, larva, pupa and adult of sugarcane stem borer

#### Egg stage

Behaviour of egg laying	Time of egg laying	Oviposition period (day)	Number of eggs in total life	Egg size	Egg colour
On 1 <sup>st</sup> , 2 <sup>nd</sup> and 3 <sup>rd</sup> top leaves, leaf sheath in rows of 2-5 tiers (Plate 2).	Evening	4	500-800 and eggs are uncovered	Length: 1.2 mm Width: 0.8 mm	Dirty white with light greenish tinge but reddish at hatching

#### Larval stage

Crochet on the prolegs	Average larval period (days)	Number and colour of stripes	Number of instar	Average larvae size	Larval colour
Complete crochets on planta of prolegs are arranged in triple series (triordinal) (Plate 4).	27-70	Four pinkish brown stripes are alike (Plate 3).	5	Young: 1-3 mm Mature: 3 mm	White with black/orange head, later stage creamy. Dark brown spots bearing hairs on body segments.

#### Pupal stage

Characteristics	Average pupal period (days)	Average pupal size	Pupal colour
1. Front of fronto-clypeal region extends upward forming hard ridge like projections. 2. The 5 <sup>th</sup> , 6 <sup>th</sup> and 7 <sup>th</sup> abdominal segments with prominent hard and branched spines (Plate 5).	7-15 up to 22 days in cool weather	Long: 16 – 20 mm	Brown

#### Adult stage

Behaviour of egg laying	Adult longevity (day)	Adult colour	Width at wingspan		Difference at ♀ and ♂	
			♀ (mm)	♂ (mm)	♀	♂
Nocturnal	3-8	Yellowish with narrow brown lines on forewing (Plate 1).	25-40	18-30	Anal part U-shaped	Thick hair on anal part

**Appendix II. Standard Key for Identification of sugarcane borer pests**

Particulars	Stalk borer ( <i>Chilo auricilius</i> )	Plassey borer ( <i>Chilo tumidicostalis</i> )	Internode borer ( <i>Chilo sachariphagous indicus</i> )
Number of stripes	Five: Almost alike	Four: Almost alike	Four: Almost alike
Colour of stripes	Violet	Pinkish brown	Violet
Colour of tubercles	Grey	Grey	Jet black
Crochet on the prolegs	Complete crochets with biordinal spines in the prolegs	Complete crochets with triordinal spines in the prolegs	Complete crochets with triordinal spines in the prolegs
Egg	Egg laid in egg mass of 2-5. Total of 47-146 eggs.	Egg lay in 4-5 tiers. Total of 500-800 eggs. Eggs are uncovered.	Egg laid cluster in 2-3 parallel rows. Total eggs of few to 414.
larvae	Head black	Black/ orange colour head	Head black
Adult	Forewing straw colour with golden spot. Hindwing straw colour with silvery fringe	Forewing dark brown scales. Terminal series of black spots present. Hindwing light brown scales. Seven spots present in terminal side.	Straw colour with slightly dark spot on forewing.

According to Butani (1956)

## COMBINING ABILITY AND GENE ACTION IN INDIGENOUS BITTER GOURD (*Momordica charantia* L.)

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### ABSTRACT

The investigation was carried out to get information regarding magnitude of combining ability and nature of gene action for fruit yield and several other yield attributing traits following line × tester mating design involving 12 lines and 3 testers and their 36 hybrids tested in two environments viz., summer and rainy season 2010 at Research Farm of Department of Vegetable Science, N. D. University of Agriculture & Technology, Kumarganj, Faizabad, UP, India. From the estimate of gca effects, among the parental lines NDBT-13, NDBT-15 and NDBT-19 were identified as superior donor for both seasons and NDBT-10 for summer season and among the testers Kalyanpur Sona for summer season and Pusa Do Mausami for rainy season for fruit yield per plant and its yield contributing traits like number of fruits per plant and average fruit weight. Eight crosses displayed desirable significant sca effects in both seasons for fruit yield per plant. Among these eight crosses the best cross combinations based on desirable sca effects for fruit yield per plant were NDBT-19 × Pusa Do Mousami in summer season while NDBT-8 × Pusa Do Mousami, NDBT-15 × NDBT-12 and NDBT-10 × Pusa Do Mousami in rainy season. These crosses have more number of fruits per plant, average fruit weight, fruit diameter and other component traits in both seasons.

**Key words:** Gene action, General combining ability (gca), Hybrids, Line × Tester mating design, *Momordica charantia* L., RBD, Specific combining ability (sca)

### INTRODUCTION

Bitter gourd (*Momordica charantia* L.) is one of the most nutritive and commercially important vegetable grown throughout the country. The importance of bitter gourd has been recognized due to its high nutritive value and medicinal

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properties. Bitter gourd still remains an unexploited crop from genetic and breeding point of view. In bitter gourd, Indian variability is quite distinct than that of African/S.E. Asian region (Seshadri and Chatterjee, 1996). Wide range of variability in respect of vegetative and fruit character is available. Many varieties and  $F_1$  hybrids have been developed utilizing those variability. It is a monoecious and highly cross pollinated crop and has been known to offer good potentialities for increased yield. Considering these facts, it is essential and desirable to carry out a successful breeding programme utilizing the land races available in Indian subcontinent. Therefore, this study was conducted to generate information about general and specific combining ability effects for different economic characters.

### MATERIALS AND METHODS

The present investigation involving 36  $F_1$ s (derived through line  $\times$  tester mating design) and their parents i.e., 12 lines (NDBT-1, NDBT-2, NDBT-3, NDBT-4, NDBT-5, NDBT-6, NDBT-7, NDBT-8, NDBT-10, NDBT-13, NDBT-15 and NDBT-19) and 3 testers (Kalyanpur Sona, NDBT-12 and Pusa Do Mausami) was undertaken for estimating the gca (general combining ability) and sca (specific combining ability) for the 14 characters. Accordingly the experiment was set up in a complete randomized block design (RBD) with three replications in two seasons, namely summer (S) season and rainy (R) season 2010 at the Research Farm of Department of Vegetable Science, ND University of Agriculture & Technology, Kumarganj, Faizabad, UP, India. Sowing was done with a 3.0 m for row to row and, 0.5 m plant to plant spacing was maintained. All the recommended agronomic package and practices, plant protection measures were followed to raise a good and healthy crop. Among the 14 characters the observations were recorded on plot basis for node number to anthesis of first staminate flower, node number to anthesis of first pistillate flower, days to anthesis of first staminate flower, days to anthesis of first pistillate flower and days to first fruit harvest, where as for fruit length (cm), fruit diameter (cm), average fruit weight (g), number of fruits per plant, fruit yield per plant (kg), number of primary branches per plant, number of nodes per plant, internodal length (cm) and vine length (m) data were recorded on 10 randomly selected plants. The estimation of combining ability and gene action were carried out as per the procedure of Kempthorne (1957).

### RESULT AND DISCUSSION

The estimates of sca variances were found to be greater than corresponding gca variances for all the characters in both summer and rainy seasons indicating preponderance of non-additive gene action (Table 1). The ratio of gca to sca variances (average degree of dominance) was observed to be more than unity for all the characters (revealing over dominance) except for node number to anthesis of first pistillate flower in summer and for fruit yield per plant in rainy season (nearly equal to unity, suggests existence of complete dominance) indicating that predominance of

non-additive gene effects representing non-fixable dominance and epistatic components of genetic variance. This indicated that maintenance of heterozygosity would be highly fruitful for improving the characters. These results are in accordance with Singh et al. (2006) who reported non additive type of gene action for yield and yield related traits in their material. However Mishra et al. (1994) found that both additive and non additive gene actions were involved in the expression of yield and yield related characters. Similar gene actions were also observed by Devadas and Ramadas, 1997, Ram et al., 1999 and Jadhav et al., 2010. The differences in the results might have been due to the differences in the genetic material studied.

### **General combining ability**

As per table 2, the general combining ability of parents indicated that four lines i.e. NDBT-10, NDBT-13, NDBT-15 and NDBT-19 had significant and positive gca effects for fruit yield per plant during both seasons and among the testers, Kalyanpur Sona and PDM had significant and positive gca effects during summer and rainy seasons, respectively for this trait. Furthermore, these four lines were found to be good general combiners for number of fruits per plant during both seasons except NDBT-15, which showed negative effect in summer. Among the 12 lines, two lines NDBT-13 and NDBT-19 were found to have significantly positive gca effect for majority of the characters in both seasons like days to anthesis of first pistillate flower, days to first fruit harvest, number of fruits per plant, fruit yield per plant number of nodes per plant, inter-nodal length and vine length. The line NDBT-13 was also good combiner for node number of first staminate flower (significantly negative), fruit length, fruit diameter and average fruit weight (significantly positive) during both seasons. The line NDBT-15 was also found to be a good combiner for days to anthesis of first staminate flower, average fruit weight and inter-nodal length during both the seasons and also found to be a good combiner for days to anthesis of first pistillate flower, number of fruits per plant during rainy season along with node number to anthesis of first pistillate flower in summer season. The line NDBT-2 was found good combiner for earliness and other desirable traits like; node number to anthesis of first staminate flower, node number to anthesis of pistillate flower, days to anthesis of first staminate flower, days to anthesis of first pistillate flower, days to first fruit harvest. Among the testers, Kalyanpur Sona was found to be a good combiner for node number to anthesis of first staminate flower, node number to anthesis of first pistillate flower in both seasons and for days to anthesis of first pistillate flower, days to fruit harvest indicating earliness, and also more number of fruit and fruit yield per plant in summer season. Pusa Do Mousami (PDM) was also found to be a good combiner for fruit length, fruit diameter and vine length in both the seasons and for fruit yield per plant and number of fruits per plant in rainy season. The parents mentioned above may be used in a future breeding programme for bringing improvement in bitter gourd. Similar results were observed in bitter gourd by Ram et al., 1999; Kumar et al., 2005; Sundharaiya and Venkatesan, 2006 and Sundaram, 2008.

### Specific combining ability effects

Estimation of specific combining ability effects are given in table 3. Among thirty-six crosses, eight in summer and eight in rainy season exhibited significant positive sca effect for increased fruit yield and among these four are common in both seasons. It was also observed that the majority of the crosses which showed significant sca effects for increased yield also involved at least one parent having high and significant gca estimates. The high significant positive sca effects in respect of fruit yield per plant were observed in a cross NDBT-3  $\times$  Pusa Do Mousami followed by NDBT-1  $\times$  Kalyanpur Sona, NDBT-4  $\times$  NDBT-12, NDBT-19  $\times$  Pusa Do Mousami and NDBT-7  $\times$  Kalyanpur Sona in summer season, and NDBT-15  $\times$  NDBT-12 followed by NDBT-10  $\times$  Pusa Do Mousami, NDBT-1  $\times$  Kalyanpur Sona, NDBT-8  $\times$  Pusa Do Mousami, NDBT-7  $\times$  Kalyanpur Sona in rainy season where as NDBT-1  $\times$  Kalyanpur Sona and NDBT-7  $\times$  Kalyanpur Sona NDBT-4  $\times$  NDBT-12 and NDBT-8  $\times$  PDM were found desirable in both seasons. These crosses also showed consistently favourable sca effects for other yield component characters. However, the best crosses varied with the characters. Based on sca effects NDBT-19  $\times$  Pusa Do Mousami in summer season, NDBT-8  $\times$  Pusa Do Mousami, NDBT-15  $\times$  NDBT-12 and NDBT-10  $\times$  Pusa Do Mousami in rainy season were found to be the best specific combiners for fruit yield per plant. The cross NDBT-19  $\times$  Pusa Do Mousami exhibited significantly the highest sca effects for other characters like average fruit weight and number of fruits per plant in summer season. The cross NDBT-3  $\times$  Pusa Do Mousami was also found to be having the highest significant sca effects for fruit diameter and number of fruits per plant in both seasons. The cross NDBT-1  $\times$  Kalyanpur Sona was also found along with the significant sca effect for number of primary branches per plant and number of nodes per plant in summer season. The cross NDBT-4  $\times$  NDBT-12 was also found to be having significant negative sca effects for node number to anthesis of first pistillate flower and vine length in both seasons where as it was positively significant for days to first fruit harvest and fruit length in summer season. The cross NDBT-7  $\times$  Kalyanpur Sona was also found to have significant and negative sca effects for days to anthesis of first pistillate flower, days to anthesis of first staminate flower, days to first fruit harvest indicating earliness for these characters, whereas number of fruits per plant, number of nodes per plant and vine length had significant positive sca effect in both seasons. The cross NDBT-6  $\times$  Pusa Do Mousami had negative sca effects significant for days to anthesis of first pistillate flower and days to first fruit harvest in rainy season. The highest sca effect for days to first harvest was observed in cross combination NDBT-13  $\times$  Kalyanpur Sona in rainy season. For fruit diameter, the highest positive and significant sca effects were observed in the cross combination NDBT-5  $\times$  Kalyan Sona followed by NDBT-3  $\times$  Pusa Do Mousami, NDBT-7  $\times$  NDBT-12, NDBT-13  $\times$  Kalyan Sona, NDBT-8  $\times$  PDM and NDBT-2  $\times$  NDBT-12 in both seasons. For fruit length the highest significant sca effect was observed in a cross combination of NDBT-5  $\times$  Kalyanpur Sona followed by NDBT-7  $\times$  NDBT-12 in summer, while as

NDBT-3 × Pusa Do Mousami followed by NDBT-4 × Kalyanpur Sona in rainy season were the desirable crosses. Similar results were observed in bitter gourd by Ram et al., 1999, Sundharaiya and Venkatesan, 2006, Sundaram, 2008.

From these studies, it was indicated that sca effects of certain crosses were related with gca of their parents as the best cross combination for most of the characters involved at least one parent with high or average gca effects for particular traits in both or any one season (summer or rainy). Hence, these parents may be utilized for development of new hybrids. The crosses NDBT-1 × Kalyanpur Sona and NDBT-7 × Kalyanpur Sona exhibited high sca effects for fruit yield per plant, days to anthesis of first staminate flower, days to anthesis of first pistillate flower, days to first fruit harvest, average fruit weight, number of fruits per plant, number of primary branches per plant, number of nodes per plant and vine length and these crosses can be exploited for development of desirable hybrids for either both (summer and rainy) or summer/rainy season.

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**Table 1: Estimates of genetic components (variances) and heritability narrow sense for 14 traits of bitter gourd during summer and rainy seasons**

S. No.	Characters	gca variance ( $\sigma^2_g$ )		sca variance ( $\sigma^2_s$ )		Av. Degree of dominance ( $\sigma^2_s/2\sigma^2_g$ )		$\sigma^2_A$		$\sigma^2_D$	
		Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy
1.	Node no. to anthesis of first staminate flower	0.02	0.51	1.90	2.75	5.89	1.64	0.05	1.02	1.90	2.76
2.	Node no. to anthesis of first pistillate flower	0.52	0.03	0.67	1.71	0.80	4.88	1.04	0.07	0.67	1.71
3.	Days to anthesis of first staminate flower	0.44	0.91	3.40	4.51	1.95	1.57	0.89	1.81	3.40	4.51
4.	Days to anthesis of first pistillate flower	2.00	1.36	3.45	6.21	0.92	1.51	4.00	2.73	3.45	6.21
5.	Days to first fruit harvest	1.56	1.37	4.36	7.31	1.18	1.63	3.14	2.74	4.36	7.31
6.	Fruit length (cm)	-0.04	0.05	1.46	0.87	4.22	2.79	-0.09	0.11	1.46	0.87
7.	Fruit diameter (cm)	-0.02	-0.01	0.25	0.25	2.99	3.75	-0.02	-0.01	0.25	0.25
8.	Average fruit weight (g)	4.19	5.48	19.71	21.51	1.54	1.41	8.38	10.97	19.71	21.51
9.	Number of fruits per plant	1.48	2.51	5.89	5.23	1.42	1.02	2.95	5.04	5.89	5.23
10.	Fruit yield per plant (kg)	0.01	0.03	0.03	0.03	1.25	0.87	0.02	0.02	0.03	0.03
11.	No. of primary branches per plant	0.36	1.04	5.85	2.82	2.86	1.16	0.72	2.08	5.85	2.82
12.	No. of nodes per plant	3.98	4.18	33.56	21.52	2.05	1.61	7.96	8.37	33.56	21.42
13.	Internodal length(cm)	0.07	0.06	0.43	0.38	1.70	1.74	0.15	0.12	0.43	0.38
14.	Vine length (m)	0.04	0.03	0.11	0.12	1.18	1.37	0.07	0.06	0.11	0.12

**Table 2: Estimate of gca effects of parents for 14 characters during summer and rainy seasons in bitter gourd**

Characters	Node no. to anthesis of first staminate flower		Node no. to anthesis of first pistillate flower		Days to anthesis of first staminate flower		Days to anthesis of first pistillate flower		Days to first fruit harvest		Fruit length		Fruit diameter	
	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy
NDBT-1	-0.56	-1.92**	-2.16**	-1.21**	-0.20	-2.81**	0.83	-3.53**	0.14	-3.68**	-0.04	-0.69**	0.23**	-0.06
NDBT-2	-1.00**	-1.81**	-3.49**	-1.88**	-1.20**	-2.48**	-4.06**	-4.53**	-4.64**	-5.01**	-0.78**	0.25	-0.39**	-0.34**
NDBT-3	-1.11**	-0.69	-0.49	-0.10	-1.54**	-1.26**	-0.61	-2.08**	0.03	-2.68**	0.41*	0.39	0.38**	0.30**
NDBT-4	0.11	0.86*	0.84*	1.12**	-0.43	-0.48	-1.28*	-3.08**	-1.86**	-2.68**	0.42*	-0.59*	0.16**	0.10
NDBT-5	-0.11	0.08	0.06	-0.99*	-1.76**	-1.93**	-1.28*	-0.86	-1.34*	0.21	0.16	-0.59*	-0.09	-0.03
NDBT-6	1.00**	1.75**	0.40	-0.10	0.02	-0.93**	-1.63**	-0.08	-0.86	0.32	-0.27	-0.54*	-0.25**	0.08
NDBT-7	0.08	0.86*	-0.16	-0.10	-1.31**	-1.26**	-0.17	0.81	-0.64	1.10	-0.61**	0.12	0.04	-0.13
NDBT-8	0.67	2.31**	-0.16	1.12*	-0.87*	2.74**	-0.50	-0.97*	-0.42	-0.45	-0.41*	-0.05	-0.56**	-0.41**
NDBT-10	1.00**	0.75	0.95**	1.90**	3.69**	2.41**	2.17**	4.58**	2.81**	4.55**	-0.12	0.70**	0.01	-0.05
NDBT-13	-0.78*	-1.81**	0.51	0.12	-0.76	-1.48**	1.83**	2.69**	2.36**	2.77**	0.86**	1.25**	0.57**	0.63**
NDBT-15	-0.11	-1.25**	1.40**	-0.99*	1.69**	2.30**	1.06	3.03**	1.03	2.21	0.31	-0.69**	-0.05	0.06
NDBT-19	0.89*	0.86*	2.40**	1.12*	2.69**	5.19**	3.61**	4.03**	3.36**	3.32**	0.07	0.44	-0.03	-0.18*
E(gi) Lines	0.35	0.40	0.42	0.45	0.42	0.43	0.61	0.48	0.65	0.56	0.20	0.24	0.05	0.08
SE(gi-gj) Lines	0.50	0.57	0.59	0.63	0.60	0.61	0.86	0.68	0.93	0.79	0.28	0.34	0.08	0.12
Testers														
ζ. Sona	-0.47**	-0.58**	-0.57**	-0.21	-0.34	-0.59**	-1.42**	0.75**	-1.22**	0.96**	-0.42**	-0.44**	-0.02	-0.08*
√DBT-12	0.56**	0.92**	0.29	0.37	-0.43*	-0.06	-0.14	-0.69**	-0.14	-0.87**	0.05	0.18	-0.02	-0.07
ρDM	-0.08	-0.33	0.29	-0.16	0.77**	0.66**	1.56**	-0.06	1.36**	-0.09	0.37*	0.26*	0.04	0.15**
SE(gi) Testers	0.17	0.20	0.21	0.22	0.21	0.21	0.30	0.24	0.32	0.28	0.10	0.12	0.02	0.04
SE(gi-gj) Testers	0.25	0.28	0.29	0.31	0.30	0.30	0.43	0.34	0.46	0.39	0.14	0.17	0.04	0.06
NDBT-1	-10.07**	-12.21**	-1.98**	-2.98**	-0.30**	-0.39**	1.67**	0.77*	-1.39	-1.94*	-1.03**	-0.89**	-0.53**	-0.62**
NDBT-2	2.59**	2.79**	-1.53**	-0.93*	-0.09**	-0.04	-0.68*	2.33**	5.28**	3.72**	-1.18**	-0.83**	-0.40**	-0.23**

Characters	Node no. to anthesis of first staminate flower		Node no. to anthesis of first pistillate flower		Days to anthesis of first staminate flower		Days to anthesis of first pistillate flower		Days to first fruit harvest		Fruit length		Fruit diameter	
	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy
NDBT-3	2.70**	3.79**	-4.05**	-2.17**	-0.32**	-0.14**	-3.67**	-3.20**	2.50**	2.83**	-0.57**	-0.49**	-0.19**	-0.03**
NDBT-4	-2.52**	-3.99**	0.22	0.32	-0.01	-0.04	1.34**	0.88*	2.94**	0.39	-0.53**	-0.50**	-0.20**	-0.18**
NDBT-5	0.70	-0.88	-0.87*	-2.30**	-0.03	-0.16**	1.50**	1.88**	2.06*	1.28	-0.46**	-0.66**	-0.21**	-0.26**
NDBT-6	-3.74**	-3.66**	1.50**	-2.25**	0.02	-0.24**	0.13	0.22	-2.28*	-2.39**	-0.56**	-0.64**	-0.42**	-0.52**
NDBT-7	-2.07**	-1.88	-0.16	-0.88*	-0.05*	-0.11**	0.28	-2.10**	-6.83**	-5.72**	0.38**	0.36**	-0.03	-0.13**
NDBT-8	-3.96**	-3.44**	0.23	1.16**	-0.04	0.02	0.04	-1.54**	-9.17**	-9.06**	0.59**	0.68**	-0.04	-0.05
NDBT-10	0.70	1.68	2.03**	0.86*	0.19**	0.08**	-1.25**	-1.29**	3.72**	3.06**	1.20**	1.11**	0.65**	0.63**
NDBT-13	10.93**	12.79**	1.69**	1.97**	0.36**	0.38**	0.21	-0.49	8.17**	9.94**	0.59**	0.28**	0.74**	0.67**
NDBT-15	9.26**	8.90**	-1.06**	1.77**	0.07**	0.30**	-0.38	-0.22	-11.72**	-8.61**	0.49**	0.47**	-0.28**	-0.23**
NDBT-19	-4.52**	-3.88**	3.97**	5.49**	0.22**	0.34**	0.82*	2.82**	6.72**	6.50**	1.08**	1.12**	0.90**	0.94**
SE(gi)Lines	0.85	0.93	0.39	0.43	0.02	0.03	0.34	0.38	0.89	0.87	0.07	0.05	0.04	0.03
SE(gi-gj) Lines	1.21	1.32	0.55	0.61	0.03	0.04	0.48	0.54	1.27	1.23	0.10	0.07	0.06	0.05
Testers														
K.Sona	-0.07	-0.05	1.52**	-1.25**	0.11**	-0.11**	-1.10**	0.08	-0.69	-0.56	-0.02	-0.05*	-0.03	-0.05**
NDBT-12	0.48	-0.16	-0.59**	-0.59**	-0.03**	-0.04**	0.75**	0.96**	-0.31	-0.42	-0.01	0.01	-0.07**	-0.01
PD M	-0.41	0.20	-0.94**	1.84**	-0.08**	0.16**	0.35*	-1.04**	1.00*	0.97*	0.03	0.04	0.08**	0.06**
SE(gi) Testers	0.42	0.46	0.19	0.21	0.01	0.01	0.17	0.19	0.44	0.43	0.03	0.02	0.02	0.01
SE(gi-gj) Testers	0.60	0.66	0.27	0.30	0.01	0.02	0.24	0.27	0.63	0.61	0.05	0.03	0.03	0.02

\*- Significant at 5 per cent probability level; \*\* - Significant at 1 per cent probability level

**Table 3: Estimate of sca effects of 36 F<sub>1</sub> hybrids for 14 characters during summer and rainy seasons in bitter gourd**

Character	Node no. to anthesis of first staminate flower		Node no. to anthesis of first pistillate flower		Days to anthesis of first staminate flower		Days to anthesis of first pistillate flower		Days to first fruit harvest		Fruit length		Fruit diameter	
	Hybrids	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer
NDBT-1 x K.Sona	-0.63	-0.86	-0.53	-0.56	-0.99	-3.51**	0.63	-1.19	0.55	-0.51	0.61	-0.12	0.24*	-0.00
NDBT-1 x NDBT-12	-0.66	-0.02	1.60**	-0.14	-0.57	0.62	-0.63	-0.08	-0.52	-1.35	0.02	0.29	0.20*	0.23
NDBT-1 x PDM	1.30*	0.88	-1.46*	0.71	1.56*	2.89**	0.00	1.27	-0.02	1.87	-0.69*	-0.16	-0.44**	-0.22
NDBT-2 x K.Sona	0.13	0.02	0.12	1.10	1.00	1.51*	-0.13	1.13	-1.00	-0.55	1.35**	0.11	0.13	0.05
NDBT-2 x NDBT-12	-0.55	-0.80	0.93	-0.48	0.42	-0.04	1.91	0.58	3.25**	1.31	-0.94**	0.70	0.27**	0.39**
NDBT-2 x PDM	0.41	0.77	-1.06	-0.62	-1.49*	-1.43	-1.77	-1.72*	-2.28*	-0.79	-0.45	-0.84*	-0.40**	-0.44**
NDBT-3 x K.Sona	-0.08	-0.41	0.46	0.99	-1.65*	-1.07	-0.58	1.02	-0.66	1.48	-0.02	-0.42	-0.24*	-0.24
NDBT-3 x NDBT-12	-0.77	-0.25	-1.49*	-0.25	0.75	1.39	-1.52	-0.19	-2.08	-0.68	-0.70*	-1.54**	-0.31**	-0.20
NDBT-3 x PDM	0.86	0.67	0.93	-0.73	0.89	-0.32	2.11**	-0.83	2.75*	-0.79	0.72*	1.96**	0.55**	0.44**
NDBT-4 x K.Sona	1.69**	3.02**	2.24**	2.10**	-0.76	0.14	2.08	-0.63	1.55	-0.51	-1.64**	1.48**	0.13	0.10
NDBT-4 x NDBT-12	0.02	0.19	-1.62*	-2.48**	0.98	0.28	-1.86	0.80	-2.29*	0.64	1.31**	-0.89*	-0.21*	0.11
NDBT-4 x PDM	-1.69**	-3.22**	-0.62	0.37	-0.21	-0.43	-0.22	-0.16	0.63	-0.12	0.33	-0.58	0.08	-0.21
NDBT-5 x K.Sona	0.91	0.80	0.24	0.21	-0.10	-0.74	1.08	-0.19	1.33	-0.40	2.24**	-0.04	0.80**	0.98**
NDBT-5 x NDBT-12	-0.11	-1.02	-0.28	-1.37	-1.50*	-0.93	-0.52	0.25	-0.08	0.42	-1.82**	-0.60	-0.74**	-0.76**
NDBT-5 x PDM	-0.80	0.22	0.04	1.15	1.45	1.67*	-0.55	-0.05	-1.25	-0.01	-0.41	0.64	-0.06	-0.21
NDBT-6 x K.Sona	-1.29*	0.80	0.57	2.32**	0.45	1.25	-1.91	2.02*	-1.77	2.58**	-0.51	-0.52	-0.38**	-0.39
NDBT-6 x NDBT-12	0.77	-1.69*	-0.28	-1.59	0.20	-0.26	1.80	0.80	1.80	0.31	1.06**	-0.14	0.43**	0.27
NDBT-6 x PDM	0.41	0.88*	-0.27	-0.73	-0.65	-0.99	0.11	-2.83**	-0.02	-2.79**	-0.55	0.66	-0.04	0.12
NDBT-7 x K.Sona	-0.19	-0.63	-0.53	-0.67	-1.54*	-2.07**	-5.36**	-3.19**	-6.00**	-4.29**	-0.07	-0.10	0.34**	0.22
NDBT-7 x NDBT-12	-0.22	-0.13	0.26	1.07	1.87*	1.06	3.69**	-1.41	3.25**	-1.12	1.73**	0.59	0.51**	0.57**
NDBT-7 x PDM	0.41	0.77	0.26	-0.39	-0.32	1.00	1.66	4.61**	2.75*	5.42**	-1.65**	-0.49	-0.86**	-0.79**
NDBT-8 x K.Sona	-1.86**	-0.08	-0.87	-1.89*	-1.99**	0.92	1.63	-0.75	1.77	0.59	-0.20	1.14**	-0.11	-0.30
NDBT-8 x NDBT-12	2.77**	1.75*	-0.39	0.85	0.75	-1.93*	-2.82**	-0.97	-0.97	0.42	-0.94**	0.26	-0.24*	-0.62**
NDBT-8 x PDM	-0.91	-1.66*	1.26	1.04	1.23	1.00	0.66	1.72*	-0.80	-1.01	1.08**	-1.40**	0.36**	0.92**
NDBT-10 x K.Sona	-0.52	-0.19	0.68	0.99	6.12**	5.92**	1.63	0.69	1.55	0.59	-0.11	-0.37	-0.33**	-0.33
NDBT-10 x NDBT-12	1.77**	1.63*	-0.84	-0.25	-3.12**	-2.60**	0.69	-1.19	-0.19	-1.57	0.54	0.20	-0.03	0.03
NDBT-10 x PDM	-1.25*	-1.44*	0.15	-0.73	-2.99**	-3.32**	-2.83**	0.50	-1.36	0.98	-0.43	0.17	0.37**	0.29
NDBT-13 x K.Sona	1.68**	1.36*	-1.53*	-0.89	0.23	0.48	1.30	6.25**	1.00	6.37**	0.49	0.78	0.48**	0.51**
NDBT-13 x NDBT-12	-1.64**	-0.47	0.93	1.51	0.31	0.95	0.02	-0.97	0.91	-1.12	-1.05**	-0.73	-0.09	-0.10

Character	Node no. to anthesis of first staminate flower		Node no. to anthesis of first pistillate flower		Days to anthesis of first staminate flower		Days to anthesis of first pistillate flower		Days to first fruit harvest		Fruit length		Fruit diameter	
	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy
NDBT-13 x PDM	-0.13	-0.88	0.60	-0.62	-0.54	-1.43	-1.33	-5.27**	-1.91	-5.24**	0.55	-0.05	-0.38**	-0.40**
NDBT-15 x K.Sona	0.25	-0.86	-0.42	-1.78*	-0.21	-1.96*	-0.25	-3.08**	0.33	-4.40**	-0.62	-0.38	-0.54**	-0.12
NDBT-15 x NDBT-12	0.88	1.63*	0.37	1.62*	-0.12	1.50*	-1.19	1.36	-1.75	2.75**	0.69*	1.18**	-0.07	-0.15
NDBT-15 x PDM	-1.13	-0.77	0.04	0.15	0.34	0.45	1.44	1.72*	1.41	1.64	-0.03	-0.70	0.61**	0.28
NDBT-19 x K.Sona	-0.08	-2.97**	-0.42	-1.89*	-0.54	-0.85	-0.13	-2.08*	1.33	-0.85	-1.49**	-1.52**	-0.52**	-0.48**
NDBT-19 x NDBT-12	-2.44**	-0.80	0.71	1.51	-0.12	-0.04	-0.08	1.02	-1.41	-0.01	0.01	0.77	0.31**	0.23
NDBT-19 x PDM	2.52**	3.77**	-0.28	0.37	0.67	0.89	0.22	1.05	0.08	0.87	1.48**	0.76	0.20*	0.25
SE(Sij-Skl)	0.86	0.99	1.03	1.10	1.05	1.06	1.49	1.19	1.61	1.37	0.49	0.59	0.14	0.21
SE(Sij-Sik)	1.80	2.06	2.16	2.29	2.18	2.22	3.12	2.48	3.36	2.86	1.02	1.24	0.30	0.43
NDBT-1 x K.Sona	3.95**	3.60*	3.48**	3.63**	0.27**	0.22**	2.89**	-2.40**	6.13**	0.11	-0.15	0.05	0.17*	0.18**
NDBT-1 x NDBT-12	-2.37	-1.62	-2.10**	-1.95*	-0.15**	-0.16**	-4.42**	-0.69	0.75	1.97	0.04	0.00	0.11	0.06
NDBT-1 x PDM	-1.48	-1.98	-1.38*	-1.68*	-0.11**	-0.06	1.56**	3.10**	-6.88**	-2.08	0.10	-0.06	-0.27**	-0.24**
NDBT-2 x K.Sona	-0.81	-0.73	-1.39*	-0.94	-0.13**	-0.07	-1.46	0.39	-0.52	-4.88**	-0.07	-0.00	-0.00	-0.31**
NDBT-2 x NDBT-12	-2.03	-2.28	1.42*	-0.19	0.08*	-0.06	2.51**	-1.89**	-1.25	-1.36	-0.01	0.05	-0.17*	0.06
NDBT-2 x PDM	2.98*	3.01	-0.02	1.13	0.05	0.16**	-1.04	1.50*	1.77	6.25**	0.08	-0.05	0.18*	0.25**
NDBT-3 x K.Sona	2.07	1.60	-2.03**	-3.20**	-0.15**	-0.21**	-3.20**	-0.47	-6.41**	-3.00	-0.11	-0.18	-0.27**	-0.18**
NDBT-3 x NDBT-12	-0.14	0.37	-3.26**	1.36	-0.25**	0.07	0.67	-0.16	3.52*	2.86	-0.05	-0.09	0.08	0.10
NDBT-3 x PDM	-1.92	-1.98	5.28**	1.83*	0.41**	0.13	2.53**	0.63	2.88	0.13	0.17	0.27**	0.18*	0.08
NDBT-4 x K.Sona	2.96*	3.37*	-0.97	-1.72	-0.00	-0.07	-0.68	1.55*	-3.52*	-3.22	-0.08	-0.19*	-0.18*	-0.15
NDBT-4 x NDBT-12	1.07	1.49	2.56**	1.17	0.17**	0.16**	0.05	-1.34*	3.75*	4.63**	0.17	0.18*	0.33**	0.29**
NDBT-4 x PDM	-4.03**	-4.87**	-1.58*	0.55	-0.16**	-0.09	0.62	-0.23	-0.22	-1.41	-0.09	-0.01	-0.14	-0.14*
NDBT-5 x K.Sona	3.40**	4.30**	-0.65	-0.50	-0.01	0.10	-2.44**	0.65	0.02	-0.11	-0.02	0.15	-0.02	0.03
NDBT-5 x NDBT-12	-0.14	0.04	-0.17	1.66*	0.05	0.16**	0.49	0.17	-0.02	-1.58	-0.16	-0.19*	-0.07	-0.11
NDBT-5 x PDM	-3.25*	-4.31**	0.83	-1.16	-0.03	-0.20**	1.95**	-0.83	0.00	1.69	0.19	0.03	0.09	0.08
NDBT-6 x K.Sona	-5.45**	-3.28*	0.94	0.80	-0.03	0.03	-2.54**	1.58*	2.36	2.88	0.00	0.04	0.11	0.15*
NDBT-6 x NDBT-12	8.29**	5.49**	-2.24**	-1.78*	0.00	-0.06	1.73**	0.37	3.97*	4.41**	0.06	-0.06	0.15	0.12*

Character	Node no. to anthesis of first staminate flower		Node no. to anthesis of first pistillate flower		Days to anthesis of first staminate flower		Days to anthesis of first pistillate flower		Days to first fruit harvest		Fruit length		Fruit diameter	
	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy	Summer	Rainy
NDBT-6 x PDM	-2.81	-2.20	1.30	0.98	0.03	0.02	0.80	-1.96**	-6.33**	-7.30**	-0.07	0.02	-0.27**	-0.27**
NDBT-7 x K.Sona	-2.14	-3.39*	2.26**	2.97**	0.15**	0.19**	2.01**	-0.55	8.23**	7.55**	0.20	0.20*	0.39**	0.41**
NDBT-7 x NDBT-12	4.62**	4.71**	0.109	-1.51*	0.06	-0.03	-0.30	0.61	-2.13	-1.91	0.33**	0.36**	0.17*	0.09
NDBT-7 x PDM	-2.48	-1.31	-2.37**	-1.48	-0.22**	-0.16**	-1.70**	-0.06	-6.11**	-5.63**	-0.53**	-0.57**	-0.57**	-0.51**
NDBT-8 x K.Sona	-0.92	0.15	0.60	0.85	0.03	0.06	0.92	-2.06**	-0.41	1.22	-0.97**	-0.98**	-0.49**	-0.48**
NDBT-8 x NDBT-12	-7.81**	-9.39**	0.05	-1.23	-0.14**	-0.26**	-1.66**	2.13**	1.19	-0.25	-0.74**	-0.65**	-0.29**	-0.31**
NDBT-8 x PDM	8.74**	9.24**	-0.66	0.37	0.11**	0.20**	0.73	-0.07	-0.77	-0.97	1.71**	1.63**	0.78**	0.80**
NDBT-10 x K.Sona	-6.25**	-8.62**	2.10**	1.15	0.06	-0.07	1.48	-1.02	-11.97**	-6.88**	1.51**	1.28**	0.23**	0.27**
NDBT-10 x NDBT-12	5.51**	5.49**	-0.94	-3.76**	0.03	-0.19**	-2.34**	2.21**	5.30**	1.63	-0.58**	-0.29**	-0.14	-0.07
NDBT-10 x PDM	0.74	3.12	-1.16	2.60**	-0.10*	0.27**	0.85	-1.18	6.66**	5.25**	-0.92**	-0.99**	-0.06	-0.20**
NDBT-13 x K.Sona	2.18	2.60	-2.57**	-1.99**	-0.17**	-0.13	-1.34	2.69**	0.58	0.55	0.12	0.01	0.00	-0.03
NDBT-13 x NDBT-12	-4.70**	-5.28**	4.29**	1.66*	0.27**	0.03	2.76**	-1.77**	-2.80	-2.25	0.24*	-0.05	-0.15	-0.14*
NDBT-13 x PDM	2.51	2.68	-1.71*	0.23	-0.09*	0.09	-1.41	-0.93	2.22	1.69	-0.35**	0.03	0.15	0.17**
NDBT-15 x K.Sona	1.85	1.15	-0.43	-1.04	0.07	-0.06	1.15	0.48	4.13**	2.11	-0.60**	-0.60**	-0.03	-0.17**
NDBT-15 x NDBT-12	-1.37	-0.39	1.14	4.40**	-0.01	0.33**	1.45	-1.59*	-11.58**	-8.02**	0.45**	0.41**	-0.19*	-0.25**
NDBT-15 x PDM	-0.48	-0.75	-0.707	-3.36**	-0.06	-0.26**	-2.61**	1.10	7.44**	5.91**	0.15	0.18*	0.23**	0.43**
NDBT-19 x K.Sona	-0.70	-0.73	-1.35*	-0.09	-0.09*	-0.00	3.23**	-0.84	1.36	3.66	0.17	0.17	0.13	0.29**
NDBT-19 x NDBT-12	-0.92	1.37	-0.84	0.14	-0.07	0.07	-0.95	1.92**	-0.69	-0.13	0.26*	0.33**	0.17*	0.18**
NDBT-19 x PDM	1.62	-0.64	2.20**	-0.04	0.16**	-0.07	-2.28**	-1.07	-0.66	-3.52	-0.43**	-0.50**	-0.31**	-0.45**
SE(Sij-Skl)	2.10	2.29	0.95	1.06	0.06	0.08	0.833	0.95	2.20	2.13	0.17	0.12	0.11	0.09
SE(Sij-Sik)	4.37	4.77	1.99	2.21	0.13	0.16	1.73	1.98	4.58	4.45	0.36	0.26	0.24	0.19

\* - Significant at 5 per cent probability level; \*\* - Significant at 1 per cent probability level

## NUTRIENT BUDGETING, ECONOMICS AND ENERGETICS OF COWPEA UNDER POTASSIUM AND PHOSPHORUS MANAGEMENT

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### ABSTRACT

Low availability of phosphorus (P) and potassium (K) in acidic soil is a major constraint for crop production. Cowpea was grown under split-plot design with three levels of K and P (viz., 100%, 75% and 50% of recommended dose). Green pod and stover yield was 20.9 and 16.9% higher with 100% K over 50% K application. Similarly, 100% P had 20.2% higher green pod and 16.6% higher stover yield over 50% P. Production efficiency was found higher with 100% K and P (48.6 and 49.0 kg ha<sup>-1</sup> day<sup>-1</sup> respectively) followed by 75% K. The actual gain of N, P and K, and balance were higher with the increase of K and P levels but reverse in case of gross and net return, and BCR. The energy productivity and energy use efficiency were higher on 100% K and P. Cowpea on acid soil along with 100% recommended dose of 40 kg K and 60 kg P ha<sup>-1</sup> showed highest yield along with soil fertility restoration as well as reduced the cost of inorganic fertilizer, higher BCR (1.89) and energy requirement.

**Key words:** Cowpea, economics, energy, phosphorus, potassium

### INTRODUCTION

Cowpea (*Vigna unguiculata* L.) is an important grain legume in rainfed regions and marginal areas of the tropics and subtropics. It is particularly important in India because this crop can be used as pulse, vegetable and fodder. Cowpea is grown by most farmers due to its short growing cycle. The grain is a good source of human protein, while the haulms are valuable source of livestock protein (Mpeperekki et al., 2000). Apart from these, the cowpea reduces the soil erosion, enriches the soil by atmospheric nitrogen (N)-fixation and reduces the weed growth by smothering effect, and therefore, it is considered as important legume cum vegetable crop. Cowpea can fix more than 50% of its N from N<sub>2</sub>-fixation (Khan et al., 2002).

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Crop productivity on more than 40% of earth's arable land is limited by poor plant availability of P (Vance, 2001). The generally poor food situation in Eastern Himalayan Region (EHR) is related to the low availability of P in the soil. Moderate applications of P fertilizer often have only marginal effect on yields due to P fixation by Iron (Fe) and Al-oxides in the soils (Sample et al., 1980). Most of the soil-P is bound in sparingly soluble P pools and not immediately available to support plant growth. The low availability of soil phosphorus may limit N<sub>2</sub>-fixation and also uptake of K. Pulse crop required higher amount of P to develop their root system. Positive effects of grain legumes on yields of cereals grown in rotation may also be due to enhanced P-availability through secretion of enzymes and acids in the legume rhizosphere (Schlecht et al., 2006). The functions of potassium in plant-water relations, energy and enzyme synthesis and tolerance against diseases are well documented. Application of P and K fertilizer is essential for optimal crop yields when soils are deficient with them. The agronomic efficiency of inorganic P fertilizers has been reported to be only 10–25% within the first year of application, as a large portion of applied fertilizer P is fixed by soil and therefore become unavailable to plant (Thierry, 2008). Cowpea is an important leguminous crop, but the crop is grown under poor soil with or without chemical inputs, therefore, the productivity is very low. An attempt was therefore, required to estimate the nutritional need of the crop to harvest higher yield, nutrient budgeting, economics and energy involvement in production.

## MATERIALS AND METHODS

### Climatic condition and experimental site

A field experiment was carried out at the experimental farm of ICAR Research Complex for NEH Region, Arunachal Pradesh Centre, Basar, located at West Siang District of Arunachal Pradesh, India during 2009 and 2010. Experimental site was at 27°95' N latitude, 94°76' E longitude with 660 m above MSL and falls under humid sub tropical climate. The daily temperature during the study period varied widely between minimum of 20°C and maximum of 35°C, received average rainfall of 1300 mm from March to June. The physical and chemical properties of the soils are silty loam in texture, acidic in reaction (pH 5.3), high in organic carbon (Walkaley and Black, 1.50 g kg<sup>-1</sup>), low in available N (alkaline permanganate N, 205.6 kg ha<sup>-1</sup>), low in available phosphorus (Bray P, 8.3 kg ha<sup>-1</sup>) and medium in available K (Neutral normal ammonium acetate K, 260 kg ha<sup>-1</sup>).

### Treatment details

The experiment was laid out in split-plot design with three replications. Main plots were subjected to potassium with three levels (*viz.* 100, 75 and 50% of recommended dose (RD)) and sub-plots were subjected to phosphorus with three levels (*viz.* 100, 75 and 50% of RD) i.e. 40, 30 and 20 kg K and 60, 45 and 30 kg P ha<sup>-1</sup>. The recommended dose of fertilizers (25:60:40 kg NPK ha<sup>-1</sup>) was applied in the

form of urea, single super phosphate and muriate of potash, respectively. Five tones of well decomposed manure were applied at final land preparation and all the fertilizer as per treatments were applied prior to sowing. Seeds were sown with spacing of 45 cm x 20 cm and all other cultural operations were done as per recommendation.

### Observations on crop and soil

Green pod and stover yield was recorded from the net plot area of (3.0 m<sup>2</sup>). Production efficiency was calculated by dividing green pod yield by 85 days duration of cowpea. Soil samples were collected before and after the experiment and plant samples were collected at the time of final harvest for chemical analysis. The contents of N, P and K in the plant were analyzed by standard procedure and the total uptake of nutrients by plants was calculated from their contents in the plants multiplied by dry matter yield and expressed in kg ha<sup>-1</sup>. N, P and K use efficiency was calculated by nutrient uptake by crop per hectare to the nutrient applied to the individuals.

### Energetic parameters

Energy input and output was calculated using energy equivalents as suggested by Devasanapathy et al. (2009). Energy use efficiency (kg pod MJ<sup>-1</sup> and kg dry matter MJ<sup>-1</sup>) was calculated by dividing green pod yield (kg) by total input energy (MJ) and total dry matter produced (kg) by total input energy (MJ). The details of energy parameters and their estimation are given in table 1.

**Table 1: Energy parameters and their estimation**

Parameters	Formulae	Description
Total energy (MJ ha <sup>-1</sup> )	$T_e = \sum_{i=1}^n (E_g + E_s)$	<i>T<sub>e</sub></i> : total energy, <i>E<sub>g</sub></i> : energy from green pod; <i>E<sub>s</sub></i> : energy from straw
Energy use efficiency green pod (%)	$EE_g = \sum_{i=1}^n \left( \frac{E_g}{T_e} \right)$	<i>EE<sub>g</sub></i> : energy use efficiency of green pod
Energy use efficiency stover (%)	$EE_s = \sum_{i=1}^n \left( \frac{E_s}{T_e} \right)$	<i>EE<sub>s</sub></i> : energy use efficiency of stover
Energy productivity (kg MJ <sup>-1</sup> )	$E_p = \sum_{i=1}^n \left( \frac{G_o}{E_i} \right)$	<i>E<sub>p</sub></i> : Energy productivity, <i>G<sub>o</sub></i> : green pod yield, <i>E<sub>i</sub></i> : energy input
Net energy (MJ ha <sup>-1</sup> )	$N_e = \sum_{i=1}^n (E_o - E_i)$	<i>N<sub>e</sub></i> : Net energy, <i>E<sub>o</sub></i> : energy output

### **Economic and statistical analysis**

Economic analysis was worked out as per prevailing market prices of the inputs and produce. The price (₹ t<sup>-1</sup>) of cowpea pod and stover were ₹ 8000 and 500 respectively [USD 1(\$)= ₹48.0] constant for both the years. The analysis of variance of the data was carried out by using SAS 9.2 software. Treatment mean differences were separated by the least significant difference (LSD) test at 0.05 probability level.

## **RESULTS AND DISCUSSION**

### **Total dry matter and yield**

Total dry matter (TDM) content significantly varied with K and P levels in cowpea (Table 2). TDM was highest with the application of K with 100% of recommendation. There was trend to decrease TDM with the decrease of K levels. Similar trend was followed in both the years. Similarly, 100% P had highest TDM in both the years. This is because of higher growth attributes viz. leaf number and leaf area (data not presented), which helped the plant to produce more photosynthesis and accumulated in different plant parts (Abayomi et al., 2008). Green pod and stover yields were significantly highest with 100% K which showed 20.2 and 21.5% of green pod, and 11.2 and 22.6% of stover yield for 2009 and 2010, respectively higher over 50% K. Though, 100 and 50% P application showed at par in 2010. Phosphorus application with 100% P recorded 19.8 and 20.5% higher green pod yield and 11.5 and 19.6% higher stover yield for 2009 and 2010, respectively over 50% P. Similar results were also observed by Neumann and George, 2009.

### **Production efficiency**

Data presented in table 2 clearly depicts that production efficiency (PE) significant ( $P < 0.05$ ) differed with K and P levels. PE in K application with 100 and 75% showed identical in 2010 but differed in 2009 where 100% P application registered significantly higher. PE was 16.5 and 9.8% higher on 100 and 75% K, respectively over 50% K. Similarly, with P levels, PE was 14.4 and 3.4% higher on 100 and 75% P, respectively over 50% P. In both the years P application with 100% gave significantly higher PE. This might be due to better growth and yield attributes and finally leads to higher yield. The applied nutrients were efficiently taken up by plants. Similar finding was also reported by Singh et al. (2010).

### **Nitrogen uptake and balance**

The N uptake significantly ( $P < 0.05$ ) differed with the different levels of K and P which clearly indicates that the N uptake was highest with the application of 100% K in both the years. However, lowest N uptake was recorded with 50% K (Table 3). Increased N uptake due to higher yield was also confirmed by Tanwar et al. (2010). The nitrogen budgeting was greatly influenced by the levels of K and P applied. It can be visualized that among the K levels, 100% of applied K recorded 71.3% higher actual gain of N followed by 75% (39.3%). However, the balance was recorded 92.6% higher in 100% K followed by 46.3% with 75% of K over the 50% of K. It

clearly indicates that the application of higher levels of K helped the plant to synthesize more N than the lower level of K.

Among the P levels, higher N uptake was recorded with 100% P followed by 75% P than 50% P. Similarly, actual N gain was 110.6% higher with 100% followed by 75% (55.8% higher) over the 50% P. This also clearly depicts that as P levels increased the N fixation significantly increased and more N was left over in the soil. During 2009 the N fixation and actual gain of N was comparatively higher than 2010 which might be due to higher rainfall during 2010 (Figure 1). The high rainfall during the crop period might restrict the plants to fix N in lower quantities or created unfavourable condition might have leached some of the fixed N.

#### **Phosphorus uptake and balance**

Uptake of P varied significantly ( $P < 0.05$ ) with K and P levels. Among the K levels, the higher P was taken up when crop was applied with 100% K followed by 75% K during both the years (Table 4). However, the least P uptake was recorded with 50% K application. This might be due to the applied P was fixed by the Fe- and Al-oxides in the soils (Sample et al., 1980). But still, as K levels increased from 50 to 100% the actual P gain was recorded 25 and 10% higher in 100 and 75% K, respectively. Similarly, as P balance increased from 50 to 100% K levels increased by 14.9 and 6.8% in 100 and 75% P, respectively.

Among the P levels, uptake of P was higher with 100% P followed by 75% P during both the years. P balance increased with 60.7 and 32.9% in 100 and 75% P over 50% P. On the other hand, least P uptake was recorded with 50% of P application. The present study also indicated that plant roots were able to induce transformation and uptake of non-labile soil-P within the soil volume exploited by roots and root hairs. Plants facing a withdrawal of inorganic P can adapt their physiology and development in order to efficiently use the lower supply of P (Thierry, 2008).

#### **Potassium uptake and balance**

Uptake of K varied significantly ( $P < 0.05$ ) with K and P levels. Among the K levels, the higher K was taken up when crop was supplied with 100% K followed by 75% K (Table 5). However, the least uptake was recorded with 50% K application. The K balance is greatly influenced by the applied K and P levels. The actual gain was recorded higher as K levels increased in order of 100% > 75% > 50%. However, K balance was recorded 81.8% higher with 100% K followed by 75% (43.8% higher).

Among the P levels, uptake of K was higher on 100% P followed by 75% P during both the years. As P levels increased the balance of K also increased significantly in order of 100% > 75% > 50%. However, it was recorded that 100% P level increased the K balance with 25.7% followed by 14.7% in 75% P over 50% P application. This might be due to better uptake and efficient utilization of applied nutrients for producing growth and yield attributes and finally leads to higher yield. Similar finding was also reported by Pandey et al. (2006). The positive balance of K

was observed due to release of K from its non exchangeable pool of the soil to meet the demand of crop.

There were significant and positive correlation between N and P uptake ( $R^2 = 0.791$ ) and K uptake ( $R^2 = 0.722$ ) (Figure 2). This clearly indicates that as N uptake increases, it augments the uptake of P and K. Similarly, NUE had the positive correlation with PUE ( $R^2 = 0.43$ ) and followed the exponential relationship, whereas, NUE have the negative linear correlation with KUE ( $R^2 = 0.139$ ) (Figure 3).

### **Energetic analysis**

The energy output was higher for stover as compared to green pod because of higher production of stover. Energy input was higher in sequence 100% > 75% > 50% for K and P levels (Table 6). This is due to more fertilizer input was involved in production. However, energy output was higher for 100% K followed by 75% and least by 50% K. Energy of green pod was highest with higher levels of K and P. This was due to highest green pod yield was obtained by supplying the higher levels of K and P which directly contributed to highest energy output. The Net energy and energy output: input ratio followed the similar trend to energy output. However, the energy output: input ratio ranged from 7.76 to 8.73 in 2009 and 7.08 to 8.78 during 2010 for the K levels and similar trend was noticed for P levels. The EP was 0.217 kg MJ<sup>-1</sup> with 100% K followed by 0.202 kg MJ<sup>-1</sup> in 75% K and least with 0.181 kg MJ<sup>-1</sup> in 50% K. Among the P levels EP were higher for 100% P followed by 75% P (0.219 and 1.96 kg MJ<sup>-1</sup>, respectively) and least with 50% P (0.186 kg MJ<sup>-1</sup>). Similarly, energy use efficiency of green pod and stover were higher with 100% K (146.9 and 829.4, respectively) followed by 75% and least with 50% P. The energy productivity was 19.6% higher in 100% K followed by 11.6% in 75% K over 50% K. Similarly with P levels EP was 17.5 and 5.1% higher for 100% and 75% P over 50% P. On the other hand, energy use efficiency of green pod and stover was recorded 19.4 and 15.4% respectively higher in 100% K followed by 11.1 and 5.9% in 75% K (Table 7). Among the P levels, energy use efficiency of green pod and stover was 17.4 and 12.7% respectively higher with 100% K and 5.2 and 6.8% higher in 75% of P over 50% of P. It is very much clear from the above findings that as the K and P levels increased the energy productivity and energy use efficiency of green pod and stover also increased. Similar findings were also reported by Ozkan et al. (2004) in cowpea and Padhi et al. (2010) in cereals and legumes.

### **Economic analysis**

It is evident from the table 8 that the return and B: C ratio is greatly influenced by the levels of K and P. Gross return, net return and BCR was significantly highest with 100% K application in 2009 but at par to 100 and 75% K application in 2010. Similar trend was followed in case of P application but BCR was identical to all P doses. Among the K levels, the sequence of return and BCR followed 100% > 75% > 50%. Similar trend was recorded for P levels and highest with 100% P followed by 75% and least by 50% P. During both the years the BCR followed the similar trend but all the three P and K levels did not significantly differ (Table 8).

However, the interaction of K and P were statistically similar. It was found that cowpea had direct benefit of N<sub>2</sub>- fixation and also increased the benefit of residual soil fertility. Application of higher level of K and P significantly increased the yield and also helped the budget of N, P and K. Higher level of K (40 kg ha<sup>-1</sup>) and P (60 kg ha<sup>-1</sup>) helped efficient production as well as utilization of energy and economic benefit.

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**Table 2: Dry matter accumulation, pod and stover yield and production efficiency as influenced by potassium and phosphorus**

Treatment	Total dry matter (g plant <sup>-1</sup> )		Green pod yield (t ha <sup>-1</sup> )		Stover yield (t ha <sup>-1</sup> )		Production efficiency (kg ha <sup>-1</sup> day <sup>-1</sup> )	
	2009	2010	2009	2010	2009	2010	2009	2010
<i>Potassium</i>								
100%	26.33 <sup>a</sup>	23.87 <sup>a</sup>	4.23 <sup>a</sup>	4.01 <sup>a</sup>	12.43 <sup>a</sup>	12.82 <sup>a</sup>	49.97 <sup>a</sup>	47.30 <sup>a</sup>
75%	23.33 <sup>b</sup>	22.50 <sup>b</sup>	3.77 <sup>b</sup>	3.86 <sup>a</sup>	11.64 <sup>b</sup>	11.38 <sup>b</sup>	45.31 <sup>b</sup>	46.40 <sup>a</sup>
50%	21.00 <sup>c</sup>	21.44 <sup>c</sup>	3.52 <sup>b</sup>	3.30 <sup>b</sup>	11.16 <sup>b</sup>	10.46 <sup>c</sup>	43.09 <sup>b</sup>	40.38 <sup>b</sup>
LSD (P=0.05)	1.71	0.95	0.34	0.24	0.69	0.87	4.01	2.88
<i>Phosphorus</i>								
100%	27.67 <sup>a</sup>	24.11 <sup>a</sup>	4.23 <sup>a</sup>	4.12 <sup>a</sup>	12.41 <sup>a</sup>	12.53 <sup>a</sup>	49.76 <sup>a</sup>	48.33 <sup>a</sup>
75%	23.00 <sup>b</sup>	22.31 <sup>b</sup>	3.76 <sup>b</sup>	3.64 <sup>b</sup>	11.70 <sup>ab</sup>	11.64 <sup>ab</sup>	45.02 <sup>b</sup>	43.61 <sup>b</sup>
50%	20.00 <sup>c</sup>	21.39 <sup>b</sup>	3.53 <sup>b</sup>	3.42 <sup>b</sup>	11.13 <sup>b</sup>	10.48 <sup>b</sup>	43.59 <sup>b</sup>	42.14 <sup>b</sup>
LSD (P=0.05)	2.72	1.39	0.40	0.40	0.85	1.29	4.25	4.27

K levels: 100% (40 kg); 75% (30 kg) and 50% (20 kg);

P levels: 100% (60 kg); 75% (45 kg) and 50% (30 kg);

Values with the same letter within each variables group are not significantly different ( $P < 0.05$ )

**Table 3: Balance sheet of N (kg ha<sup>-1</sup>) as influenced by potassium and phosphorus management in cowpea**

Treatment	Initial soil N status (a)		N added (b)		N uptake by crop (c)		Soil N status after crop harvest (d)		Nitrogen fixation [(c+d)-(a+b)]		Actual gain/ loss over initial status (a-d)		N balance [(a+b)-(c+d)]	
	2009	2010	2009	2010	2009	2010	2009	2010	2009	2010	2009	2010	2009	2010
<i>Potassium</i>														
100%	205.6	239.1	50.0	50.0	71.53 <sup>a</sup>	68.78 <sup>a</sup>	239.1	268.77	55.03	48.45	-33.5	-29.67	-55.03	-48.45
75%	205.6	231.9	50.0	50.0	64.67 <sup>b</sup>	62.67 <sup>b</sup>	231.9	256.97	40.97	37.65	-26.3	-25.07	-40.97	-37.65
50%	205.6	224.6	50.0	50.0	58.97 <sup>c</sup>	56.89 <sup>c</sup>	224.6	243.47	27.97	25.76	-19.0	-17.87	-27.97	-25.76
LSD (P=0.05)	NS	NS	NS	NS	2.95	5.47	NS	NS	NS	NS	NS	NS	NS	NS
<i>Phosphorus</i>														
100%	205.6	242.7	50.0	50.0	72.89 <sup>a</sup>	70.00 <sup>a</sup>	242.7	274.9	59.99	52.20	-37.1	-32.20	-59.99	-52.20
75%	205.6	232.3	50.0	50.0	64.14 <sup>b</sup>	62.11 <sup>b</sup>	232.3	256.87	40.86	36.68	-26.7	-24.57	-40.86	-36.68
50%	205.6	221.1	50.0	50.0	58.13 <sup>c</sup>	56.22 <sup>b</sup>	221.1	238.51	23.63	21.56	-15.5	-17.41	-23.63	-21.56
LSD (P=0.05)	NS	NS	NS	NS	5.97	6.76	NS	NS	NS	NS	NS	NS	NS	NS

Values with the same letter within each variables group are not significantly different ( $P < 0.05$ )

**Table 4: Balance sheet of P (kg ha<sup>-1</sup>) as influenced by potassium and phosphorus management in cowpea**

Treatment	Initial soil P status (a)		P added (b)		P uptake by crop (c)		Soil P status after crop harvest (d)		Actual gain/ loss over initial status (a-d)		P balance (a+b)-(c+d)	
	2009	2010	2009	2010	2009	2010	2009	2010	2009	2010	2009	2010
<i>Potassium</i>												
100%	8.3	9.8	55.0	55.0	15.79 <sup>a</sup>	14.34 <sup>a</sup>	9.8	10.8	-1.5	-1.0	37.71	39.66
75%	8.3	9.7	55.0	55.0	12.69 <sup>b</sup>	11.87 <sup>b</sup>	9.7	10.5	-1.4	-0.8	40.91	42.33
50%	8.3	9.7	55.0	55.0	10.02 <sup>c</sup>	9.10 <sup>c</sup>	9.7	10.3	-1.4	-0.6	43.58	45.30
LSD (P=0.05)	NS	NS	NS	NS	1.30	1.02	NS	NS				
<i>Phosphorus</i>												
100%	8.3	11.6	70	70	17.09 <sup>a</sup>	16.29 <sup>a</sup>	11.6	13.1	-3.3	-1.5	49.61	52.21
75%	8.3	9.4	55	55	12.66 <sup>b</sup>	11.23 <sup>b</sup>	9.4	10.2	-1.1	-0.8	41.24	42.97
50%	8.3	8.1	40	40	8.76 <sup>c</sup>	7.79 <sup>c</sup>	8.1	8.4	0.2	-0.3	31.44	31.91
LSD (P=0.05)	NS	NS	NS	NS	2.62	2.38	NS	NS				

Values with the same letter within each variables group are not significantly different ( $P < 0.05$ )

**Table 5: Balance sheet of K (kg ha<sup>-1</sup>) as influenced by potassium and phosphorus management in cowpea**

Treatment	Initial soil K status (a)		K added (b)		K uptake by crop (c)		Soil K status after crop harvest (d)		Actual gain/loss over initial status (a-d)		K balance (a+b)-(c+d)	
	2009	2010	2009	2010	2009	2010	2009	2010	2009	2010	2009	2010
<i>Potassium</i>												
100%	260.0	310.8	65	65	86.19 <sup>a</sup>	84.89 <sup>a</sup>	310.8	352.9	-50.8	-42.1	-72.0	-62.0
75%	260.0	293.7	55	55	81.34 <sup>b</sup>	78.89 <sup>a</sup>	293.7	315.8	-33.7	-22.1	-60.0	-46.0
50%	260.0	272.8	45	45	74.16 <sup>c</sup>	71.89 <sup>b</sup>	272.8	276.9	-12.8	-4.1	-42.7	-31.0
LSD (P=0.05)	NS	NS	NS	NS	2.79	6.48	NS	NS				
<i>Phosphorus</i>												
100%	260.0	292.5	55.0	55.0	85.61 <sup>a</sup>	84.56 <sup>a</sup>	292.5	315.4	-32.5	-22.9	-63.1	-52.5
75%	260.0	292.4	55.0	55.0	81.36 <sup>a</sup>	78.89 <sup>ab</sup>	292.4	315.2	-32.4	-22.8	-58.8	-46.7
50%	260.0	292.2	55.0	55.0	74.72 <sup>b</sup>	72.22 <sup>b</sup>	292.2	315.1	-32.2	-22.7	-51.9	-40.1
LSD (P=0.05)	NS	NS	NS	NS	5.59	7.15	NS	NS				

Values with the same letter within each variables group are not significantly different ( $P<0.05$ )

**Table 6: Energetic parameters (MJ ha<sup>-1</sup>) as influenced by potassium and phosphorus management in cowpea**

Treatment	Energy output (pod)		Energy output (stover)		Total energy output		Energy input		Net energy		Output: input	
	2009	2010	2009	2010	2009	2010	2009	2010	2009	2010	2009	2010
<i>Potassium</i>												
100%	28702 <sup>a</sup>	27195.3 <sup>a</sup>	155417 <sup>a</sup>	160278 <sup>a</sup>	184119 <sup>a</sup>	187430 <sup>a</sup>	18896	19146	165223 <sup>a</sup>	168327 <sup>a</sup>	8.73 <sup>a</sup>	8.78 <sup>a</sup>
75%	25538 <sup>b</sup>	26140.7 <sup>a</sup>	145556 <sup>b</sup>	142222 <sup>b</sup>	171094 <sup>b</sup>	168363 <sup>b</sup>	18762	19012	152331 <sup>b</sup>	149351 <sup>b</sup>	8.12 <sup>b</sup>	7.84 <sup>b</sup>
50%	23881 <sup>b</sup>	22374.0 <sup>b</sup>	139583 <sup>b</sup>	130694 <sup>c</sup>	163464 <sup>c</sup>	153068 <sup>c</sup>	18671	18921	144793 <sup>c</sup>	134148 <sup>c</sup>	7.76 <sup>b</sup>	7.08 <sup>c</sup>
LSD (0.05)	2307.7	1641.8	8637.3	10852	7365.4	11616			7365.4	11616	0.387	0.59
<i>Phosphorus</i>												
100%	28702 <sup>a</sup>	27903 <sup>a</sup>	155139 <sup>a</sup>	156667 <sup>a</sup>	183841 <sup>a</sup>	184570 <sup>a</sup>	18998.3	19248.3	164843 <sup>a</sup>	165322 <sup>a</sup>	8.67 <sup>a</sup>	8.58 <sup>a</sup>
75%	25463 <sup>b</sup>	24649 <sup>b</sup>	146250 <sup>ab</sup>	145556 <sup>ab</sup>	171713 <sup>b</sup>	170205 <sup>ab</sup>	18772.4	19022.4	152940 <sup>b</sup>	151182 <sup>ab</sup>	8.14 <sup>ab</sup>	7.92 <sup>ab</sup>
50%	23956 <sup>b</sup>	23157 <sup>b</sup>	139167 <sup>b</sup>	130972 <sup>b</sup>	163123 <sup>b</sup>	154130 <sup>b</sup>	18558.2	18808.2	144564 <sup>b</sup>	135321 <sup>b</sup>	7.80 <sup>b</sup>	7.20 <sup>b</sup>
LSD (0.05)	2687	2719.1	10634	16141	11563	17914			11491	17837	0.57	0.90

Values with the same letter within each variables group are not significantly different ( $P<0.05$ )

**Table 7: Energy productivity and use efficiency as influenced by potassium and phosphorus management in cowpea**

Treatment	Energy productivity		Energy use efficiency (pod)		Energy use efficiency (stover)	
	2009	2010	2009	2010	2009	2010
Potassium						
100%	0.224 <sup>a</sup>	0.209 <sup>a</sup>	151.83 <sup>a</sup>	141.92 <sup>a</sup>	822.31 <sup>a</sup>	836.54 <sup>a</sup>
75%	0.201 <sup>b</sup>	0.203 <sup>a</sup>	136.00 <sup>b</sup>	137.41 <sup>a</sup>	775.51 <sup>b</sup>	747.61 <sup>b</sup>
50%	0.188 <sup>b</sup>	0.174 <sup>b</sup>	127.81 <sup>b</sup>	118.20 <sup>b</sup>	747.29 <sup>b</sup>	690.47 <sup>c</sup>
LSD (0.05)	0.018	0.013	12.20	8.59	45.49	56.94
Phosphorus						
100%	0.223 <sup>a</sup>	0.214 <sup>a</sup>	151.02 <sup>a</sup>	144.88 <sup>a</sup>	816.48 <sup>a</sup>	813.58 <sup>a</sup>
75%	0.200 <sup>b</sup>	0.191 <sup>b</sup>	135.58 <sup>b</sup>	129.57 <sup>b</sup>	778.94 <sup>ab</sup>	764.84 <sup>ab</sup>
50%	0.190 <sup>b</sup>	0.182 <sup>b</sup>	129.03 <sup>b</sup>	123.09 <sup>b</sup>	749.69 <sup>b</sup>	696.20 <sup>b</sup>
LSD (0.05)	0.020	0.020	13.75	13.75	53.85	81.59

Values with the same letter within each variables group are not significantly different ( $P < 0.05$ )

**Table 8: Economic parameters (₹ ha<sup>-1</sup>) as influenced by potassium and phosphorus management in cowpea**

Treatment	Cost of cultivation		Gross return		Net return		B:C	
	2009	2010	2009	2010	2009	2010	2009	2010
Potassium								
100%	13500	13756	40083 <sup>a</sup>	38500.0 <sup>a</sup>	26583 <sup>a</sup>	24743.7 <sup>a</sup>	1.97 <sup>a</sup>	1.79 <sup>a</sup>
75%	13067	13306	35946 <sup>b</sup>	36533.3 <sup>a</sup>	22889 <sup>b</sup>	23227.7 <sup>a</sup>	1.75 <sup>b</sup>	1.74 <sup>a</sup>
50%	12700	12955	33761 <sup>b</sup>	31627.8 <sup>b</sup>	21061 <sup>b</sup>	18672.1 <sup>b</sup>	1.66 <sup>b</sup>	1.44 <sup>b</sup>
LSD (0.05)			2514.3	2150.4	2514.3	2150.4	0.186	0.157
Phosphorus								
100%	14067	14308	40072 <sup>a</sup>	39191 <sup>a</sup>	26006 <sup>a</sup>	24883 <sup>a</sup>	1.85 <sup>a</sup>	1.73 <sup>a</sup>
75%	13067	13305	35894 <sup>b</sup>	34907 <sup>b</sup>	22828 <sup>b</sup>	21602 <sup>ab</sup>	1.78 <sup>a</sup>	1.62 <sup>a</sup>
50%	12133	12404	33833 <sup>b</sup>	32563 <sup>b</sup>	21700 <sup>b</sup>	20159 <sup>b</sup>	1.74 <sup>a</sup>	1.62 <sup>a</sup>
LSD (0.05)			3296.4	3634.8	3026.4	3362.1	0.198	0.215

Values with the same letter within each variables group are not significantly different ( $P < 0.05$ )

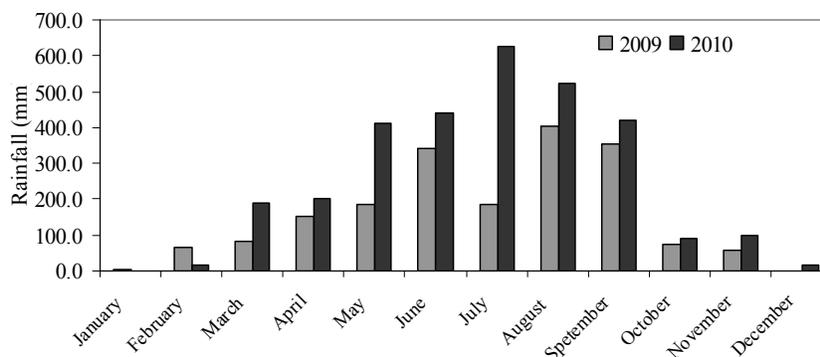


Figure 1: Rainfall distribution of the experimental site during 2009 and 2010

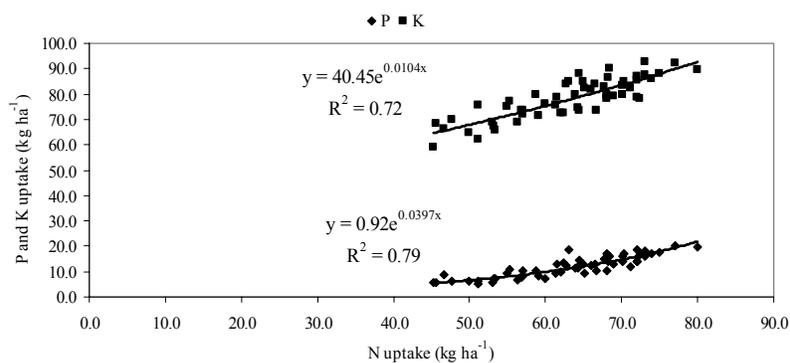


Figure 2: Relationship between nitrogen uptake ( $\text{kg ha}^{-1}$ ) with phosphorus in  $\text{kg ha}^{-1}$  (◆P  $y = 40.45e^{0.0104x}$ ,  $R^2=0.72$ ) and potassium uptake in  $\text{kg ha}^{-1}$  (■K  $y = 0.92e^{0.0397x}$ ,  $R^2= 0.79$ )

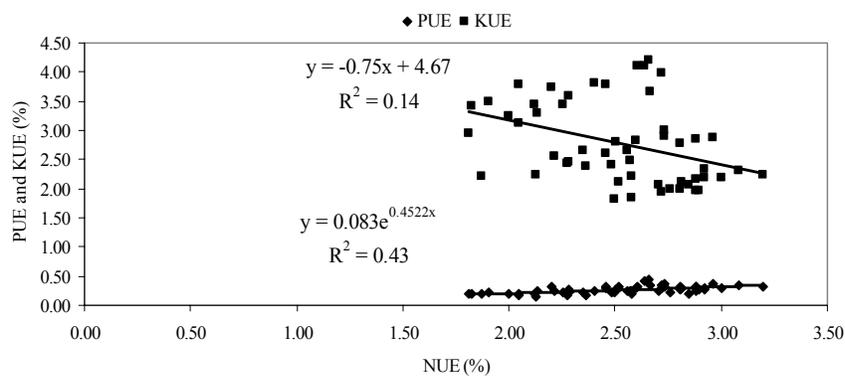


Figure 3: Relationship between nitrogen use efficiency (%) with phosphorus use efficiency (◆PUE  $y = 0.083e^{0.4522x}$ ,  $R^2=0.72$ ) and potassium use efficiency (■KUE  $y = -0.75x + 4.67$ ,  $R^2= 0.14$ )

## EFFECT OF AZOTOBACTER ON GROWTH AND YIELD OF MAIZE

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### ABSTRACT

A field experiment was conducted to study the effect of Azotobacter on growth and yield of maize (variety Rampur composite) at farmland of National Maize Research Program, Rampur, Chitwan, Nepal during the winter season of 2007-08 and 2008-09. The experiment was laid out in randomized complete block design with eight treatments each replicated three times. The treatments were control, 120:60:40 kg  $\text{NP}_2\text{O}_5\text{K}_2\text{O}$   $\text{ha}^{-1}$ , Azotobacter seed inoculation, Azotobacter soil application, Azotobacter +10 t FYM  $\text{ha}^{-1}$ , 10 t FYM  $\text{ha}^{-1}$ , 120:60:40 kg  $\text{NP}_2\text{O}_5\text{K}_2\text{O}$   $\text{ha}^{-1}$  + Azotobacter, 120:60:40 kg  $\text{NP}_2\text{O}_5\text{K}_2\text{O}$   $\text{ha}^{-1}$  + Azotobacter + 10 t FYM  $\text{ha}^{-1}$ . Analysis of variance showed that grain yield, plant height, ear height, ear length, kernel per rows and 1000 grain weight were significantly affected with treatments. Only inoculation of Azotobacter increased 15 to 35% grain yield over non inoculated treatments. The benefit of Azotobacter inoculation was higher in the absence of chemical fertilizer application.

**Key words:** Azotobacter, chemical fertilizer, FYM, grain yield, seed inoculation

### INTRODUCTION

The worldwide spread of inflation, initiated by several fold rises in Petroleum price thereby depicting its striking influence on the prices of chemical nitrogenous fertilizers. The prices of nitrogenous fertilizers have nearly doubled during the last 3-4 years. This has necessitated searching for cheaper source of nitrogen to meet the needs of crops. Farmers use chemical fertilizers to increase production to meet their needs, but the excessive use of fertilizers leads to contamination of soil and groundwater and reduce soil fertility. On the other hand, for marginal farmers in Nepal, the purchase of chemical fertilizers is difficult and expensive. So, biofertilizers can replace partially chemical fertilizers. Hence, there is a need to

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search for alternative strategies to improve soil health without causing damage to environment as well as soil. Therefore, biofertilizers are gaining importance as they are ecofriendly, non hazardous and nontoxic products (Sharma et al., 2007).

Biofertilizers include mainly the nitrogen fixing, phosphate solubilizing and plant growth promoting microorganisms. Biofertilizers benefiting the crop production are *Azotobacter*, *Azospirillum*, *blue green algae*, *Azolla*, *P-solubilizing microorganisms*, *mycorrhizae* and *sinorhizobium* (Selvakumar et al, 2009). Among the biofertilizers, *Azotobacter* represents the main group of heterotrophic, non symbiotic, gram negative, free living nitrogen-fixing bacteria. They are capable of fixing an average 20 kg N/ha/year. The genus *Azotobacter* includes 6 species, with *A. chroococcum* most commonly inhabiting in various soils all over the world (Mahato et al., 2009). Besides nitrogen fixation, *Azotobacter* also produces thiamin, riboflavin, indole acetic acid and gibberellins. When *Azotobacter* is applied to seeds, seed germination is improved to a considerable extent, so also it controls plant diseases due to above substances produced by *Azotobacter*. The exact mode of action by which *Azotobacteria* enhances plant growth is not yet fully understood. Three possible mechanisms have been proposed: N<sub>2</sub> fixation; delivering combined nitrogen to the plant; the production of phytohormone-like substances that alter plant growth and morphology, and bacterial nitrate reduction, which increases nitrogen accumulation in inoculated plants (Mrkovacki and Milic, 2001). In Nepal very limited studies were carried out on the effect of *Azotobacter* on maize growth and productivity. Therefore, a field experiment was carried out to study the effect of *Azotobacter* on maize growth and productivity.

## MATERIALS AND METHODS

The experiment was conducted at farm land of National Maize Research Program, Rampur, Chitwan, Nepal. The experiment was carried out during the winter season of 2007-08 and 2008-09. The treatment consist of T<sub>1</sub>= Control, T<sub>2</sub>= Recommended dose of chemical fertilizer (RDF; 120:60:40 kg NP<sub>2</sub>O<sub>5</sub>K<sub>2</sub>O ha<sup>-1</sup>), T<sub>3</sub>= *Azotobacter* seed inoculation, T<sub>4</sub>= *Azotobacter* soil inoculation, T<sub>5</sub>= *Azotobacter* + 10 t FYM ha<sup>-1</sup>, T<sub>6</sub>= 10 t FYM ha<sup>-1</sup>, T<sub>7</sub>= RDF (120:60:40 kg NP<sub>2</sub>O<sub>5</sub>K<sub>2</sub>O ha<sup>-1</sup>) + *Azotobacter*, T<sub>8</sub>= RDF (120:60:40 kg NP<sub>2</sub>O<sub>5</sub>K<sub>2</sub>O ha<sup>-1</sup>) + *Azotobacter* + 10 t FYM ha<sup>-1</sup>. The experiment was laid out in randomized complete block design (RCBD) with three replications. Maize open pollinated variety named Rampur composite was planted in 12 m<sup>2</sup> plot with the row to row spacing 75 cm and 25 cm plant to plant spacing. The sources of chemical fertilizer were Urea, DAP and MOP. Seed inoculation with the *Azotobacter* was carried out by using 10% sugar solution carrier. The sugar solution was prepared by adding 100 g sugar in 1 litre water and boiled then. After cooling the solution, maize seeds were put in the solution pot and taken out and the inoculants were thoroughly mixed with the maize seeds. These seeds were kept in shade before planting. The inoculation was done just few hours before seed sowing. Soil sampling was done before sowing and analyzed for total N,

available P, available K, Organic matter and pH. The soil was alluvial sandy loam in texture. The initial total N content was low (0.05%), available  $P_2O_5$  was high (173kg  $ha^{-1}$ ), available  $K_2O$  was medium (102 kg  $ha^{-1}$ ), organic matter was low (2.12%) and medium acidic in pH (6.0). All the intercultural operations were carried out as per need. ANOVA was carried out using Genstat 13.2. The significant differences among the means were tested using least significance difference (LSD) at 5% significance level.

## RESULTS AND DISCUSSION

### Growth and harvested ear

Inoculation with Azotobacter significantly influenced the plant and ear height of maize during 2007-08 and 2008-09 (Table 1). Two years mean revealed that maximum plant height (120.5) was recorded with the application of recommended dose of chemical fertilizer plus 10 t FYM  $ha^{-1}$  and Azotobacter inoculation. During 2008-09 the overall plants were taller as compared to 2007-08. Similarly, the ear placement height showed similar trend as like plant height. The numbers of harvested ears per  $m^2$  were significantly affected. The highest number of ears per  $m^2$  (5.41) was recorded in the treatment applied with recommended dose of chemical fertilizer along with Azotobacter inoculation and 10 t FYM  $ha^{-1}$ . It seems that chemical combination of organic and inorganic fertilizer increase the number of ear per  $m^2$ . The numbers of unfilled ears per  $m^2$  was not influenced by the Azotobacter inoculation and other treatments in compared to control during both years.

### Yield attributes

Yield parameters specially ear length, kernels per rows and 1000 grains weight. were highly influenced by Azotobacter inoculation. The mean ear length varied from 10.4 cm to 14.1 cm (Table 2). The maximum ear length (14.1) was seen in the treatment combination consisting of Azotobacter inoculation along with recommended dose of chemical fertilizer and 10 t FYM  $ha^{-1}$ . In contrast, the minimum ear length (10.4) was seen in control. The number of kernel rows per ear was not affected by the treatment during both years. The numbers of kernels per row was significantly influenced by Azotobacter inoculation in 2007-08. However, in 2008-09 kernels per row were not affected by treatments. In 2007-08, maximum number of kernels per rows was obtained in the treatment consisting of Azotobacter inoculation along with recommended dose of chemical fertilizer and 10 t FYM  $ha^{-1}$ . From this result, it was confirmed that the effect of Azotobacter, chemical fertilizer and FYM were positive and additive on numbers of kernel setting per ear. Thousand grains weight was significantly influenced by the treatment in both the years. Two years mean 1000 grain weight was varied from 432 to 490.8 g. The highest 1000 grains weight was measured in the treatment inoculated with Azotobacter along with recommended dose of chemical fertilizer and 10 t FYM  $ha^{-1}$  and lowest 1000 grains weight. was measured in the treatment of 10 t FYM  $ha^{-1}$  applied. Kader et al, (2002)

reported that *Azotobacter* increases N availability in the soil which could enhance the numbers of grains and 1000 grains weight. Application of recommended dose of chemical fertilizer showed major influencing factor in 1000 grain weight in 2007-08. The variation of influence of *Azotobacter* on yield attributes such as 1000 grains weight in 2008-09 than previous year may be the *Azotobacter sp.* populations affected by soil chemical (e.g. organic matter, pH, temperature, soil depth, soil moisture) and microbiological (e.g. microbial interactions) properties (Ridvan, 2009).

### **Grain and stover yield**

Grain and stover yield were significantly influenced by the *Azotobacter* inoculation during both years. The grain yield varied from 2.83 t ha<sup>-1</sup> to 6.62 t ha<sup>-1</sup> (Table 3). The highest grain yield (6.42 t ha<sup>-1</sup>) was recorded with the *Azotobacter* inoculation along with recommended dose of chemical fertilizer and 10 t FYM ha<sup>-1</sup> treatment. In contrast the lowest grain yield (2.83 t ha<sup>-1</sup>) was recorded in control. Two years mean grain yield revealed that only inoculation of *Azotobacter* increased grain yield 0.3 to 35% more as compared to non inoculated treatment combination. Similarly, the highest stover yield (11.9 t ha<sup>-1</sup>) and lowest stover yield (3.83 t ha<sup>-1</sup>) were seen with the same treatments which produced highest grain yield and lowest grain yield. Peng et al. (2013) also reported the positive effect of manure and *Azotobacter* application on maize biomass. The stover and grain yield were highly correlated in the experiment (not shown). The highest benefit of 35% grain yield increment was obtained with the seed inoculation treatment over the control followed by 34% in soil inoculated treatment over the control treatment. Inoculation of *Azotobacter* in addition to only 10 t FYM ha<sup>-1</sup> increased 15% more grain yield over the treatment only applied 10 t FYM ha<sup>-1</sup>. The lowest benefit of 0.3 % increase in yield by *Azotobacter* inoculation was obtained in the treatment applied with recommended dose of chemical fertilizer and *Azotobacter* inoculation over the recommended chemical fertilizer applied treatment. This result was in line with the Peng et al. (2013), they concluded that *Azotobacter chroococcum* inoculation with maize seeds not only economically most efficient at lower doses of N but also saved N when applied in combination with FYM. Biari et al. (2008) also found positive effect of *Azotobacter* application on maize grain yield increase at organic field condition. The benefit of *Azotobacter* applying with FYM could be as *Azotobacter* uses carbon for its metabolism from simple or compound substances of carbonaceous in nature. Besides carbon, *Azotobacter* also requires calcium for nitrogen fixation. Similarly, a medium used for growth of *Azotobacter* is required to have presence of organic nitrogen, micro-nutrients and salt in order to enhance the nitrogen fixing ability of *Azotobacter*.

### CONCLUSION

Inoculation with *Azotobacter* significantly increased plant height, ear height, number of ears per m<sup>2</sup>, ear length, kernel per row, 1000 grain weight, grain and stover yield of maize. Only inoculation of *Azotobacter* increased maize grain yield upto 35% over non inoculated treatment. The benefit of *Azotobacter* inoculation was higher when chemical fertilizer was not used. A positive additive (15% yield increased) effect of 10 t FYM ha<sup>-1</sup> with *Azotobacter* inoculation was seen. Therefore, it was concluded that *Azotobacter* could be one of the biofertilizer option for sustainable and environmental ecofriendly for maize production where chemical fertilizer is limited.

### ACKNOWLEDGEMENT

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**Table 1: Effects of Azotobacter on growth and yield attributes of maize during 2007-08 and 2008-09 at Rampur**

Treatments	Plant height (cm)			Ear height (cm)			Harvested ears/m <sup>2</sup>			Unfilled ear/m <sup>2</sup>		
	2007-08	2008-09	Mean	2007-08	2008-09	Mean	2007-08	2008-09	Mean	2007-08	2008-09	Mean
T1	150	184	167.0	65	92	78.5	4.36	4.21	4.28	1.03	1.16	1.09
T2	194	213	203.5	85	112	98.5	5.56	5.05	5.30	0.20	0.50	0.35
T3	177	216	196.5	78	115	96.5	4.73	4.43	4.58	0.73	0.88	0.80
T4	184	209	196.5	68	108	88.0	4.13	4.66	4.39	1.35	0.83	1.09
T5	188	211	199.5	90	113	101.5	4.33	4.00	4.16	0.96	1.38	1.17
T6	186	214	200.0	96	111	103.5	5.08	4.60	4.84	0.33	0.93	0.63
T7	206	211	208.5	94	111	102.5	4.86	4.83	4.84	0.41	0.76	0.58
T8	205	216	210.5	100	113	106.5	5.33	5.50	5.41	0.21	0.16	0.18
F-test	**	**		*	**		*	*		NS	NS	
CV%	5.2	2.92		9.5	4.95		14.2	10.3		24.7	28.80	
LSD(0.05)	13.6	10.7		14.3	9.51		1.00	0.84		-	-	

NS, p >0.05; \*, p<0.05; \*\*, p>0.01

**Table 2: Effects of Azotobacter on yield attributes of maize during 2007-08 and 2008-09 at Rampur**

Treatments	Ear length (cm)			Kernel rows/ear			Kernels/row			1000 grain wt.(g)		
	2007-08	2008-09	Mean	2007-08	2008-09	Mean	2007-08	2008-09	Mean	2007-08	2008-09	Mean
T1	9.5	11.3	10.4	12.2	13.2	12.7	22.4	26.8	24.6	404.8	353.4	379.1
T2	12.4	13.8	13.1	13.4	13.4	13.4	29.8	28.6	29.2	480.6	437.6	459.1
T3	12.6	13.0	12.8	13.6	13.7	13.7	25.2	28.4	26.8	456.2	437.8	447.0
T4	11.2	12.4	11.8	11.8	13.7	12.7	26.2	26.8	26.5	444.6	434.4	439.5
T5	13.3	12.4	12.8	14.6	14.0	14.3	29.3	27.5	28.4	479.8	445.3	462.5
T6	11.2	12.9	12.0	13.6	13.0	13.3	24.5	27.4	25.9	457.2	406.8	432.0
T7	12.4	14.2	13.3	13.0	14.0	13.5	29.4	30.7	30.0	474.4	458.2	466.3
T8	14.2	14.0	14.1	12.8	14.4	13.6	30.8	31.1	30.9	488.6	493.0	490.8
F-test	*	**		NS	NS		*	NS		*	*	
CV%	5.2	4.98		11.2	5.08		4.4	6.17		13.8	6.72	
LSD(0.05)	1.20	1.13		-	-		6.4	-		46.8	51.01	

NS, p >0.05; \*, p<0.05; \*\*, p>0.01

**Table 3: Effects of Azotobacter on stover and grain yield of maize during 2007-08 and 2008-09 at Rampur**

Treatments	Stover yield (t ha <sup>-1</sup> )			Grain Yield (t ha <sup>-1</sup> )			Grain yield increase (%) by inoculation over non-inoculation
	2007-08	2008-09	Mean	2007-08	2008-09	Mean	
T1	2.80	4.86	3.83	2.75	2.91	2.83	-
T2	9.28	10.05	9.67	5.40	6.30	5.85	-
T3	5.45	7.32	6.39	3.27	4.39	3.83	35
T4	5.80	7.48	6.64	3.12	4.49	3.81	34
T5	7.90	6.03	6.97	4.90	4.62	4.76	15
T6	6.10	6.77	6.44	4.22	4.06	4.14	-
T7	9.46	10.63	10.05	5.36	6.38	5.87	0.3
T8	12.8	11.04	11.92	6.22	6.62	6.42	
F-test	**	**		**	**		
CV%	13.1	15.46		12.8	15.47		
LSD (0.05)	1.99	2.18		1.23	1.31		

NS, p &gt;0.05; \*, p &lt;0.05; \*\*, p &gt;0.01

## **HYDRO-PRIMING OF SEED IMPROVES THE WATER USE EFFICIENCY, GRAIN YIELD AND NET ECONOMIC RETURN OF WHEAT UNDER DIFFERENT MOISTURE REGIMES**

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### **ABSTRACT**

Wheat is the second most important staple food crop of India and contributes a major share to food basket of the country. Since majority of the area under wheat is irrigated, it consumes huge quantity of fresh water for its cultivation. The availability of good quality water for irrigation is decreasing over a period of time due to vagaries of monsoon, urbanization and industrialization. The biggest challenge on this front is to improve the efficiency and productivity of water being used in existing cropping system. Therefore, it is the need of hour to improve water use efficiency for wheat production. In the milieu, the present investigation was taken with an objective of studying the effect of pre-germinated seed in crop establishment under sub-optimal soil moisture conditions by using the residual soil moisture after harvesting of rice in Indo-Gangetic plains, so that pre sowing irrigation requirement for crop establishment may be cut and reduced in time period which require from pre-sowing irrigation to field preparation. This experiment was conducted for two consecutive years 2010-11 and 2011-12 to evaluate the influence of hydropriming on the water use efficiency and grain yield of wheat (*Triticum aestivum* L.) under moisture stress. The experiment was conducted in split plot design with three replications keeping moisture stress treatments (optimum moisture, sub-optimal moisture and dry soil followed by irrigation) in main plots and seed priming treatments (dry seed, hydropriming, and pre-germinated seeds) in subplots. Pre-germinated seed produced significantly higher grain yield ( $5.49 \text{ t ha}^{-1}$ ), which was statistically similar to hydropriming ( $5.30 \text{ t ha}^{-1}$ ). Various seeding methods were statistically at par. The hydro-primed and pre-germinated seeds established earlier than dry seeds leading to better crop establishment under optimum, sub optimum soil moisture as well as dry soil conditions leading to higher tillering and grain yield. The results

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of experiment showed that priming with plain water and pre-germinated seeds improved germination indices, seedling growth and crop establishment. Since priming with plain water and to have pre-germinated seeds is simple and cheap method, which can increase germination percentage and homogeneity of seedling emergence under water stress conditions and it can be easily used by farmers. Interactive effect of different seed priming techniques along with seeding at sub optimal soil moisture level proved to be an efficient technique for enhancing water productivity of wheat crop.

**Keywords:** Economics, seed priming, yield, water productivity and wheat

## INTRODUCTION

Wheat (*Triticum aestivum* L.), is the most important cereal crop and is also staple food for millions of people around the world. Irrigation water decreasing and drought are the major factor limiting crop growth and productivity in many regions of the world, the loss of which is more than any other single environmental factor (Farooq et al., 2009a). Wheat yields in the South Asia where rice-wheat cropping system is following, are suffering due to delayed sowing owing to late harvesting of rice, short period of winter seasons, less developed facilities of irrigation and poor crop stands due to lack of optimal moisture. The availability of good quality water is decreasing with time due to higher population pressure, urbanization and industrialization. However, changing global climate is making the situation more serious (IPCC 2007). Water deficit during initial stage of crop results in delayed and erratic seedling emergence and stand establishment and in severe cases, complete inhibition of seedling emergence may also result (Kaya et al., 2006). The biggest challenge on this front is to improve the efficiency and productivity of water being used in existing systems. Harris (1996) demonstrated that simply soaking seeds in plain water before sowing could increase the speed and homogeneity of germination and emergence, leading to better crop stands, and stimulated seedlings to grow much more vigorously. Hydro priming, a simple hydration technique to a point of pre-germination metabolisms without actual germination (Farooq et al., 2009b), is one of the most pragmatic, simple, economic and short-term approaches to combat the effects of drought (Kaya et al., 2006) and other abiotic stresses (Jafar et al., 2012) on seedling emergence and crop establishment. Hydro primed seeds usually have early, higher and synchronized germination owing to reduction in the lag time of imbibitions otherwise required much time (Brocklehurst and Dearman 2008) and build-up of germination enhancing metabolites (Farooq et al., 2006). Preliminary research has also identified a number of opportunities for priming to be used as a vehicle to introduce biofertilizers, micronutrients and crop protection agents into seeds. Studies also suggested that it is possible to prime seeds with small amount of phosphate to good effect in the early root growth is stimulated allowing more effective uptake of available P in the soil (Johanson et al., 2004). Rajpar et al., (2006)

reported that hydro-priming improved wheat yield under non-saline conditions. Good establishment increases competitiveness against weeds, increases tolerance to abiotic stress especially dry spells and ultimately maximizes the yields (Clark et al., 2001). Direct benefits due to seed priming includes, faster emergence, better and more uniform stands, more vigorous plants, better drought tolerance, earlier flowering and higher grain yield in many crops (Harris et al., 1999; Harris and Hollington, 2001).

Wheat is the second most important staple food crop of India and contributes major share to food basket (93.9 MT in 2011-12) of the country (Sharma et al., 2013). The present water productivity of wheat is 1500 litres of water/kg of wheat grain (FAO, 2012). Since 90 per cent of wheat crop is grown in irrigated condition, thus it consumes huge quantity of fresh water for its production (Agricultural Statistics at a glance 2010). It is, therefore, the need of hour to improve water use efficiency for wheat production. Keeping this fact in view, the present investigation was undertaken with an objective to study the effect of seed priming and pre germinated seeds in crop establishment under sub optimal soil moisture conditions in Indo-Gangetic plains, so that pre-sowing irrigation requirement of wheat crop and time period (from pre sowing irrigation to field preparation) may be reduced.

### MATERIALS AND METHODS

Wheat genotype DBW 17 was used for this study and seed was obtained from Seed Project, Directorate of Wheat Research, Karnal, India. The present study was conducted at the research farm of Directorate of Wheat Research, Karnal, Haryana, India. The region is characterized by sub tropical and semi arid climate with a hot dry summer (March-June), wet monsoon season (late June – mid September) and a cool dry winter (November-February). Average annual rainfall of Karnal location is 744 mm of which about 80 percent is received during the monsoon. The climate is sub-tropical with mean maximum temperature ranging in between 34-39<sup>0</sup>C in summer and mean minimum temperature ranging in between 6-7<sup>0</sup>C in winter. The crop season received 129.7 mm and 36.3 mm rainfall during 2010-11 and 2011-12, respectively. The soil of experimental field was sandy clay loam with pH 7.3 (1:2.5 soils to water). The soil had organic carbon 0.4 %, available N 190 kg ha<sup>-1</sup>, available P 17.8 kg ha<sup>-1</sup> and available K 165 kg ha<sup>-1</sup> at the beginning of the experiment. A field experiment comprising three main plot treatments viz. 1- seeding at optimum moisture level (17.5 %), 2- seeding at sub optimal moisture (10.9 %) and 3- seeding in dry soil followed by irrigation (4.2 %); and three sub plot treatments viz., 1- no seed priming, 2- seed priming with plain water and 3- pre germinated seed was conducted in split plot design with three replications during Rabi (winter) seasons of 2010-11 and 2011-12. To impose the moisture stress treatments pre sowing irrigations were given in treatment optimum moisture level, sub optimal moisture level was obtained by using the residual soil moisture after rice harvesting (no pre sowing irrigation), whereas treatment dry soil followed by irrigation was achieved three times plowing the field to air dry the field (without pre sowing irrigation). Seed priming treatment, seeds

were soaked in normal water for about 8 hours before sowing and for pre-germinated seeds treatment; seeds were soaked in normal water over night, spread on a gunny bag and covered by another wet gunny bag for next 12 hours. The sowing of seeds were done on 18 and 6 November in 2010 and 2011, respectively and harvested on 28 April & 1 May in 2011 and 2012, respectively. The row spacing was kept 20 cm. Recommended doses of fertilizer (150:60:40 kg of N, P<sub>2</sub>O<sub>5</sub> and K<sub>2</sub>O ha<sup>-1</sup>) were applied as part of nutrient management. Full doses of P and K and 1/3 dose of N through NPK mixture (12:32:16) and urea were applied at sowing and remaining N was applied equally in two splits at first (25 days after sowing) and second (45 days after sowing) irrigations. Soon after seed sowing one light irrigation was imposed in treatment dry sowing followed by irrigation. Subsequent irrigations were given at critical growth stages of wheat crop. For weed control one spray of Sulfosulfuron (Leader 75 WG) @ 25 g ha<sup>-1</sup> were done 10 days after first irrigation. Subsequently one hand weeding was done to keep the experimental area free from weeds. Other management practices were adopted as per recommendations of the crop under irrigated conditions. Germination count per meter square was recorded at 4<sup>th</sup> day after sowing. Number of productive tillers per square meter from the centre of plot was measured in each plot at maturity. Plant height data were recorded by measuring the height of ten representative plants from each plot randomly. A random sample of ten spikes was taken from each plot to determine spike length at maturity. A net plot of size 6 m<sup>2</sup> was harvested manually to obtain biomass and yield data. Thousand grains were randomly selected from each sub plot and weighed.

Water productivity and economic analysis combines physical accounting of water with yield or economic output to assess how much value is being obtained from the use of water (Abdullaev et al., 2007). For this analysis, physical water productivity was calculated by the following equation:

$$WP = \text{Output}/Q$$

Where,

WP is the productivity of water in kg m<sup>-3</sup>, output is the productivity of wheat in kg ha<sup>-1</sup> and Q is water used by the crop in m<sup>-3</sup> ha<sup>-1</sup>.

Total amount of water used by wheat was calculated by multiplying the discharge from the tube well with total time taken for irrigating the crop throughout the season (total irrigating hours multiplied by volume of water drawn out per hour). Based on this, the average discharge of tubewell with 10 HP was estimated at 600m<sup>-3</sup> ha<sup>-1</sup> per irrigation (Kaur et al., 2012).

For statistical analysis SAS version 10.3 had been used to analyze the observations and differences among means were further grouped into significant classes by Duncan's New Multiple Range Test at five percent probability.

## RESULTS AND DISCUSSION

### Germination Indices

Pre-germinated seeds and seed priming with normal water gave beneficial effect for seed germination, seedling emergence and subsequently crop establishment. Pre-germinated seeds and seed priming in this study resulted significantly higher germination 153 & 167 per square meter which were significantly higher over dry seeds (136). In moisture regime treatments maximum germination was recorded for the treatment dry sowing followed by irrigation (205 per square meter) which was significantly superior to optimum and sub optimal soil moisture treatments. It seems that primed seeds decreased the time of emergence by about 50 per cent. Priming of seeds with normal water makes them rapidly imbibe water and revive metabolism and germination. This then results in higher germination rate, improved stand establishment, increased stress tolerance and ultimately higher yield. The importance of early germination, seedling emergence and rapid stand establishment and growth is quite essential to compete for water, light and nutrients.

### Yield Attributes

Yield attributes viz. number of effective /m<sup>2</sup> (500), ear head length (10.8) and 1000 grain weight (39.08 g) were the highest in pre-germinated seeds followed by hydro primed seeds and found significantly higher than unprimed seeds. The highest yield attributes in pre-germinated seeds may be ascribed to higher number of effective tillers which lead to higher dry matter accumulation (biomass), translocation and conversion of photosynthesis in to reproductive parts. Seeds can be soaked before sowing to meet the initial seed imbibitions requirement. These results indicate that seed priming and pre-germinated seeds caused early and vigorous germination, which led into proper crop establishment under sub optimal soil moisture conditions, there by, producing higher yield attributes. Similar beneficial effects of seed priming were reported by Rajpar et al. (2006) and revealed that seedlings were significantly faster in emergence, took fewer days to mature and gave significantly higher grain yield. Among seeding methods (seeding at optimum moisture level, sub optimal moisture and in dry soil followed by irrigation) except number of effective tillers, all yield attributing characters were found non-significant. Highest number of effective tillers /m<sup>2</sup> (499) were recorded for the treatment seeding in dry soil followed by irrigation which was statistically superior over seeding at optimum moisture level (491) and seeding at sub optimal moisture (457).

### Yield

The grain (5.49 t ha<sup>-1</sup>) and biomass (12.7 t ha<sup>-1</sup>) yields of wheat varied significantly due to pre-germinated seeds. These were the highest under pre-germinated seeds sown with optimal soil moisture and sub optimal soil moisture conditions and were significantly higher than unprimed seeds. Pre-germinated seeds

and hydro primed seeds registered 9.2 and 5.2 % higher grain yield compared to unprimed seeds, respectively. Increase in grain yield of wheat under pre-germinated seeds and seed priming treatments could be attributed to higher yield attributes whereas, the increase in biological yield was due to higher plant density and plant height. Higher grain and biomass yield in pre-germinated seeds could also be attributed to early germination and vigorous growth, consequently good crop establishment. Higher grain yield with seed priming of wheat has been also reported by Harris et al. (2001b) and Rashid et al. (2002). Harris et al. (1999) reported that early emergence and maturity in seed priming treatment could be due to advancement in metabolic state. Musa et al. (1999) also concluded that priming improve plant stand and provide benefits in term of maturity. This view was further supported by Rashid et al. (2002).

### **Water productivity**

The highest yield was observed in  $M_1S_3$  ( $6.39 \text{ t ha}^{-1}$ ) during 2010-11 and  $M_3S_3$  ( $4.95 \text{ t ha}^{-1}$ ) during 2011-12. The information in table 4 indicates the total irrigation water (tube well water + rainfall) applied for crop under different treatments. During both the years of study, water productivity per hectare was highest in the case of pre-germinated seeds technique coupled with seeding at sub optimal soil moisture level ( $M_2S_3$ ). Productivity was higher in 2011-12 in spite of low rainfall in comparison to 2010-11 and it could be due to early sowing as compared to the year 2010-11.

Cost of irrigating per hectare of wheat was more in 2011-12 owing to marginal increase in the wages of labourers (Table 5). The quantum of water used for irrigation was low in the case of seeding at sub optimal soil moisture level under seed priming techniques. The cost of irrigation ranged from as high as INR 2232  $\text{ha}^{-1}$  to as low as INR 1860/ha due to sacrificing irrigation at the rate of  $600 \text{ m ha}^{-3}$  for seeding at sub optimal soil moisture level under different seed priming techniques.

On the basis of these two years study, it can be revealed that pre-germinated and hydro primed seed sowing may be adopted to enhance seed germination, emergence, and higher biomass as well as grain yield of wheat in Indo-Gangetic plains of India. In addition this technology helps advancing the wheat sowing by about 10-15 days for avoiding the pre sowing irrigation time period require from pre sowing irrigation to field preparation. 10-15 days time period is very crucial in rice-wheat cropping system, the largest cropping system of South Asia. In this region wheat planting usually gets delayed in Rice-Wheat system due to late harvesting of rice which can cause wheat yield penalty. Secondly poor establishment of crop due to lack of optimum soil moisture could be a major constraint for proper crop establishment in other crop sequences. Integrating pre-germinated seeds, seed priming with surface seeding can be a cost effective strategies for enhancing productivity in these rice fallows. On the other hand, good establishment increases competitiveness against weeds, increases tolerance to dry spells maximizes yields and save the time and cost of one irrigation which can also advance the sowing of

wheat after rice crop. Clark et al., 2001 also expressed their views in the same manner. The benefits accrued for the country due to saving irrigation water was estimated in view of the overall deteriorating water situation. The average productivity of wheat was 4.56 t ha<sup>-1</sup> in 2010-11 and 5.85 t ha<sup>-1</sup> in 2011-12 (Table 4). Production was estimated at 45.58 MT in 2010-11 and 58.47 MT in 2011-12 and the value of production stood at INR 533247 million and INR 751340 million respectively for 2010-11 and 2011-12. Following the water conservation technique, about 20 percent of irrigation water @ 600 m ha<sup>-3</sup> and 20 percent of energy @ 24 units ha<sup>-1</sup> could be saved for the country as a whole under rice-wheat system.

### CONCLUSION

Pre-germinated seeds and seed priming is helpful in reducing the risk of poor stand establishment under sub optimal soil moisture conditions. Pre-germinated seeds and hydro priming of seeds are the simple, cost effective and useful technique, which can advance the wheat sowing by about 10 days and beneficial for both crop thereby increasing wheat yield.

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**Table 1: Effect of seed priming and seeding method on yield attributing characters of wheat under sub-optimal soil moisture conditions**

Treatments	Germination/m <sup>2</sup>			Productive tillers/m <sup>2</sup>			Spike length (cm)			Plant height (cm)			1000 grain weight (g)		
	2010-11	2011-12	Pooled	2010-11	2011-12	Pooled	2010-11	2011-12	Pooled	2010-11	2011-12	Pooled	2010-11	2011-12	Pooled
<i>A. Seeding Method</i>															
M1*	140	142	141	444	539	491	10.2	10.3	10.2	83.4	90.8	87.8	36.7	41.0	38.9
M2	108	112	110	392	521	456	10.1	10.2	10.2	85.4	91.3	88.4	36.5	40.3	38.4
M3	204	205	205	440	568	504	9.5	9.5	9.5	84.2	92.1	88.0	37.2	40.3	38.8
SEm±	6.0	5.0	5.4	4.6	12.1	7.3	0.23	0.22	0.224	0.72	0.83	0.92	0.36	0.24	0.18
CD (P=0.05)	23.4	19.7	21.1	18.2	47.9	28.5	0.90	0.8	NS	2.8	2.3	NS	1.4	0.9	NS
<i>B. Seed Priming</i>															
S1*	135	136	136	420	509	464	9.6	9.5	9.6	84.1	90.2	87.8	36.6	40.1	38.3
S2	151	154	153	424	540	482	10.1	10.2	10.1	84.8	91.3	88.3	36.7	40.5	38.7
S3	165	169	167	432	579	500	10.3	10.2	10.2	83.9	92.2	87.9	37.1	41.1	39.1
SEm±	5.2	7.7	6.4	4.2	10.5	6.1	0.16	0.17	0.175	0.48	0.62	0.17	0.16	0.25	0.13
CD (P=0.05)	16.2	23.8	19.7	12.9	32.3	19.1	0.51	0.55	NS	1.4	1.4	NS	0.5	0.7	0.4

**Table 2: Effect of seed priming and seeding method on yield of wheat under sub-optimal soil moisture conditions**

Treatments	Grain Yield (t ha <sup>-1</sup> )			Biomass Yield (t ha <sup>-1</sup> )		
	2010-11	2011-12	Pooled	2010-11	2011-12	Pooled
<i>A. Seeding Method</i>						
M1*	4.67	5.89	5.28	10.63	14.29	12.49
M2	4.56	5.85	5.20	10.54	14.21	12.45
M3	4.66	6.01	5.33	10.60	14.88	12.79
SEm±	0.053	0.048	0.043	0.049	0.081	0.019
CD (P=0.05)	NS	NS	NS	NS	0.315	0.076
<i>B. Seed Priming</i>						
S1*	4.44	5.61	5.02	10.56	14.15	12.37
S2	4.64	5.97	5.30	10.58	14.52	12.58
S3	4.81	6.17	5.49	10.62	14.72	12.74
SEm±	0.076	0.071	0.057	0.083	0.089	0.072
CD (P=0.05)	0.235	0.219	0.175	NS	0.273	0.221

\* M1: Seeding at optimum moisture Level, M2: Seeding at sub optimal soil moisture level

M3: Seeding in dry soil followed by irrigation

S<sub>1</sub>: No seed priming, S<sub>2</sub>: Seed priming, S<sub>3</sub>: Sprouted seed

**Table 3: Interaction effects of seed priming and seeding method on crop establishment, growth and yield of wheat during the year 2011-11 and 2011-12 (Pooled analysis)**

Treatments	No. of Tillers m <sup>2</sup>	Earhead Length (cm)	Plant Height (cm)	Biomass (t ha <sup>-1</sup> )	Grain Yield (t ha <sup>-1</sup> )	1000 grain weight (g)
M <sub>1</sub> S <sub>1</sub>	469.68	9.90	87.57	12.25	4.98	38.68
M <sub>1</sub> S <sub>2</sub>	485.11	10.25	88.77	12.66	5.25	38.17
M <sub>1</sub> S <sub>3</sub>	520.41	10.57	86.94	12.56	5.61	39.78
M <sub>2</sub> S <sub>1</sub>	424.72	9.71	88.52	12.53	5.09	38.36
M <sub>2</sub> S <sub>2</sub>	449.99	10.73	88.09	12.17	5.16	38.65
M <sub>2</sub> S <sub>3</sub>	496.10	10.01	88.50	12.55	5.37	38.24
M <sub>3</sub> S <sub>1</sub>	500.55	9.22	87.44	12.33	5.01	37.92
M <sub>3</sub> S <sub>2</sub>	512.35	9.43	88.17	12.91	5.51	39.18
M <sub>3</sub> S <sub>3</sub>	485.68	9.95	88.42	13.12	5.49	39.22
SEm±	10.73	0.30	0.59	0.12	0.09	0.23
CD (P=0.05)	33.06	NS	0.95	0.38	NS	0.72

\* M1: Seeding at optimum moisture Level, M2: Seeding at sub optimal soil moisture level

M3: Seeding in dry soil followed by irrigation

S<sub>1</sub>: No seed priming, S<sub>2</sub>: Seed priming, S<sub>3</sub>: Sprouted seed

**Table 4: Effect of sprouted seeds and seed priming on water productivity of wheat crop**

Treatment	Total irrigation water* application (m ha <sup>-3</sup> )			Water productivity per ha (kg grain/m <sup>3</sup> water)		
	2010-11	2011-12	Pooled	2010-11	2011-12	Pooled
M <sub>1</sub> S <sub>1</sub>	3642	3612	3627	1.22	1.52	1.37
M <sub>1</sub> S <sub>2</sub>	3642	3612	3627	1.30	1.60	1.45
M <sub>1</sub> S <sub>3</sub>	3642	3612	3627	1.32	1.77	1.55
M <sub>2</sub> S <sub>1</sub>	3042	3012	3027	1.47	1.89	1.68
M <sub>2</sub> S <sub>2</sub>	3042	3012	3027	1.50	1.91	1.70
M <sub>2</sub> S <sub>3</sub>	3042	3012	3027	1.52	2.02	1.77
M <sub>3</sub> S <sub>1</sub>	3642	3612	3627	1.21	1.56	1.38
M <sub>3</sub> S <sub>2</sub>	3642	3612	3627	1.27	1.76	1.52
M <sub>3</sub> S <sub>3</sub>	3642	3612	3627	1.36	1.67	1.51

Note: \* Calculated on the basis of irrigation water @ 600 m ha<sup>-3</sup> per irrigation + rainfall during the crop season, 2010-11 = 129.7 mm, and 2011-12 = 36.3mm.

**Table 5: Cost of irrigation influenced by seeding of hydro-primed and pre-germinated wheat seeds**

Treatment	Irrigation water					
	2010-11		2011-12		Pooled	
	Quantity (m ha <sup>-3</sup> )	Cost (INR ha <sup>-1</sup> )	Quantity (m ha <sup>-3</sup> )	Cost (INR ha <sup>-1</sup> )	Quantity (m ha <sup>-3</sup> )	Cost (INR ha <sup>-1</sup> )
M <sub>1</sub> S <sub>1</sub>	3600	2172	3600	2292	3600	2232
M <sub>1</sub> S <sub>2</sub>	3600	2172	3600	2292	3600	2232
M <sub>1</sub> S <sub>3</sub>	3600	2172	3600	2292	3600	2232
M <sub>2</sub> S <sub>1</sub>	3000	1810	3000	1910	3000	1860
M <sub>2</sub> S <sub>2</sub>	3000	1810	3000	1910	3000	1860
M <sub>2</sub> S <sub>3</sub>	3000	1810	3000	1910	3000	1860
M <sub>3</sub> S <sub>1</sub>	3600	2172	3600	2292	3600	2232
M <sub>3</sub> S <sub>2</sub>	3600	2172	3600	2292	3600	2232
M <sub>3</sub> S <sub>3</sub>	3600	2172	3600	2292	3600	2232
Mean	3400	2051	3400	2164	3400	2108

\* M1: Seeding at optimum moisture Level, M2: Seeding at sub optimal soil moisture level, M3: Seeding in dry soil followed by irrigation, S<sub>1</sub>: No seed priming, S<sub>2</sub>: Seed priming, S<sub>3</sub>: Sprouted seed

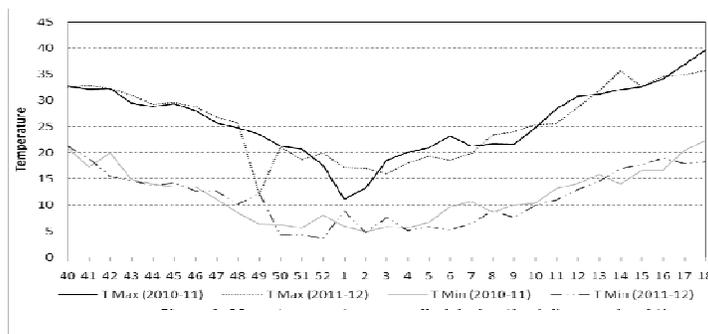


Figure 1: Weather data for the *Rabi* season of 2010-11 & 2011-12

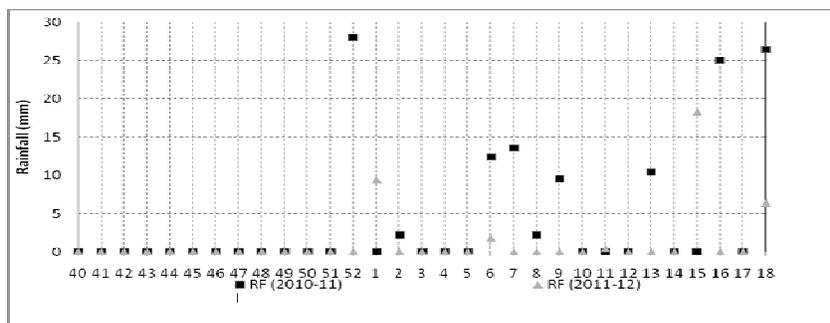


Figure 2: Mean rainfall prevailed during the Julian weeks of the crop season

## GENETIC VARIABILITY AND CHARACTER ASSOCIATION IN DOLICHOS BEAN

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### ABSTRACT

Field bean (*Dolichos lablab* L.) is an important vegetable crop throughout India due to its local acceptability by the people. It has wide genetic variability for various traits like plant habit, branching habit, stem pigmentation, pod colour and pod characters etc. In the past, very little attention was given by the scientists on systematic crop improvement in Dolichos bean. The present investigation was laid out in a simple randomized complete block design with three replications during *rabi* 2011-2012 to study the variability present in 23 genotypes of dolichos bean. Analysis of variance revealed that there were significant differences among the genotypes for all the characters studied. The difference between PCV and GCV was narrow for all characters except percentage of pod set. Hence, these characters were less influenced by environment. High GCV and heritability estimates were associated with greater genetic advance for the nine traits *viz.*, percentage of pod set, number pods per cluster, number pods per plant, green pod length, green pod width, individual green pod weight, pod yield, crude protein and crude fibre indicating that these characters had additive gene effect and were more reliable for effective selection in the improvement of dolichos bean. It is inferred from correlation and path analysis that the individual green pod weight, percentage of pod set, number of flowers per cluster and number of pods per plant exhibited significant positive correlation and direct positive effect on yield. Hence, these traits may be used as selection indices for yield improvement of dolichos bean.

**Key words:** multivariate analysis, pod characters, selection, sem

### INTRODUCTION

Dolichos bean [*Lablab prupureus* (L.) Sweet] is an important leguminous vegetable of India and is mainly grown for its tender pods which are cooked and

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consumed as vegetable. It is also called as sem, hyacinth bean, field bean and avarai. Being leguminous vegetable, the immature green pods of dolichos bean is a good source of protein, minerals and vitamins (Basu et al., 1999). Based on historical evidences, India is considered as origin and primary centre of diversity for dolichos bean. It is one of the most ancient crop among the cultivated plants grown as either pure or mixed with other crops, such as finger millet, groundnut, castor, corn or sorghum. It is also grown in homestead. It is a multipurpose crop grown as pulse, vegetable and forage. Field bean is a drought tolerant crop grown in dry lands with limited rainfall. The crop prefers relatively cool season when sowing done in July-August. It starts fruiting in winter and continues indeterminately in spring (Savitha, 2008). Despite having many good attributes, the crop has remained unexploited owing to low productivity, long duration, photosensitivity and indeterminate growth habit. The consumer preference also varies with pod size, shape, colour and aroma (Sogadu). The efforts of improving the crop by utilizing indigenous and exotic germplasm have been useful in breaking the yield barriers resulting in compact plant type, reduced duration and photo-insensitive types. Hence, comprehensive germplasm collection and evaluation, identification of suitable genotypes for pure crop and investigation of its value as an intercrop with other food and forage crops are essential. The crop is mainly grown for its green pods, while the dry seeds are used in various vegetable preparations. It is one of the major sources of protein in South Indian dietary. A wide range of variations exist for the plant and pod characters amongst the accessions grown all over the country. The success of any breeding programme in general and improvement of specific trait through selection in particular, totally depends upon the genetic variability present in the available germplasm of a particular crop (Parmar et al. 2013). Since, many of the plant characters are governed by polygenes and greatly influenced by environmental conditions; the progress of breeding is, however, conditioned by the magnitude, nature and interrelationship of genotypic and non-genotypic variation. Among the quantitative characters, yield is a complex character, which is dependent on a number of yield contributing characters (Savitha, 2008). The knowledge of the association of yield components and their relative contribution shown by path analysis has practical significance in selection. The study of the association between pairs of characters and yield provide basis for further breeding programme. With a view to measuring the variability among the 23 genotypes for flower and pod characters, getting information on the associations of flower and pod characters with pod yield and assessing the degree of divergence among the genotypes as well as assessing the relative contribution of different components, the present study was undertaken with the following objectives. (i) to study the nature and magnitude of variability in dolichos genotypes; (ii) to understand the degree and direction of association between yield and its components and among themselves; (iii) to assess the direct and indirect effects of traits on yield.

## MATERIALS AND METHODS

The present evaluation of Dolichos bean genotypes suitable for coastal region of Karaikal in U.T. of Puducherry was conducted in the college orchard, Department of Horticulture, Pandit Jawaharlal Nehru College of Agriculture and Research Institute, Karaikal during *rabi* season 2012. The experimental material consisted of 23 genotypes of dolichos bean collected from NBPGR (National Bureau of Plant Genetic Resources), New Delhi; TNAU (Tamil Nadu Agricultural University); IIHR (Indian Institute of Horticultural Research), and local varieties of Tamil Nadu and Puducherry. The details of the accessions are given in table 1. The experiment was laid out in the Eastern block of PAJANCOA & RI, Karaikal during November-March, 2012. The experimental site is situated between 10°49' and 11°01' N Latitude and 78°43' and 79°52' E Longitude with an altitude of four meters from the sea level. The soil type of the experimental plot was clay loam with medium fertility. Karaikal has tropical climate and receives an average annual rainfall of 915.5 mm in 44 rainy days. The farm pond water was used for irrigating the experimental field. The field experiment was laid out in a randomized block design with three replications during *rabi* season 2012. The plants were spaced 60 cm between row and 30 cm between plants in a row. Recommended horticultural practices and plant protection measures were followed uniformly. The observations were recorded from ten randomly selected plants per replication on each genotype for 15 characters *viz.*, plant height, number of branches per plant, days to 50 percent flowering, number of flowers per cluster, number of clusters per plant, percentage of pod set, days to maturity, number of pods per cluster, number of pods per plant, green pod length, green pod width, individual green pod weight, pod yield per plant, crude protein content and crude fibre content. Crude protein content of the pods was estimated by Bremner (1965). Crude fibre content of pods was estimated by the method of Sadasivam and Manickam (1996). The mean values were subjected to statistical analysis. Correlation, variability and path coefficient analysis were carried out by using GENRES software.

## RESULTS AND DISCUSSION

### (I) Variability studies

Variability of a character is measured by range and genotypic coefficient of variation (Rahman et al., 2002). The range in the mean values reflects the extent of phenotypic variability present in the entries. In this way, the values include genotypic, environmental and genotype x environmental components. So, the estimation of genetic (heritable) and environmental (non-heritable) components of the total variability are required as these help us in the choice of suitable breeding programme. In the present investigation the difference between PCV and GCV was compared for different traits and it was found to be relatively narrow for almost all the characters except for percentage of pod set, indicating the presence of low degree

of environmental influence on these characters. Regarding the percentage of pod set, the difference between PCV and GCV was moderately higher which indicates that this trait is much influenced by environment. Since GCV represents the heritable components of total variation, it would be more appropriate to use this parameter for comparing variability of different characters among genotypes in dolichos bean.

In the present investigation, high GCV was recorded for percentage of pod set, number of pods per cluster, number of pods per plant, green pod length, green pod width, individual green pod weight, green pod crude protein, green pod crude fiber and pod yield (Table 2). This indicated the maximum variability existing in the genotypes for these characters and offers good scope for improvement of these traits by simple selection. Similar findings were reported by Golani et al. (2007) and Mohan et al. (2009) in dolichos bean. Moderate GCV was observed for plant height, number of branches per plant, number of flower per cluster, number of cluster per plant and days to maturity, whereas lowest GCV was observed for days to 50 percent flowering. These lowest and moderate GCV estimates for these traits revealed that the extent of response of these traits for selection would be lesser than that of the other traits.

High genotypic coefficient of variation (GCV) alone is not sufficient for determination of the heritable variation, as it simply measures the extent of genetic variability present for a character. Hence, GCV together with heritability estimates would give the best picture of the advance to be expected by selection. In the present investigation, high heritability estimates were observed for all the characters studied (Table 2). High heritability ranged from 74.270 for number of branches per plant to 99.730 for green pod crude fiber. High heritability values obtained for most of the traits suggested that this might generally be governed by additive gene action and hence the phenotype could provide fairly reliable measure of genotypic effect. Hence, selection could be exercised on the phenotypic performance. Higher estimates of genetic advance ( $\geq 40\%$ ) as percent of mean were observed for 9 traits out of 15 (Table 2). Moderate genetic advance as percentage mean was observed for the rests. Heritability estimates in broad sense alone do not serve as the true indicator of genetic potentiality of the genotypes since the scope is restricted by their interaction with environment. High heritability coupled with high genetic advance as percent of mean was observed for 9 traits viz., percent of pod set, number pods per cluster, number pods per plant, green pod length, green pod width, individual green pod weight, green pod crude protein, green pod crude fiber and pod yield per plant. These traits have also high GCV. This indicates the presence of additive genes and consequently there is an ample scope of improving these characters by simple selection. High heritability coupled with moderate genetic advance as percent mean was observed for plant height, number of flower per cluster, number of cluster per plant, days to maturity, number of branches per plant and days to 50 percentage of flowering, indicate the presence of dominance and epistatic gene action effects in controlling these characters. Similar trends were observed by Bendale et al. (2004) in lablab bean.

**(ii) Correlation studies**

Although variability estimates provide information on the extent of improvement through selection but they do not throw much light on the extent and nature of relationship between the characters which could be utilized for selection following hybridization. The correlation studies revealed higher estimates of genotypic correlation coefficients than their corresponding phenotypic correlation coefficients, suggesting a strong inherent relationship between different traits and that the environmental factors had not played much role in expression of the phenotypic.

**Genotypic correlation coefficient for yield and quality traits**

The genotypic correlation of pod yield with yield components revealed that the pod yield per plant was positively and significantly associated with green pod width, number of flowers per cluster, pod length and individual green pod weight (Table 3). It implied that these characters strongly influenced the pod yield per plant. Percent of pod set and number of pods per plant exhibited non-significant positive correlation with pod yield. A similar trend was reported by Ali et al. (2005) and Roy et al. (2006).

Negative correlation was observed for days to 50 percent flowering and days to maturity. This association indicated that these characters shall be taken into consideration for improvement of dolichos bean crop for earliness. Significant negative correlation of other characters with green pod yield implied that there was a little scope for consideration of these characters for the improvement of green pod yield in dolichos bean. Based on the above discussion, it could be inferred that the most important pod characters such as green pod length, number of flowers per cluster, green pod width and individual green pod weight may be considered for intentional selection in simultaneous improvement on pod yield per plant. In addition to these characters, percentage of pod set and number of pods per plant can also be taken into consideration for improvement on pod yield. The parameters such as days to 50 percent flowering and days to maturity were negatively correlated with yield and hence these characters could be considered for earliness.

In the inter correlation among the components characters, plant height was significantly and positively correlated with days to 50 percent flowering and days to maturity. This result indicated that increase in plant height caused late flowering and consequently the maturity was also delayed. The percentage of pod set was positively correlated with number of pods per cluster and number of pods per plant. These results clearly indicated that the percentage of pod set is one most important trait for increasing the production of more number of pods per plant. However, increased percentage of pod set and increased number of pods per cluster as well as number of pods per plant were negatively correlated with green pod length, green pod width and individual green pod weight. The length of pod had positive association with green pod width and individual green pod weight. Therefore, knowledge on the inter correlation association of component traits with pod yield per plant will help making selection more precise and accurate.

### **Phenotypic correlation coefficient for yield and quality traits**

In the present investigation the phenotypic correlation coefficients also showed similar trend as exhibited by genotypic correlation wherein the green pod width had strong positive correlation with yield. In addition, the green pod width, green pod length, number of flowers per cluster, number of pods per plant and percent pod set had direct positive effect on yield (Table 3). It indicated that these characters can be considered for selection in the improvement of pod yield per plant. Similarly, the inter correlation among components traits, the flower characters as well as pod traits were inter associated among themselves.

From the above discussion it may be concluded that the most important traits viz., green pod length, green pod width, number of flowers per cluster and individual green pod weight were found to be positively and significantly correlated both at genotypic and phenotypic levels, indicating that these traits may be considered as very important in selection and any improvement in these four characters would bring about an enhancement in yield. For earliness, parameters such as days to 50 percent flowering and days to maturity may be taken into consideration. It is also suggested that hybridization of genotypes possessing combination of such characters will be useful for obtaining desirable segregants. Similar reports were reported by Sonali et al. (2009) in pea.

### **(iii) Path analysis**

Selection of superior genotypes based on yield as such may not be effective for the enhancement of yield and hence selection should be made for component traits as well. Association of yield components thus assumes a unique prominence as the basis for selecting desirable genotypes with high yield potential. In addition, knowledge on presence of association among component characters reveals that some of them may serve as indicators of yield. In the present study, the path analysis reveals that individual green pod weight exhibited very high direct positive effect (3.72) on pod yield per plant. In addition, percent pod set, number of clusters per plant, pods per plant and flowers per cluster and days to 50 percent flowering also showed very high direct positive effect on pod yield (Table 4). This indicated that these characters played an important role in increasing the pod yield. Therefore, the direct selection of these characters may about bring an overall improvement in pod yield. These findings were in agreement with the results reported Desai et al. (2003) in dolichos bean, where the number of branches per plant, green pod crude protein showed high and direct positive effects respectively on pod yield.

The days to maturity, pod length, plant height, green pod width and number of pods per cluster had negatively direct effect on yield per plant. The residual effect determines how best the causal factor accounts for the variability of the dependent factor that is yield per plant in this study. In the present investigation the residual effect of path coefficient analysis was 0.1519, which clearly indicated that the 15 characters taken for this investigation were sufficient for genetic analysis in dolichos

bean. Only 15% of the variability was controlled by other traits besides these 15 characters. Based on the above results the traits such as, individual green pod weight and number flowers per cluster, number of clusters per plant, percentage of pod set and number of pods per plant were important yield contributing traits which were positive and significantly correlated with pod yield and also showed high direct effect on pod yield. For earliness, days to 50 percent flowering had to be considered. Therefore, these traits shall be used as selection criteria for the improvement of yield directly in dolichos bean.

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**Table 1: Sources of Dolichos bean**

S. No.	Genotype designation used in text and tables	Accession name	Source
1	DB1	Dolichos Diana	Ashoka Seeds Private Limited, Bangalore
2	DB2	Dolichos Nandini	Prabhakar Seeds Private Limited, Bangalore
3	DB3	PHS Dhoni	Prabhakar Seeds Private Limited, Bangalore
4	DB4	Rohini	Rasi Seeds Private Limited, Salem
5	DB5	Ankur Goldy	Ankur Seeds Private Limited, Nagpur
6	DB6	Konkan	Dr. Balasheb Sawant Konkan Krishi Vidyapeeth, Dapoli, Maharashtra
7	DB7	Dolichos Ruchi	Indosun Agri Genetics Private Limited, Hyderabad
8	DB8	Dolichos Nandi	Aditya Seeds Private Limited, Bangalore
9	DB9	Arka Jay	Indian Institute of Horticultural Research, Bangalore
10	DB10	Arka Vijay	Indian Institute of Horticultural Research, Bangalore
11	DB11	COGB -14	Tamil Nadu Agricultural University, Coimbatore
12	DB12	Dolichos Senthil	Senthil Seeds & Company Private Limited, Bangalore
13	DB13	PHS Dolichos	Prabhakar Seeds Private Limited, Bangalore
14	DB14	Kumbakonam local	Kumbakonam, Tamil Nadu
15	DB15	Madagadipet local	Pondicherry
16	DB16	Ottanchathiram local	Tamil Nadu
17	DB17	IC 354334	National Bureau of Plant Genetic Resources, New Delhi.
18	DB18	IC 354336	National Bureau of Plant Genetic Resources, New Delhi.
19	DB19	IC 354337	National Bureau of Plant Genetic Resources, New Delhi.
20	DB20	Lakshmi	Super Seeds Private Limited, Hyderabad
21	DB21	Gold 24	Sun Star Seeds Private Limited, Hyderabad
22	DB22	Dolli	Rasi Seeds Private Limited, Haryana
23	DB23	Theni local	Tamil Nadu

**Table 2: Estimation of variability parameters for dolichos bean genotypes**

SI. No	Character	Phenotypic variance (PV)	Genotypic Variance (GV)	Phenotypic coefficient of variation (PCV %)	Genetic coefficient of variation (GCV %)	Heritability	Genetic advance	Genetic advance as percent of mean
1	Plant height	241.019	214.472	16.332	15.407	88.990	28.459	29.939
2	Number of branches per plant	0.648	0.481	12.811	11.041	74.270	1.232	19.602
3	Days to 50% flowering	16.130	15.741	8.180	8.081	97.590	8.074	16.445
4	Number of flowers per cluster	11.967	8.930	16.942	14.635	74.620	5.318	26.043
5	Number of clusters per plant	1.806	1.568	14.517	13.527	86.830	2.404	25.966
6	Percent pod set	42.264	33.011	31.049	27.440	78.110	10.460	49.958
7	Days to maturity	38.304	38.078	11.312	11.279	99.410	12.674	23.166
8	Number pods per cluster	5.318	4.911	29.656	28.498	92.350	4.387	56.416
9	Number pods per plant	83.499	72.614	24.008	22.389	86.960	16.370	43.010
10	Green pod length	2.779	2.626	21.475	20.876	94.500	3.245	41.805
11	Green pod width	0.203	0.185	25.764	24.569	90.940	0.845	48.263
12	Individual green pod weight	1.325	1.260	28.720	28.007	95.100	2.255	56.264
13	Green pod crude protein	4.313	4.258	23.102	22.954	98.730	4.224	46.985
14	Green pod crude fiber	8.43	8.411	39.646	39.593	99.730	5.966	81.453
15	Pod yield per plant	827.571	797.648	20.373	20.002	96.380	57.118	40.452

Variability in field bean

**Table 3: Genotypic correlation coefficient for different characters of dolichos bean**

	2	3	4	5	6	7	8	9	10	11	12	13	14	15
1	0.049	0.488*	-0.222	0.130	-0.274	0.148*	-0.352	-0.408*	0.274	0.277	0.331	0.290	0.008	-0.163
1a	0.025	0.454*	-0.147	0.094	-0.234	0.393	-0.303	-0.352	0.225	0.260	0.294	0.284	0.005	-0.139
2		-0.278	-0.213	0.063	-0.098	-0.361	0.076	-0.190	-0.160	-0.010	-0.056	-0.133	0.245	-0.262
2a		-0.241	-0.169	0.018	-0.061	-0.306	0.051	-0.171	0.119	-0.025	-0.050	-0.126	0.213	-0.225
3			-0.144	0.014	0.140	0.944**	0.147	0.010	0.229	-0.286	-0.201	0.504*	0.143	-0.343
3a			-0.107	0.027	0.114	0.927**	0.151	0.014	-0.215	-0.255	-0.191	0.494*	0.138	-0.332
4				0.065	-0.311	-0.047	-0.074	0.165	-0.185	0.316	-0.033	-0.154	-0.280	0.416*
4a				0.071	-0.421*	-0.039	-0.073	0.111	-0.117	0.307	-0.007	-0.139	-0.241	0.357
5					-0.452	0.086	0.254	0.045	-0.228	-0.307	-0.301	0.162	-0.248	-0.523**
5a					-0.420*	0.083	0.255	0.070	-0.211	-0.268	-273	0.151	-0.232	-0.459*
6						0.051	0.513*	0.750**	-0.349	-0.425*	-0.561**	-0.090	0.266	0.150
6a						0.044	0.468*	0.702**	-0.331	-0.370	-0.541**	-0.077	0.233	0.132
7							0.083	0.046	-0.241	-0.261	-0.163	0.487*	0.017	-0.285
7a							0.079	0.043	-0.235	-0.252	-0.159	0.482*	0.017	-0.277
8								0.681**	-0.734**	-0.730**	-0.854**	0.042	0.332	-0.404*
8a								0.661**	-0.685**	-0.654**	-0.819**	0.047	0.315	-0.361
9									-0.677**	-0.525**	-0.821**	-0.106	0.004	0.126
9a									-0.620**	-0.453*	-0.811**	-0.099	0.002	0.146
10										0.613**	0.835**	-0.092	-0.149	0.409*
10a										0.589**	0.806**	-0.089	-0.186	0.390
11											0.763**	-0.287	-0.243	0.520**
11a											0.711**	-0.277	-0.234	0.489*
12												-0.160	-0.224	0.406*
12a												-0.155	-0.216	0.385
13													0.190	-0.304
13a													0.188	-0.298
14														-0.302
14a														-0.297

\*, \*\* Significant at 5 and 1% levels. 'a' denotes phenotypic correlation co-efficients.

1 = Plant height, 2 = Number of branches per plant, 3= Days to 50%age flowering, 4 = Number of flowers per cluster, 5 = Number of cluster per plant, 6 = Percent pod set, 7 = Days to maturity, 8 = Number of pods per cluster, 9 = Number of pods per plant, 10 = Green pod length, 11 = Green pod width, 12 = Individual green pod weight, 13 = Green pod crude protein, 14 = Green pod crude fiber, 15 = Pod yield per plant.

**Table 4. Path coefficient for direct and indirect effects of different traits**

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
<b>1</b>	<b>-0.5017</b>	0.0321	0.5871	-0.2256	0.1479	-0.4994	-0.4426	0.0830	-0.4582	-0.2089	-0.0671	1.2318	0.1579	0.0005	-0.163
<b>2</b>	-0.0246	<b>0.6547</b>	-0.3348	-0.2164	0.0721	-0.1789	0.3817	-0.0180	-0.2133	-0.1220	0.0023	-0.2090	-0.0723	0.0163	-0.262
<b>3</b>	-0.2448	-0.1822	<b>1.2030</b>	-0.1469	0.0161	0.2559	-0.9996	-0.0346	0.0116	0.1741	0.0693	-0.7492	0.2747	0.0095	-0.343
<b>4</b>	0.1113	-0.1393	-0.1737	<b>1.0174</b>	0.0741	-0.5673	0.0492	0.0174	0.1854	0.1412	-0.0766	-0.1209	-0.0838	-0.0186	0.416*
<b>5</b>	-0.0650	0.0414	0.0170	0.0661	<b>1.1406</b>	-0.8250	-0.0910	-0.0600	0.0507	0.1739	0.0744	-1.1180	0.0885	-0.0165	-0.523**
<b>6</b>	0.1373	-0.0642	0.1687	-0.3164	-0.5157	<b>1.8245</b>	-0.0544	-0.1210	0.8411	0.2656	0.1030	-2.0870	-0.0492	0.0177	0.150
<b>7</b>	-0.2098	-0.2361	1.1362	-0.0473	0.0981	0.0938	<b>-1.0584</b>	-0.0197	0.0511	0.1840	0.0631	-0.6072	0.2656	0.0011	-0.285
<b>8</b>	0.1765	0.0498	0.1764	-0.0750	0.2902	0.9354	-0.0882	<b>-0.2359</b>	0.7643	0.5592	0.1769	-3.1782	0.0228	0.0221	-0.404*
<b>9</b>	0.2048	-0.1244	0.0124	0.1681	0.0516	1.3675	-0.0482	-0.1607	<b>1.1222</b>	0.5160	0.1271	-3.0526	-0.0578	0.0003	0.126
<b>10</b>	-0.1375	0.1048	-0.2749	-0.1885	0-0.2602	-0.6359	0.2556	0.1731	-0.7598	<b>-0.7621</b>	-0.1485	3.1053	-0.0499	-0.0129	0.409*
<b>11</b>	-0.1389	-0.0062	-0.3442	0.3217	-0.3505	-0.7758	0.2759	0.1722	-0.5887	-0.4671	<b>-0.2422</b>	2.8366	-0.1564	-0.0162	0.520**
<b>12</b>	-0.1661	-0.0368	-0.2423	-0.0331	-0.3428	-1.0236	0.1728	0.2016	-0.9209	-0.6362	-0.1847	<b>3.7200</b>	-0.0873	0.0149	0.406*
<b>13</b>	-0.1453	-0.0868	0.6060	-0.1563	0.1852	-0.1647	-0.5156	-0.0099	-0.1190	0.0698	0.0695	-0.5954	<b>0.5453</b>	0.0127	-0.304
<b>14</b>	-0.0039	0.1602	0.1714	-0.2850	-0.2828	0.4854	-0.0178	-0.0783	0.0042	0.1476	0.0588	-0.8323	0.1036	<b>0.0666</b>	-0.302

\*\* Significant at 1% level \* Significant at 5% level

1 = Plant height, 2 = Number of branches per plant, 3= Days to 50%age flowering, 4 = Number of flowers per cluster, 5 = Number of cluster per plant, 6 = Percent pod set, 7 = Days to maturity, 8 = Number of pods per cluster, 9 = Number of pods per plant, 10 = Green pod length, 11 = Green pod width, 12 = Individual green pod weight, 13 = Green pod crude protein, 14 = Green pod crude fiber, 15 = Pod yield per plant.

**Short Note**

**INTEGRATED NUTRIENT MANAGEMENT ON  
PRODUCTIVITY OF CARROT AND FERTILITY OF SOIL**

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Carrot (*Daucus carota* L.) belongs to the family umbelliferae and is an important root vegetable grown throughout the world. Among various factors responsible for low production of carrot, nutrient management is of prime importance for maintaining higher yield and soil fertility. It has been reported that neither the chemical fertilizer alone nor the organic manure are able to sustain the crop productivity and soil fertility. The increasing use of chemical fertilizers to increase vegetable production has been widely recognized but its long run impact on soil health, ecology and other natural resources are detrimental which affect living organisms including beneficial soil microorganism and human being. The escalating prices of chemical fertilizers and its detrimental impact on the soil, environment and human health urged the farmer to adoption of integrated plant nutrient that offers the sustainable crop production and soil fertility (Sentiyangla et al., 2010). Besides fertilizers, there are several sources of plant nutrients like organic manures, biofertilizers etc. These nutrients sources not only reduce quantity of chemical fertilizers but also improve soil fertility (Chumyani et al., 2012). Use of organic manures in INM help mitigating multiple nutrient deficiencies. Application of organic manures to acidic soil reduces the soluble and exchangeable Al temporarily by forming complex and provides better environment for growth and development in addition to improvement in physical, chemical and biological properties of soil (Avitoli et al., 2012). Biofertilizers have also emerged as promising components of nutrient supply system. Application of biofertilizers which is environment friendly and low cost input, with organic and inorganic fertilizers as part of an integrated nutrient management strategy and play significant role in plant nutrition.

A field experiment was conducted during 2011 - 2012 at the Experimental Farm of SASRD, Medziphema campus, Nagaland University, Nagaland. The field is located at the altitude of 304.8 m above mean sea level with geographical location at 20° 45' 43" N latitude and 93 ° 53' 04" E longitudes. The soil of the experimental site was sandy loam having soil pH 4.4, organic carbon 1.60 % and available N, P and K content of 305.76, 17.00 and 225.25 kg ha<sup>-1</sup> respectively. The experiment was laid out in a randomized block design with three replications. Plot size measured 1.8 m x

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1.8 m and spacing was maintained at 30 x 10 cm. Seeds were sown on 5 October, 2011 and thinning was done 10 days after sowing to maintain spacing. The treatments consisted of T<sub>1</sub> - Control, T<sub>2</sub> - FYM @ 20 t ha<sup>-1</sup>, T<sub>3</sub> - Pig manure @ 15 t ha<sup>-1</sup>, T<sub>4</sub> - Vermicompost @ 5 t ha<sup>-1</sup>, T<sub>5</sub> - 100% recommended dose of NPK (80:40:40 kg ha<sup>-1</sup>), T<sub>6</sub> - 50% NPK + 50% FYM, T<sub>7</sub> - 50% NPK + 50% Pig manure, T<sub>8</sub> - 50% NPK + 50% Vermicompost, T<sub>9</sub> - 50% NPK + 50% FYM + Biofertilizers, T<sub>10</sub> - 50% NPK + 50% Pig manure + Biofertilizers, T<sub>11</sub> - 50% NPK + 50% Vermicompost + Biofertilizers. N, P and K were given through Urea, SSP and MOP respectively. Full dose of P and K and half dose of N were applied at the time of transplanting and remaining half dose of N was given 45 days after transplanting. Manures viz., FYM, pig manure and vermicompost were incorporated as per treatment to respective plots prior to transplanting. Biofertilizers (*Azospirillum* and *Phosphotika*) were inoculated to seeds prior to sowing as seed treatment methods @ 500 g ha<sup>-1</sup> each. Observations on plant height, number of leaves, root length, root diameter, root weight, root yield and carotene content were recorded at harvesting. Carotene content was determined by spectrophotometer method (Rangama, 1977) and expressed in mg 100<sup>-1</sup>g.

Soil samples were collected before and after harvest of crop from different locations of the experimental plot to a depth of 15 cm with the help of screw type auger. The collected soil samples were mixed and reduced to 500 g, dried under shade and ground to pass 2 mm sieve. Soil samples were analysed for pH, organic carbon, available nitrogen, phosphorus and potassium which were determined by Digital pH meter, Walkley and Black Rapid titration method, Alkaline potassium permanganate method, Olsen's method, flame photometer method, respectively (Jackson, 1973). Statistical analysis was carried out as per standard procedure.

Economics of the treatments were calculated as per prevailing market price of input and output. Gross income was calculated by yield multiplied with whole sale rate of carrot (Rs. 10000 t<sup>-1</sup>). Net income was estimated by deducting the total cost of cultivation (fixed cost + treatment cost) from gross income of the particular treatment. Cost-benefit ratio was worked out by dividing net return from total cost of cultivation.

Improvement in growth characters is considered to be a pre-requisite to increase the yield. NPK fertilizers with different organic manures along with biofertilizers alone or in combination were found to have significant effect on growth characters as compared to control (Table-1). All the treatments were found effective in increasing the plant growth over control. Application of 50% NPK + 50% FYM + biofertilizers (T<sub>9</sub>) recorded maximum plant height (25.00 cm) and number of leaves (6.98). The lowest values of plant height and number of leaves were recorded with control. The increase in vegetative growth might be due to the role of nitrogen in promoting vegetative growth and enhancing cell division and elongation as well as greater chlorophyll synthesis, phosphorus is easily mobilized in the plant and translocated to the meristematic zone and increase the activity of leaf formation and

development in carrot and potassium activates many enzymes involved in respiration and photosynthesis. The added FYM in integrated nutrient management would have improved the physical, chemical and biological properties of soil which helps better nutrient absorption and utilization by plant resulting better plant growth. Rani et al. (2006) reported that application of neem cake and castor cake in combination with half the recommended dose of NPK recorded maximum growth characters in carrot. Sentiyangla et al. (2010) reported significant increase in plant height and number of leaves in radish when applied integrated application of chemical fertilizers, organic manures and biofertilizers (50% NPK + 50% FYM + biofertilizers). Subenthung et al. (2012) reported that combined application of 50% Pig manure + 50% NPK recorded maximum plant height (50.16 cm) and number of leaves (14.43), leaf area (185.86 cm<sup>2</sup>) and root yield (522.51 t ha<sup>-1</sup>) in turnip. These results are in conformity with the finding of Chumyani et al., 2012 in tomato and Vimera et al., 2012 in king chilli, they found that maximum growth characters with 50% NPK + 50% FYM + biofertilizers.

Integrated application of chemical fertilizers, organic manures and biofertilizers alone or in combination significantly increased the yield and yield attributing characters of carrot compared to control (Table-1). Application of 50 % NPK + 50 % FYM + biofertilizers (T<sub>9</sub>) recorded maximum values of all yield attributing characters such as root length (18.88 cm), root diameter (4.14 cm), root weight (90.37 g). This result indicates positive effects of integrating NPK with manures as well as biofertilizers. Integrated application of organic manure and inorganic fertilizer increased the availability of NPK and also improved the fertility status of soil and productivity due to which yield attributing characters might have increased. Besides NPK, micronutrients might have played an important role in increasing the yield attributing characters of carrot as addition of FYM increased the availability of micronutrients. Also, biofertilizers might have played a vital role in increasing the yield and yield related attributes. Root yield per hectare was recorded highest (30.88 t) in the treatment combination of 50% NPK + 50% FYM + Biofertilizers (T<sub>9</sub>) which was significantly superior over other treatment except T<sub>10</sub> (50% NPK + 50% Pig manure + Biofertilizers) and minimum root yield was recorded in control. Application of 50% NPK + 50% FYM + Biofertilizers produced 15% higher yield over 100% recommended doze of NPK. This might be due to corresponding response to increased growth and yield attributing characters attained previously under this treatment. Sagiv et al. (1994) reported that highest yield was obtained in carrot when organic manure, composted refuses and N fertilization was applied in combination. Rani et al. (2006) reported that application of neem cake and castor cake in combination with half the recommended dose of NPK recorded maximum root yield (25.860 t ha<sup>-1</sup>) in carrot. Sentiyangla et al. (2010) observed maximum yield in radish by combination of NPK, FYM and biofertilizers. Subenthung et al. (2012) observed that maximum root yield was obtained in the combined application of 50% pig manure + 50% NPK in turnip. Similarly Chumyani

et al. (2012) and Vimera et al. (2012) also conducted an experiment on integrated nutrient management and found that 50% NPK + 50% FYM + Biofertilizers recorded maximum yield in tomato and king chilli, respectively.

Quality of carrot is usually evaluated by carotene content. Various organic manures, inorganic fertilizers and biofertilizers and their combination had a beneficial impact on carotene content in roots. It is evident from table-1 that maximum carotene (3.41 mg 100<sup>-1</sup>g) was recorded with 50 % NPK + 50 % FYM + biofertilizers (T<sub>9</sub>). The comparative higher level of carotene might be due to the action of specific soil nutrients which might be made more readily available into the soil for plant absorption as a result of mineral fertilizer + lone organic manure 'or' with biofertilizers integration effect which might have activated specific enzymes for the synthesis of carotene in carrot. Nakagawa et al. (2003) reported that carotene content in roots was increased by application of organic fertilizers. Rani et al. (2006) reported that application of neem cake and castor cake in combination with half the recommended dose of NPK recorded highest carotene content (3.96 mg 100<sup>-1</sup> g) in carrot. Sunandarani and Malareddy (2007) reported that Neem cake + 50% RD of NPK gave the highest carotene content (4.60 mg 100<sup>-1</sup> g) in carrot.

Sustainability of a cropping system is being evaluated on the basis of crop yield as well as nutrient status of the soil after harvest of the crop. Different treatments alone and their combination with biofertilizers showed profound residual effect on soil fertility after harvest. However, their intensity varied considerably because of quantum variation (Table-2). Maximum available nitrogen (314.92 kg ha<sup>-1</sup>), phosphorous (19.59 kg ha<sup>-1</sup>) and potassium (250.42 kg ha<sup>-1</sup>) was recorded with treatment 100% NPK (T<sub>5</sub>) which was found at par with 50% NPK + 50% FYM + Biofertilizers (T<sub>9</sub>), 50% NPK + 50% Pig manure + Biofertilizers (T<sub>10</sub>), 50% NPK + 50% Vermicompost + Biofertilizers (T<sub>11</sub>). Maximum available nitrogen phosphorous and potassium in treatment 100% NPK might be due to poor soil physical structure, lack of organic manures and microbial activities, thus resulting in poor utilization of NPK by plants. As such the applied NPK could bring about higher residual NPK in soil after harvest. Similar result was also reported by Vimera et al. (2012) who reported that application of 100% NPK fertilizers alone recorded maximum available NPK in soil after harvest in king chilli. Organic carbon of soil acts as a sink and source of nutrients for microbial population, which regulates the availability of different nutrients through microbial transformation. The net increase in organic carbon was much higher with organic manures in combination with biofertilizers and fertilizers over 100% NPK alone. Application of 50% NPK + 50 % FYM + biofertilizers (T<sub>9</sub>) recorded maximum organic carbon (1.85 %) and soil pH (4.65) after harvest. Application of 50% NPK + 50 % FYM + biofertilizers recorded 16 % higher organic carbon over 100% NPK alone. This might be due to increased microbial activities in the root zone which decomposed organic manures and also fixed unavailable form of mineral nutrients into available forms in soil thereby substantiated crop requirements and improved organic carbon level and stabilized

soil pH. Chaudhary et al. (2005) reported that the incorporation of biofertilizers and FYM with inorganic fertilizers significantly improved the organic carbon content and pH of the soil in tomato. Similar results were also reported by Chumyani et al. (2012) in tomato who found maximum organic carbon and soil pH under treatment of 50% NPK + 50 % FYM + biofertilizers.

It is evident from table-3 that the integration of 50% NPK + 50 % FYM + biofertilizers (T<sub>9</sub>) was found to be the most profitable treatment in carrot exhibiting highest net return Rs. 2,36,193 with cost benefit ratio of 1:3.65 followed by Rs. 2,21,723 with the application of 50% NPK + 50% Pig manure + Biofertilizers (T<sub>10</sub>). The reason of high profitability in these two modes of integration can be due to lower cost of inputs and higher yield. Similar results were also reported by Chumyani et al. (2012) in tomato and Vimera et al. (2012) in king chilli. They found highest net return with the combined application of 50% NPK + 50% FYM + Biofertilizers.

It can be concluded from the experiment that integrated application of 50% NPK + 50% FYM + Biofertilizers was found optimum for getting maximum productivity of carrot without reducing fertility status of soil. This treatment reduced 50% chemical fertilizers without any compromise on yield of carrot and fertility of soil. Therefore, 50% NPK + 50% FYM + Biofertilizers may be recommended for sustainable yield of carrot and to nourish the soil fertility under foothills condition of Nagaland.

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**Table 1: Effect of integrated nutrient management on growth, yield and quality of carrot**

Treatments	Plant height (cm)	Number of leaves	Root length (cm)	Root diameter (cm)	Root weight (g)	Root yield (t ha <sup>-1</sup> )	Carotene content (mg 10 <sup>-1</sup> g)
T <sub>1</sub> Control	17.93	6.13	14.99	3.20	35.65	11.86	2.22
T <sub>2</sub> FYM 20t ha <sup>-1</sup>	20.93	6.77	15.69	3.53	41.29	13.73	2.51
T <sub>3</sub> Pig manure 15t ha <sup>-1</sup>	20.73	6.70	16.04	3.42	43.04	14.31	2.50
T <sub>4</sub> Vermicompost 5t ha <sup>-1</sup>	20.13	6.30	15.25	3.23	40.74	13.55	2.42
T <sub>5</sub> 100% NPK ((80:40:40 kg ha <sup>-1</sup> )	21.93	6.67	16.59	3.87	78.37	26.09	2.69
T <sub>6</sub> 50% NPK + 50% FYM	22.60	6.77	17.06	3.78	76.33	25.41	2.78
T <sub>7</sub> 50% NPK + 50% Pig manure	21.67	6.70	17.01	3.71	72.60	24.17	2.76
T <sub>8</sub> 50% NPK + 50% Vermicompost	21.53	6.67	17.00	3.68	70.77	23.56	2.64
T <sub>9</sub> 50% NPK + 50% FYM + Biofertilizers	25.00	6.98	18.88	4.14	90.37	30.08	3.41
T <sub>10</sub> 50% NPK + 50% Pig manure+ Biofertilizers	23.67	6.94	18.60	4.13	86.07	28.66	3.09
T <sub>11</sub> 50% NPK + 50% Vermicompost+Biofertilizers	22.00	6.90	18.23	4.13	79.00	26.31	3.02
SEm±	0.81	0.8	0.47	0.10	1.45	0.47	0.07
CD (P=0.05)	2.47	0.24	1.49	0.34	4.39	1.46	0.21

**Table 2: Effect of integrated nutrient management on the nutrient status of the soil after harvest**

Treatments	Available N (kg ha <sup>-1</sup> )	Available P (kg ha <sup>-1</sup> )	Available K (kg ha <sup>-1</sup> )	Organic carbon (%)	Soil pH
T <sub>1</sub> Control	216.39	14.21	188.50	1.43	4.38
T <sub>2</sub> FYM 20t ha <sup>-1</sup>	264.91	18.62	236.21	1.69	4.54
T <sub>3</sub> Pig manure 15t ha <sup>-1</sup>	256.56	17.87	237.06	1.64	4.51
T <sub>4</sub> Vermicompost 5t ha <sup>-1</sup>	266.16	16.82	230.21	1.59	4.48
T <sub>5</sub> 100% NPK ((80:40:40 kg ha <sup>-1</sup> )	314.92	19.59	250.42	1.60	4.42
T <sub>6</sub> 50% NPK + 50% FYM	301.84	18.60	238.90	1.77	4.59
T <sub>7</sub> 50% NPK + 50% Pig manure	297.36	18.45	233.65	1.76	4.55
T <sub>8</sub> 50% NPK + 50% Vermicompost	287.46	18.11	230.96	1.71	4.46
T <sub>9</sub> 50% NPK + 50% FYM + Biofertilizers	309.17	19.40	246.18	1.85	4.65
T <sub>10</sub> 50% NPK + 50% Pig manure + Biofertilizers	307.98	19.38	245.94	1.81	4.60
T <sub>11</sub> 50% NPK + 50% Vermicompost + Biofertilizers	306.49	19.09	241.32	1.74	4.56
SEm <sub>±</sub>	3.70	0.29	3.46	0.03	0.01
CD (P=0.05)	11.13	0.90	10.40	0.11	0.05

**Table 3: Effect of integrated nutrient management on economics of the treatments**

Treatments	Fixed cost (Rs.)	Treatment cost (Rs.)	Total cost (Rs.)	Root yield (t ha <sup>-1</sup> )	Gross income (Rs ha <sup>-1</sup> )	Net income (Rs ha <sup>-1</sup> )	Cost benefit ratio
T <sub>1</sub> Control	55000	0	55000	11.86	118670	63670	1: 1.15
T <sub>2</sub> FYM 20t ha <sup>-1</sup>	55000	10000	65000	13.73	137330	72330	1: 1.11
T <sub>3</sub> Pig manure 15t ha <sup>-1</sup>	55000	10500	65500	14.31	143110	77610	1: 1.18
T <sub>4</sub> Vermicompost 5t ha <sup>-1</sup>	55000	50000	105000	13.55	135550	30550	1: 0.29
T <sub>5</sub> 100% NPK (80:40:40 kg ha <sup>-1</sup> )	55000	9273	64273	26.09	160880	96607	1: 3.06
T <sub>6</sub> 50% NPK + 50% FYM	55000	9636	64636	25.41	164000	99364	1: 2.93
T <sub>7</sub> 50% NPK + 50% Pig manure	55000	9886	64886	24.17	168440	103554	1: 2.73
T <sub>8</sub> 50% NPK + 50% Vermicompost	55000	29636	84636	23.56	235550	150914	1: 1.78
T <sub>9</sub> 50% NPK + 50% FYM + Biofertilizers	55000	9687	64687	30.08	300880	236193	1: 3.65
T <sub>10</sub> 50% NPK + 50% Pig manure + Biofertilizers	55000	9937	64937	28.66	286660	221723	1: 3.41
T <sub>11</sub> 50% NPK + 50% Vermicompost + Biofertilizers	55000	29687	84687	26.31	263110	178423	1: 2.10

**Short Note**

**MORPHOLOGICAL CHARACTERIZATION OF  
CAULIFLOWER VARIETIES/CULTIVARS USING DUS  
CHARACTERS**

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Cauliflower is a cool season crop; it is more exacting in its climatic requirements than most other crops in this family. The plant is extremely sensitive to unfavourable conditions, such as unusually hot weather, drought or too low temperature, which often result in the formation of premature curds. It is monogenic species whose genomic constitution is C and chromosome number is  $n=9$  belongs to Cruciferae family (Thamburaj and Singh, 2001). The variety attains acceptance when the farmers get genetically pure seed of high standard. For the purpose, each cultivar should be properly defined with suitable descriptors, so as to maintain its identity during seed production through field inspection and certification. In India, Protection of Plant Varieties and Farmer's Right Act, 2001 (PPVandFRA, 2001) envisages the registration and protection of new and notified/extant plant varieties based on the criteria of Distinctness, Uniformity and Stability (DUS) of morphological characteristics and increasing attention is being paid towards comprehensive plant genetic recourses.

The characterizations of 15 varieties were done to use as reference material for protection of other varieties under PPVand FR Rules, 2003. Therefore, the database of cauliflower varieties generated may be useful for the selection of suitable varieties to be compared against the candidate varieties developed in India as and when required. This investigation may also be helpful to the researchers with respect to breeding of cauliflower varieties for particular traits. Moreover, farmers can also get benefit with regards to selection of suitable varieties of their interest.

The present investigation was carried out for successive three years during Rabi season of 2010 to 2012 to carry out characterization of already released cauliflower varieties at Research Farm, Indian Institute of Vegetable Research, Varanasi, India. The seed materials for the present investigation comprised of 15 varieties i.e Pusa Paushja, Pusa Sharad, Pusa Himjyoti, Pusa Shukti, Pusa Meghna, Pusa Deepali, Kashi Kunwari, Kashi Agahani, CCS-80, Pant Gobhi-3, Pusa Subhra,

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Pusa Snow Ball-1 PSBK-1, PSB-16 and PSBK-25. The field experiment was laid out in a randomized block design with three replications. Each plot consisted of four rows of 5m length with spacing of 50 cm between rows and 50 cm between plants and all the recommended agronomic practices were followed to raise the good crop. The observations were recorded on 10 plants in each replication at specific stages of crop growth period when the characters under study had full expression. Varieties were evaluated for 28 DUS qualitative and quantitative characters listed in National DUS test guidelines descriptors for cauliflower *viz*; seedling: anthocyanin colouration of hypocotyl, outer stem (stalk): length (up to insertion of first leaf) , leaf attributes *viz*; leaf: attitude, length, width, shape, lobe, colour, waxiness, torsion of tip profile of upper side of blade, puckering , crimping near main vein and degree of undulation of margin, curd characteristics *viz*; curd: initiation (days to 50% of the plants with curd initiation from sowing of seed), covering by inner leaves, polar diameter, equatorial diameter, shape in longitudinal section, doming, colour, knobbing, texture, compactness, anthocyanin colouration at maturity, maturity group, flower: colour, and stalk length (Anonymous, 2009).

The characterization of cauliflower varieties is presented in table-1 and frequency distribution of each descriptor of released varieties along with example varieties is depicted in table-2.

The varieties were classified on the basis of seedling anthocyanin pigment of hypocotyls into two groups i.e. absent and present. Most of the varieties exhibited absent while, snowball group (Pusa Snowball-1, PSBK-1, PSB-16) expressed seedling anthocyanin colouration. Based on outer stem (stalk) length, cauliflower varieties were grouped as short, medium and long. Two varieties namely Pusa Sharad and Pusa Snow Ball-1 expressed medium and 13 varieties showed long stalk length while, none of the varieties depicted short outer stem (stalk) length. As per leaf attitude, three varieties i.e. Pusa Himjyoti, Pusa Snow Ball-1 and PSBK-25 expressed erect while, Pusa Meghna and Kashi Aghrahani showed horizontal and rest 10 varieties showed semi erect type. Leaf length of cauliflower varieties classified into three categories *viz*; short, medium and long. However, in the present study, only four varieties *i.e.* Pusa Meghna, Pusa Shukti, PSB-16 and Pusa Sharad had short leaf length, ten varieties showed medium leaf length and other varieties showed long leaf length. Considerable variation was also observed for leaf shape, variety like Pusa Meghna was of narrow elliptic leaf shape while, Pusa Paushja, Pusa Sharad, PSB-1, Pusa Himjyoti, Pant Gobhi-3, PSBK-25, Pusa Deepali and Kashi Agahani were of elliptic type and rest varieties *i.e.* Pusa Snow Ball K-1, Pusa Shukti, CCH-80, Pusa Subhra, PSB-16 and Kashi Kunwari had broad elliptic type leaf shape. On the basis of leaf lobe, the varieties were classified in two group *viz*; absent and present. Most of the varieties expressed leaf lobe except Pusa Paushja, Pusa Sharad, Pusa Himjyoti, Pusa Snow Ball-1 and Pusa Shukti.

The varieties were also classified on the basis of leaf colour in three groups *i.e.* light green, dark green and bluish. Most of the varieties were found to have light and dark green leaves except Pusa Paushja which showed as bluish leaf colour. Leaf waxiness was another trait with good variability. On the basis of this character, the varieties were categorized into four groups *i.e.* absent, light, medium and strong. Waxiness was absent in Pusa Meghna while, Pusa Paushja, CCS-80, Pant Gobhi-3, Pusa Subhra and PSB-16 exhibited medium waxiness. Varieties like Pusa Himjyoti, Pusa Snow Ball-1, Pusa Shukti, PSBK-1 and Pusa Meghna expressed light whereas; Pusa Paushja and PSBK-25 showed strong leaf waxiness. The trait leaf torsion of tip was categorized into four categories namely, absent, weak, medium and strong. Five varieties *i.e.* Pusa Snow Ball-1, Pusa Shukti, Pusa Meghna, Pusa Deepali and Kashi Kunwari depicted absent, while, 5 varieties expressed weak and rest varieties showed medium torsion of leaf tip. None of the varieties showed strong torsion of leaf tip. On the basis of leaf profile of upper side of blade character, the varieties were classified into three groups *i.e.* concave, flat and convex. Varieties like, Pusa Snow Ball-1, Pusa Shukti, Pusa Deepali, Kashi Kunwari and Kashi Agahani were of flat type leaf while, Pusa Paushja, Pusa Sharad, Pusa Himjyoti, CCS-80, Pant Gobhi-3 and PSBK-1 showed convex and rest varieties showed concave type leaf. Among the varieties, weak leaf puckering was absent in Pusa Himjyoti, Pusa Snow Ball-1, Pusa Meghna and Kashi Kunwari whereas, Pusa Paushja, Pusa Sharad, CCS-80, Pant Gobhi-3, Pusa Subhra, PSBK-1 and Kashi Agahani expressed medium and Pusa Shukti, PSBK-25 and Pusa Deepali showed strong leaf puckering characters. In the favour of leaf crimping near main vine, the varieties have been grouped into four categories, *viz;* seven varieties as weak, five as medium and 3 as strong whereas, none of the varieties had no leaf crimping near main vine. On the basis of curd initiation, cauliflower varieties can be classified into three categories *viz;* early (<75 days), medium (75-100 days), late (>100 days). There are three varieties with early curd initiation *e.g.* Pusa Meghna, Pusa Deepali and Kashi Kunwari while, 9 varieties *i.e.* Pusa Himjyoti, Kashi Agahani, Pusa Paushja, Pusa Sharad, Pusa Shukti, CCS-80, Pant Gobhi-3 and Pusa Subhra were in medium group and Pusa Snowball -1, PSBK-1, PSB-16 and PSBK-25 exhibited late curd initiations. Curd covering by inner leaves was another trait with good variability. On the basis of this character, the varieties were grouped into three types *viz;* not covered, partly covered, and covered. Four varieties (Kashi Kunwari, Pusa Himjyoti, Pusa Deepali and Pusa Meghna) were not covered, whereas, Kashi Agahani Pusa Paushja, Pusa Sharad, Pusa Shukti, CCS-80, Pant Gobhi-3 and Pusa Subhra exhibited partly covered and four (PSBK-1, PSB-16, PSBK-25 and Pusa Snow Ball-1) varieties covered by inner leaves. Based on polar diameter of curd, varieties were classified into 3 groups *viz;* small (<15 cm), medium (15-20cm) and large (>20cm). Proportionate distributions of varieties in small and medium group were observed. Only two varieties Pusa Shukti and PSBK-1 were in large group. Equatorial diameter of curd cauliflower varieties were grouped as small, medium and large, accordingly, 4 varieties as small, 9 as medium and 2

were observed as large equatorial diameter during experimentation. Dubey et al., (2003) reported high variability in curd size of cauliflower. Shape in longitudinal section of curd is another character with sufficient variability in cauliflower varieties. Circular shape was observed in Pusa Himjyoti, broad elliptic was observed in 11 varieties *viz*; Kashi Kunwari, Kashi Agahani, Pusa Paushja, Pusa Sharad, Pusa Shukti, CCS-80, Pant Gobhi-3, Pusa Subhra, PSBK-1, PSB-16 and PSBK-25, while, Pusa Snow Ball-1, Pusa Megna and Pusa Deepali were depicted as narrow elliptic. The character of curd doming was categorized as weak, medium and strong. In this respect variety Pusa Meghna showed weak and variety Pusa Paushja expressed have strong curd doming whereas, rest of other varieties were depicted as medium curd doming. Curd colour varied from white to creamy white in the varieties studied. Pusa Paushja, Pusa Sharad, PSBK-1 and PSBK-25 exhibited white colour whereas, rest of the varieties showed creamy white colour. Curd knobbing is important character categorized into three categories i.e. fine, medium and course. Most of the varieties showed medium knobbing except PSBK-1 which expressed fine curd knobbing. Among the 15 cauliflower varieties studied considerable variation was observed for all the characters except curd texture (Table-1) where all the varieties exhibited fine curd texture. On the basis of curd compactness, the varieties were classified into three groups *viz*; loose, medium and compact, seven varieties were found as medium type and 8 varieties of compact type. Similar findings were reported by Singh et al., (2005). Kumar et al. (2010) found significant differences among the genotypes in Indian cauliflower suggesting sufficient variability for yield and quality characters.

The trait anthocyanin colouration of curd at maturity was categorized into two categories namely, absent and present. Two varieties depicted as present and rest of 13 varieties showed absent in anthocyanin curd colouration. Curd maturity group is most important character which classified as early, mid early, mid late and late. Only one variety Pusa Deepali was found early, whereas, Kashi Kunwari, Pusa Meghna was expressed as mid early group, Pusa Paushja, Pusa Sharad, Pusa Himjyoti, Pusa Shukti, CCS-80, Pant Gobhi-3 and Pusa Subhra were exhibited as mid late group and all the snow ball group varieties were depicted as late group for maturity. Late group varieties *viz*; Pusa Snow Ball-1, PSB-16, PSBK-1, PSB-16 and PSBK-25 could not flower due to lack of chilling requirement in northern plane in India. Flower colour grouped as white, creamy white and yellow. Among the 11 flowered varieties, three varieties namely Pusa Himjyoti, Pusa Deepali and Pusa Meghna expressed creamy white, and rest of the other varieties showed yellow flower colour while, none of the variety was depicted as white colour. Similar grouping trends were reported by Singh et al., (2012) in cabbage. The trait flower stalk length was categorized into three categories namely short, medium and long. Out of these, two varieties was depicted as short stalk length and 9 varieties showed medium stalk length.

It is concluded that the DUS descriptors can be effectively used for identification and grouping of varieties and satisfying the DUS criteria for these morphological descriptors could be registered under PPV and FR Act. Pre-breeding

or genetic enhancement needs emphasis for transfer or introgression of genes and gene combinations from unadapted source into more usable breeding material. There are indications that novel and useful traits can be successfully combined from related varieties. Further, these varieties can be used in varietal improvement programme depending upon the desired characteristics. However, registration of candidate varieties only considered under DUS testing as per the legal frame work and till now, morphological characterizations are being considered as per PPV and FR Act for notified vegetable crops.

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**Table 1: Description of morphological DUS descriptors for fifteen cauliflower varieties/ cultivars.**

S. No.	Descriptor	Varieties/Cultivars															Remarks
		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	
1	Seedling: anthocyanin colouration of hypocotyl	1	1	1	9	1	1	1	1	9	1	9	1	1	1	1	1:absent, 9:present
2	Outer stem (stalk): length (up to insertion of first leaf)	7	5	7	5	7	7	7	7	7	7	7	7	7	7	7	3:short, 5: medium,7: long
3	Leaf : attitude	3	3	1	1	3	3	3	3	3	3	1	5	3	3	5	1:erect, 3: semi erect,5: horizontal
4	Leaf: length	5	3	5	3	5	5	5	5	7	3	5	3	5	5	5	3:short, 5: medium,7: long
5	Leaf: width	5	5	3	3	7	5	5	5	5	3	5	5	5	3	5	3:narrow, 5:medium, 7:broad
6	Leaf: shape	5	5	5	5	7	7	5	7	7	7	5	3	5	7	5	3:narrow elliptic, 5:elliptic,7:broad elliptic
7	Leaf: lobe	9	9	9	9	9	9	1	1	1	1	1	1	1	1	1	1:absent, 9:present
8	Leaf: colour	2	2	1	1	2	2	2	1	2	2	2	1	1	1	2	1:light green, 2:dark green,3: bluish
9	Leaf: waxiness	7	3	2	2	2	3	3	3	2	3	7	1	2	3	3	1:absent, 2:light,3:medium,7:strong
10	Leaf: torsion of tip	3	5	3	1	1	5	3	3	5	5	5	1	1	1	3	1:absent, 3:weak,5:medium,7:strong
11	Leaf: profile of upper side of blade	3	3	3	2	2	3	3	1	3	1	1	1	2	2	2	1:concave, 2:flat, 3:convex
12	Leaf: puckering	5	5	3	3	7	5	5	5	5	1	7	3	7	3	5	1:absent, 3:weak,5:medium,7:strong
13	Leaf: crimping near main vein	5	5	3	3	7	5	5	3	7	3	7	3	5	3	3	1:absent, 3:weak,5:medium,7:strong
14	Leaf: degree of undulation of margin	3	5	3	1	7	5	5	5	5	3	7	5	3	3	3	1:absent, 3:weak,5:medium,7:strong
15	Curd initiation (days to 50% of the plants with curd initiation from sowing of seed)	5	5	5	7	5	5	5	5	7	7	7	3	3	3	5	3:early, 5:medium, 7:late

S. No.	Descriptor	Varieties/Cultivars															Remarks
		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	
16	Curd: covering by inner leaves	5	5	3	7	5	5	5	5	7	7	7	3	3	3	5	3: not covered, 5: partly covered, 7: covered
17	Curd: polar diameter	5	5	3	3	7	5	5	5	7	5	5	3	3	5	5	3: small, 5: medium, 7: large
18	Curd: equatorial diameter	5	5	3	3	7	5	5	5	7	5	5	3	3	5	5	3: small, 5: medium, 7: large
19	Curd: shape in longitudinal section	3	3	1	5	3	3	3	3	3	3	3	5	5	3	3	1: circular, 3: broad elliptic, 5: narrow elliptic
20	Curd: doming	7	5	5	5	5	5	5	5	5	5	5	3	5	5	5	3: weak, 5: medium, 7: strong
21	Curd: colour	1	1	2	2	2	2	2	2	1	2	1	2	2	2	2	1: white, 2: creamy white, 3: orange
22	Curd: knobbing	5	5	5	5	5	5	5	5	3	5	5	5	5	5	5	3: fine, 5: medium, 7: coarse
23	Curd: texture	3	3	3	3	3	3	3	3	3	3	3	3	3	3	3	3: fine, 7: coarse
24	Curd: compactness	7	7	5	7	5	5	5	7	7	5	7	5	5	7	7	3: loose, 5: medium, 7: compact
25	Curd: anthocyanin colouration at maturity	1	1	1	9	1	1	1	1	9	1	1	1	1	1	1	1: absent, 9: present
26	Curd: maturity group	7	7	7	9	7	7	7	7	9	7	9	5	3	5	7	3: early, 5: mid early, 7: mid late, 9: late
27	Flower: colour	3	3	2	-	3	3	3	3	-	-	-	2	2	3	3	1: white, 2: creamy white, 3: yellow
28	Flower: stalk length	5	3	5	-	5	5	5	5	-	-	-	5	3	5	5	3: short, 5: medium, 7: long

**Varieties/cultivars:** 1: Pusa Paushja, 2: Pusa Sharad, 3: Pusa Himjyoti, 4: Pusa Snow Ball-1, 5: Pusa Shukti, 6: CCS-80, 7: Pant Gobhi-3, 8: Pusa Subhra, 9: PSBK-1, 10: PSB-16, 11: PSBK-25, 12: Pusa Meghna, 13: Pusa Deepali, 14: Kashi Kunwari, 15: Kashi Agahani

Note: (-) is Pusa Snow Ball-1, PSBK-1, PSB-16 and PSBK-25 are snowball group. Temperate region is suitable for this group flowering.

**Table 2: Frequency distribution and example varieties of some important attributes of 15 varieties/ cultivars of cauliflower**

Plant descriptors	Range in expression	No. of varieties	Example varieties
Seedling: anthocyanin colouration of hypocotyl	absent	12	Pusa Deepali, Kashi Agahani
	present	3	PSBK-1, PSBK-25, Pusa Snowball-1
Leaf : attitude	erect	3	PSBK-25, Pusa Snowball-1
	semi-erect	10	Pusa Paushja, Pusa Sharad, Pusa Himjyoti
	horizontal	2	Pusa Meghna
Leaf: length	short (<35 cm)	4	Pusa Meghna, Pusa Shukti, PSB-16
	medium (35-50 cm)	10	Pusa Sharad, Kashi Agahani, CCS-80
	large (>50 cm)	1	PSBK-1
Leaf: width	narrow (<15 cm)	4	Pusa Himjyoti, Kashi Kunwari
	medium ( 15-25 cm)	10	Kashi Agahani, Pusa Deepali, , PSBK-25
	broad (>25 cm)	1	Pusa Shukti
Leaf: shape	narrow elliptic	1	Pusa Meghna
	elliptic	8	Pusa Paushja, Pusa Sharad, PSB-1, Pusa Himjyoti
	broad elliptic	6	Pusa Snow Ball-1, Pusa Shukti
Leaf: profile of upper side of blade	concave	4	Pusa Meghna, PSB-16
	flat	5	Pusa Shukti, Kashi Kunwari, Pusa Deepali
	convex	6	Pusa Paushja, Pusa Sharad, Pusa Himjyoti
Leaf: puckering	absent	1	PSB-16
	weak	4	Kashi Kunwari, Pusa Meghna, Pusa Himjyoti
Leaf: crimping near main vein	medium	7	Kashi Agahani Pusa Paushja, Pusa Sharad,
	strong	3	Pusa Deepali, Pusa Shukti
	absent	0	Nil
	weak	7	Kashi Kunwari, Pusa Meghna
Curd initiation (days to 50% of the plants with curd initiation from sowing of seed)	medium	5	Pusa Paushja, Pusa Sharad, Pusa Deepali
	strong	3	Pusa Shukti, PSBK-25, PSBK-1
	early (<75 days)	2	Pusa Meghna, Pusa Deepali
	medium (75-100)	9	Kashi Kunwari, , Pusa Himjyoti, Kashi Agahani Pusa Paushja, Pusa Sharad, Pusa Shukti
	late (>100)	4	Pusa Snowball -1, PSBK-1, PSB-16, PSBK-25
Curd: covering by inner	not covered	4	Kashi Kunwari, Pusa Himjyoti, Pusa

Plant descriptors	Range in expression	No. of varieties	Example varieties
leaves	partly covered	7	Deepali, Pusa Meghna Kashi Agahani Pusa Paushja, Pusa Sharad, Pusa Shukti
	covered	4	PSBK-1, PSB-16, PSBK-25, Pusa Snow Ball-1
Curd: polar diameter	small (<15 cm)	4	Pusa Himjyoti, Pusa Deepali, Pusa Meghna
	medium (15-20 cm)	9	Kashi Agahani Pusa Paushja, Pusa Sharad
	large (>20cm)	2	Pusa Shukti, PSBK-1
Curd: equatorial diameter	small (<15 cm)	4	Pusa Himjyoti, Pusa Deepali, Pusa Meghna
	medium (15-20 cm)	9	Kashi Agahani Pusa Paushja, Pusa Sharad
	large (>20cm)	2	Pusa Shukti, PSBK-1
Curd: shape in longitudinal section	circular	1	Pusa Himjyoti
	broad elliptic	11	Kashi Kunwari, Kashi Agahani Pusa Paushja, Pusa Sharad, PSBK-25
	narrow elliptic	3	Pusa Deepali, Pusa Meghna PSB-1
Curd: doming	weak	1	Pusa Meghna
	medium	13	Kashi Kunwari, Kashi Agahani, PSBK-25, Pusa Sharad, Pusa Himjyoti
	strong	1	Pusa Paushja
Curd: compactness	loose	0	Nil
	medium	7	Pusa Himjyoti, Pusa Deepali, Pusa Meghna,
	compact	8	PSBK-25, Pusa Sharad, Pusa Paushja, Pusa Shukti
Curd: maturity group	early	1	Pusa Deepali
	mid-early	2	Kashi Kunwari, Pusa Meghna
	mid-late	9	Pusa Paushja, Pusa Shukti ,Pusa Sharad, Kashi Agahani
	late	3	PSBK-25, PSB-1, Pusa Snowball -1
Flower: colour	white	0	Nil
	creamy white	3	Pusa Himjyoti, Pusa Deepali, Pusa Meghna
	yellow	8	Pusa Sharad, Kashi Agahani, Pusa Himjyoti, Kashi Kunwari

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#### **Book/Bulletin/Reports/Series**

- Bhuiyan, S.I. 1982. Irrigation system management research and selected methodological issues. *IRRI research paper series no 81*. Los Banos, Manila.
- De Datta, S. K. 1981. Principles and practices of rice production. Los Banos, Manila.
- International Rice Research Institute. 2000. International rice trade: a review of 1999 and prospects for 2000. *International Rice Commission Newsletter*, IRRI, Manila.
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- Westerman, R.L. (ed.) 1990. Soil testing and plant analysis. 3rd ed. SSSA Book Ser. 3. SSSA, Madison, WI.

#### **Chapter in a Book**

- David, H. and Easwaramoorthy. 1988. Physical resistance mechanisms in insect plant interactions. p. 45-70. In T.N. Ananthkrishnan and A. Rahman (ed.), *Dynamics of insect plant interactions: Recent advances and future trends*. Oxford and IBH Publication, New Delhi.
- Johnson, D.W. and D.E. Todd. 1998. Effects of harvesting intensity on forest productivity and soil carbon storage. p. 351-363. In R. Lal et al. (ed.) *Management of carbon sequestration in soils*. Advances in Soil Science. CRC Press, Boca Raton, FL.

### **Conference/Symposium/ Proceedings**

Joshi, B.K. 2004. Crossing frequency and ancestors used in developing Nepalese mid and high hill rice cultivars: Possible criteria for yield improvement and rice genes conservation. p. 502-523. In Proc. *National Conference on Science and Technology*, 4th, Vol. 1. 23-26 Mar., 2004. NAST, Kathmandu, Nepal.

Ramanujam, S. (ed.) 1979. *Proceedings of International Wheat Genet Symposia*, 5th, New Delhi, India. 23–28 Feb. 1978. Indian Soc. Genet. Plant Breeding, Indian Agric. Res. Inst., New Delhi.

### **Dissertation**

Singh, A.A. 2005. Weed management approaches and modeling crop weed interaction in soybean. M. Sc. (Ag.) thesis. Tamil Nadu Agricultural Univ., Coimbatore.

### **Software and Software Documentation**

Minitab. 1998. MINITAB 12. Minitab, State College, PA.

### **Online publication**

Venugopal, D. 2000. Nilgiri tea in crisis: Causes consequences and possible solutions. Retrieved October 11, 2000 from <http://www.badaga.org>.

### **Online journal article**

Doerge, T.A. 2002. Variable-rate nitrogen management creates opportunities and challenges for corn producers. *Crop Manage.* doi:10.1094/cm-2002-0905-01-RS.

### **Tables**

1. Each table must be typed on a separate sheet (not to be included in the text) and numbered consecutively in the same order as they mentioned in text.
2. The title should fully describe the contents of the table and explain any symbol or abbreviation used in it as a footnote, using asterisks or small letters viz. a, b, etc.
3. Tables should be self-explanatory, not very large (< 10 columns in portrait and <14 columns in landscape formats respectively) and may cover space up to 20-25% of the text.
4. Maximum size of table acceptable is that can be conveniently composed within one full printed page of the journal. The large sized tables should be suitably split into two or more small tables.
5. Standard abbreviations of units of different parameters should be added between parentheses.

6. The data in the tables should be corrected to minimum place of decimal so as to make it more meaningful.
7. Vertical lines should not be used to separate columns. Similarly, horizontal lines should be used only where these are necessary, not in the body of the article.
8. All tables should be tagged with the main body of the text i.e. after references.

### **Figures**

1. Figures may be given in place of tables where a large number of values are presented that can be interpreted through figures. In no case the same data should be presented in both tables and figures.
2. Originals of the figures should be no larger than twice the final size, of good quality and printed clearly in black on plain white paper or in color. The figures may be sized to fit within the columns of the journal (8 cm width for single column or 17 cm for columns i.e. full page).
3. Lines should be bold enough to allow the figure to be reduced to either single or double column width in the journal.
4. Black and white photographs are also accepted if these are necessary to improve the presentation and quality of the article.

### **Some useful hints**

1. All scientific or technical names as well as all data and facts must be rechecked carefully before submitting the manuscript.
2. Dates and years may be mentioned as 28 May 2007, 28 May to 7 June, and 28-30 May instead of May 28, 2007, 28 May-7 June, and 28 to 30 May, respectively.
3. Avoid numerals and abbreviations at the beginning of a sentence; spell out or change the word order if necessary.
4. A comma may be used for data in thousands or more such as 10,000 or 2,30,000 etc. Alternatively, these data can also be presented as 10.0 or 230.0 if a common expression such as ' $\times 10^3$ ' is used in tables or figures. Avoid expressing data in 'lakhs', instead use 'thousand' or 'million'.
5. Only standard abbreviations should be used and these should invariably be explained at first mention. Avoid use of self-made abbreviations such as Rhizo., Azo., buta, isop. etc. for *Rhizobium*, *Azotobacter*, butachlor, isoproturon, respectively.
6. For names of plant protection chemicals, the first letter of the name need not be capitalized for scientific names but should be capitalized for trade names. All the names should be checked very carefully.

7. Use of unnecessary abbreviations and treatment symbols such as T1, T2 etc. under Materials and Methods or tables without actually using these under Results and Discussion should be avoided.
8. All weights and measurements must be in SI or metric units. Use  $\text{kg ha}^{-1}$ , or  $\text{t ha}^{-1}$  but not  $\text{q ha}^{-1}$ . Do not follow the style  $\text{g/ha}$ ,  $\text{mg/kg}$ ,  $\text{mg/l}$ ,  $\text{mg/g}$ ,  $\text{ml/l}$  or  $\text{g per ha}$ , etc
9. Use % after numbers, not per-cent, e.g. 7%. In a series or range of measurements, mention the units only at end, e.g. use 30, 100, 170 and 300C; 20 or 30% more instead of 30C, 100C, 170C and 300 °C; 20% or 30% more.
10. Numeral should be used whenever it is followed by a unit measure or its abbreviations e.g. 1 g, 3 m, 5 h, 6 months etc. Otherwise, words should be used for numbers one to nine and numerals for larger ones except in a series of numbers when numerals should be used for all in the series.
11. For the composition of fertilizers, manures, crops or soil, the elemental forms (K, P, Mg etc.) should be used and not the oxides.
12. Statistical analysis of data in the standard experimental design should be sound and complete in itself with both  $\pm\text{SE}$  (Standard Error of means) and CD (Critical Difference) or LSD (Least Significant Difference) ( $P=0.05$ ) values given for comparison of treatment means in tables and figures.
13. Use multiplication sign ( $\times$ ) not a alphabet x for indicating multiplication, crossing, etc

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