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## **Biochemical response of mungbean (*Vignaradiata L.*) under the influence of reduced dose of chemical fertilizer and different time and method of application of biofertilizer**

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Fundamental differences between organic and conventional production systems, particularly in soil fertility management, may affect the nutritive composition of plants, including secondary plant metabolites. The availability of inorganic nitrogen in particular has the potential to influence the synthesis of secondary plant metabolites, proteins, and soluble solids. This study was conducted to evaluate the on-farm assessment of biochemical changes in leaves of mungbean by employing different varieties and different treatments such as different reduced doses of N:P:K chemical fertilizer, different method and time of application of biofertilizer. Untreated seeds were used as control. PDM-11 variety showed higher accumulation of chlorophyll, sugar, protein in leaves and PUSHA-9531 reported higher accumulation of amino acid content in leaves. Significant level of variation was observed in the level of plant metabolites and macromolecules in seeds among the six studied mungbean varieties.

**Key words:** Biochemistry, Mungbean, Phenol, Protein

### **Introduction**

Mung bean (*Vignaradiata L.*) is a leguminous pulse crop which used is as a vegetable protein source, animal fodder and green manure, it contains isoflavonoids having estrogen and antioxidant activities that used in prevention of many diseases such as cancer, and it also exhibits antimicrobial and insecticidal activities (Aggrawalet *al.*, 1997). Imbalanced and continuous use of chemical fertilizers in cropping systems is leading to imbalance of nutrients in soil which have an adverse effect on soil health and also biochemical constituents of plant. The productivity of any crops depends on the process of photosynthesis, which in turn depends on the chlorophyll content of leaves in plants (Shah, 1959).

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Fundamental differences between organic and conventional production systems, particularly in soil fertility management, may affect the nutritive composition of plants, including secondary plant metabolites. The availability of inorganic nitrogen in particular has the potential to influence the synthesis of secondary plant metabolites, proteins, and soluble solids. Stamp (2003) reported that increased crop growth and development rates and greater biomass accumulation in well-fertilized crops would also correlate with decreased allocation of resources toward the production of starch, cellulose, and non-nitrogen-containing secondary metabolites. Phenolic compounds are an important group of bioactive compounds that are widely distributed in fruits, herbs and vegetables and their antioxidant activity has been extensively reported. Nitrogen is the nutrient that has the largest effect on the physiology and yield of cotton plants. It is an essential nutrient for production of plant dry matter, as well as many energy-rich compounds that regulate photosynthesis and plant production (Wu *et al.*, 1998). Nitrogen is also vital in protein structure and in basic N compounds such as purines, pyrimidines, porphyrins and coenzymes. Purines and pyrimidines are constituents of the nucleic acids RNA and DNA, which are essential for protein synthesis. Porphyrin is found in metabolically important compounds such as chlorophylls and the cytochrome enzymes, which are necessary for photosynthesis and respiration (Devlin, 1972). Form of nitrogen fertilizer could have an important influence on the accumulation of chlorophyll in plants. Lettuce plants fertilized with two forms of nitrogen ( $\text{NO}_3 + \text{NH}_4$ ) had significantly more chlorophylls *a* and *b* than that fertilized with nitrate nitrogen; and plants fertilized with ammonium nitrogen had the highest concentration of chlorophylls (Michalek and Rukasz, 1998).

Nitrogen fertilization stimulates accumulation of chlorophyll. Mihailovic *et al.*, (1997) reported that ammonium ions in nutrient solution increased concentration of chlorophylls *a* and *b* in wheat leaves in comparison with nitrate ions. There is an optimal relationship between plant N content and  $\text{CO}_2$  assimilation (Sinclair and Horie, 1989).

Interest in other plant secondary compounds has increased because of their potential effects on improving human health (Dumus *et al.*, 2003). Phenolic compounds play a role in plant defence mechanism to resist diseases and insects, also acts as antioxidants if consumed in food. Higher levels of phenolic compounds frequently have been reported in organically grown crops than in conventionally grown crops (Carbonaro *et al.*, 2002). Some earlier workers (Michalek and Rukasz, 1998) found that ammonium nitrogen increased accumulation of phenols. Influence of ammonium and nitrate nitrogen applied together on phenol concentration depended on cultivar. Some earlier workers (Rozek *et al.*, 1994) reported that fertilization with reduced form

of nitrogen increased content of sugars in leaves of lettuce in comparison to plants fertilized only with  $\text{NO}_3$ . The same effect was observed in corn salad leaves in the case of second growing period. The aim of the study was to investigate the biochemical changes in leaves and seeds of mungbean plants associated with different reduced dose of chemical fertilizer and supplement by the biofertilizer and different methods of biofertilizer application and subsequent grain yield.

## Materials and methods

Field experiments were conducted during summer season of 2003-04, 2004-05, and 2005-06 at Crop Research and Seed Multiplication Farm of Burdwan University, India (Latitude  $67^{\circ} 50' 51''$  N and longitude  $23^{\circ} 15' 12''$  E). The source of chemical fertilizer includes urea as nitrogen fertilizer, single super phosphate as phosphate and muriate of potash as potassium. Pure culture of *Rhizobium* was used as bacterial biofertilizer and was procured from Exploration India, Research and Development Wing, Burdwan Institute of Management and Computer sciences, Burdwan, West Bengal, India. In 2003-04, five varieties of mungbean [V<sub>1</sub>-NORMAL, V<sub>2</sub>-PDM-11, V<sub>3</sub>-PDM-54, V<sub>4</sub>-PDM-139, V<sub>5</sub>-PUSHA-9531 and V<sub>6</sub>-PANT MUNG] were cultivated under recommended dose of chemical fertilizer (Directorate of Agriculture, Government of India). In 2004-05 the treatment combination includes.

During the summer season (2004) we conducted one field experiment with different reduced dose of nitrogen(N) and phosphate(P) fertilizer along with recommended dose of potassium(K) which includes T<sub>1</sub>-20% less nitrogen + 25% less phosphate + recommended dose of potassium; T<sub>2</sub>-30% less nitrogen + 25% less phosphate + recommended dose of potassium; T<sub>3</sub>-40% less nitrogen + 25% less phosphate + recommended dose of potassium; T<sub>4</sub>-50% less nitrogen + 25% less phosphate + recommended dose of potassium; T<sub>5</sub>- 60% less nitrogen + 25% less phosphate + recommended dose of potassium and T<sub>6</sub>-recommended dose i.e. 20:40:20.

In 2005-06 the treatment combination was T<sub>1</sub>-as basal application  $1.5 \text{ kg ha}^{-1}$  + on 21 DAS at  $0.75 \text{ kg ha}^{-1}$  as spray + on 42 DAS at  $0.75 \text{ kg ha}^{-1}$  as spray; T<sub>2</sub>- basal application  $3 \text{ kg ha}^{-1}$ ; T<sub>3</sub>-as basal application at  $1.5 \text{ kg ha}^{-1}$  + on 21 DAS at  $1.5 \text{ kg ha}^{-1}$  as soil application; T<sub>4</sub>-as basal application at  $1.5 \text{ kg ha}^{-1}$  + on 21 DAS at  $1.5 \text{ kg ha}^{-1}$  as spray; T<sub>5</sub>-as basal at  $0.75 \text{ kg ha}^{-1}$  + on 21 DAS at  $1.5 \text{ kg ha}^{-1}$  as soil application + on 42 DAS at  $0.75 \text{ kg ha}^{-1}$  as soil application. The experiment was carried out in a randomized block design with different treatment in different year with three replications. The experiments were conducted in randomized block design with three replications. Individual plot size (5m x 5m); plant row to row spacing 30 cm and plant to plant 15 cm.

For biochemical estimation in leaves and seeds constituents such as total chlorophyll (Arnon, 1949), total soluble sugar (McCready *et al.*, 1952), total amino acids (Moore and Stein, 1948), total protein (Lowry *et al.*, 1950) was estimated from physiologically active leaves and other parameters like RNA, DNA (Choudhuri and Chatterjee, 1970). Ascorbic acid (along with total soluble sugar (McCready *et al.*, 1950), total amino acid (Moore and Stein, 1948), and total protein (Lowry *et al.*, 1950) was estimated from seeds.

After harvesting the biomass (as dried) of the whole plant were estimated. Then the grain yields were estimated in square meter and again it is calculated in hector.

Physiologically active leaf (3<sup>rd</sup> and 4<sup>th</sup> from the top) samples from ten different plants of each plots were plucked randomly for determination of different biochemical parameters. Composite samples of leaf were crushed separately in a mortar with 10 ml of distilled water, poured in centrifuge tube and centrifuged at 10000 rpm for ten minutes. The supernatant was used for soluble sugar determination following the method of McCready *et al.*, (1950); for estimation of free amino acid in leaf samples Moore and Stain (1948); for chlorophyll content (mg.g<sup>-1</sup>.f.w) method of Arnon (1949); for protein content (mg.100g<sup>-1</sup>) method of Lowry *et al.*, (1951).

Seeds of were imbibed in distilled water under room temperature for overnight. After that seed coat was removed and the seeds were crashed in a mortar and pestle along with distilled water and centrifuged for 10000 rpm for 10 minutes. The supernatant was used for soluble sugar determination following the method of McCready *et al.*, (1950). For determination of free amino acid in seeds Moore and stain (1948); for protein (Lowry *et al.*, 1951); for Ascorbic acid (Mukherjee and Choudhury, 1984); for nucleic acid (Choudhuri and Chatterjee, 1970) standard methods were followed.

All the experimental results of the subsequent three years were analysed by one way analysis of variance (ANOVA). All the statistical analysis were carried out with the programme SPSS 11.0 for windows. All values were expressed as mean values. The statistical significance of difference between the different treatments were established using Duncan's Multiple Range Test (DMRT).

## Results

This study pertained to assess the biochemical changes in leaves and seed of mungbean under different treatments of reduced dose of chemical fertilizer and different methods and time of application of biofertilizer and was evaluated in terms of chlorophyll, protein, amino acid, sugar content in leaves and accumulation of chlorophyll, protein, amino acid, sugar, ascorbic acid, phenol,

DNA and RNA for enhanced crop growth, metabolism and optimum protection against pathogens and pest.

In 2003, there was significant level of variation in the level of total chlorophyll in the leaves of six mungbean varieties. Minimum and maximum total chlorophyll levels were observed in V<sub>5</sub> and V<sub>3</sub> respectively (Table-1). In 2004, 30% reduction of the recommended dose of nitrogen fertilizer (T<sub>2</sub>) was stimulatory for chlorophyll content in leaves when compared to application of recommended dose of chemical fertilizer (T<sub>6</sub>). During 2005, soil application of biofertilizer was found to promote the total chlorophyll content in leaves of mungbean plants (T<sub>3</sub> and T<sub>6</sub>) over spray application (Table-2).

In 2003, there was significant level of variation in the level of total chlorophyll in the leaves of six mungbean varieties. Minimum and maximum total chlorophyll levels were observed in V<sub>6</sub> and V<sub>3</sub> respectively (Table-1). In 2004, 30% reduction of the recommended dose of nitrogen fertilizer (T<sub>2</sub>) significantly promoted the accumulation of sugar content in leaves when compared to application of recommended dose of chemical fertilizer (T<sub>6</sub>). During 2005, soil application of biofertilizer significantly promoted the accumulation of soluble sugar content in leaves of mungbean plants (T<sub>3</sub> and T<sub>6</sub>) over spray application (Table-2).

Significant levels of variation in the level of amino acid in the leaves of six mungbean varieties were observed during the present investigation of 2003. Highest and lowest accumulation of amino acid were found to be in V<sub>3</sub> and V<sub>6</sub> respectively (Table-1). In 2004, 30% reduction of the recommended dose of nitrogen fertilizer (T<sub>2</sub>) was found to be stimulatory towards biosynthesis of amino acid in leaves when compared to application of recommended dose of chemical fertilizer (T<sub>6</sub>). Soil application of biofertilizer at different phases of crop growth were found to be stimulatory towards biosynthesis of amino acid and its accumulation in leaves of mungbean plants when compared to spray application (Table-2).

Protein content in leaves varied significantly among the seven varieties of mungbean. Maximum and minimum accumulations were recorded by varieties V<sub>6</sub> and V<sub>3</sub> respectively (Table-1). In 2004, 30% reduction of the recommended dose of nitrogen fertilizer (T<sub>2</sub>) was found to be stimulatory towards biosynthesis of protein in leaves when compared to application of recommended dose of chemical fertilizer (T<sub>6</sub>). Soil application of biofertilizer at different phases of crop growth were found to be stimulatory towards biosynthesis of protein and its accumulation in leaves of mungbean plants when compared to spray application (Table-2).

The sugar content in seeds of six studied mungbean varieties showed significant variation among them. Maximum and minimum accumulation of

sugar was found to be in V<sub>3</sub> and V<sub>6</sub> variety respectively (Table-3). In 2004, 30% reduction of the nitrogen fertilizer dose was found to be stimulatory towards accumulation of sugar in freshly harvested seeds when compared to application of recommended dose of chemical fertilizer. Soil application of biofertilizer significantly promoted accumulation of sugar in seeds of freshly harvested mungbean (Table-4).

The amino acid content in freshly harvested seed varied between 15.303 g 100g<sup>-1</sup> in V<sub>3</sub>( PDM-54) and 31.80 g 100g<sup>-1</sup> in V<sub>6</sub> (PANT MUNG) during 2003-04.(Table-3). The amino acid content in seeds decreased proportionately as the rate of nitrogen fertilizer reduction increased gradually. Soil application of biofertilizer at different rate and time was found to be stimulatory towards amino acid biosynthesis leading to higher accumulation of amino acid in seeds over spray application (Table-4).

Significant levels of variation in the level of protein in the seeds of six mungbean varieties were observed during the present investigation of 2003. Highest and lowest accumulation of amino acid were found to be in V<sub>3</sub> and V<sub>6</sub> respectively (Table-3). 30% reduction of nitrogen fertilizer dose was found to be stimulatory over application of recommended dose of nitrogen fertilizer towards protein synthesis in mungbean seeds. Soil application of biofertilizer significantly promoted better protein synthesis in mungbean seeds leading to higher protein content in seeds (Table- 4).

Ascorbic acid content in seeds showed significant level of variation among the six studied mungbean varieties in 2003. PDM-54 seeds recorded the highest accumulation of ascorbic acid.(Table-3). Treatment T<sub>2</sub> recorded highest accumulation of ascorbic acid in mungbean seeds during 2004.Spray application of biofertilizer at different rate and time of crop growth was found to be inhibitive towards accumulation of ascorbic acid in mungbean seeds (Table-4).

Phenol content in seeds showed significant level of variation among the six studied mungbean varieties in 2003. PDM-54 seeds recorded the highest accumulation of ascorbic acid (Table-3). Treatment T<sub>2</sub> recorded highest accumulation of phenolics in mungbean seeds during 2004.Spray application of biofertilizer at different rate and time of crop growth was found to be inhibitive towards accumulation of phenolics in mungbean seeds (Table-4).

Nucleic acid (DNA and RNA) content in mungbean seeds varied significantly among the six varieties. The level of ribonucleic acid (RNA) content in seeds was higher than deoxy-ribonucleic acid(DNA) among the six studied varieties (Table-3). Nucleic acid content in seeds varied significantly along with different nitrogen fertilizer reduction dose in 2004 as well as with

the method, rate and time of application of biofertilizer. The RNA content in seeds was higher than DNA for both the years of 2004 and 2005 (Table-4).

## Discussion

The significant variation in the level of total chlorophyll content in physiologically active leaves of six mungbean varieties may be due to variable rate of biosynthesis of chlorophyll and photosynthesis depending upon the genetic potential of the mungbean varieties. Total chlorophyll increased in T<sub>2</sub> treatment than the recommended dose which might be due to significant influence of microorganisms as biofertilizer of producing growth promoting substances resulting in more efficient absorption of nutrients, which main components of photosynthetic pigments and consequently the chlorophyll content was increased (Gomaa and Abou-Aly, 2001). Similar results were obtained by Hassan (2009).

The soil application of bacterial biofertilizer at different phases of crop growth may have promoted increased chlorophyll content in leaves due to the supply of higher amount nitrogen to the growing tissue and organs supplied by N<sub>2</sub> fixing *Rhizobium*. (Chandrasekhar *et al.*, 2005) over spray application.

Total soluble sugar content in leaves showed considerable variation among the different mungbean varieties which may be attributed towards the variable rate of photosynthesis, leading to production of variable amount of photosynthate among the different varieties. The sugar content in leaves increased upto 30% of reduced dose of nitrogenous chemical fertilizer and then decreased further with gradual increment of reduction of nitrogen fertilizer dose which may be attributed towards higher rate of translocation of sugar transported from leaves to flowering parts and its subsequent utilization for the development of seeds (Chandrasekar BR, 2005) under reduced supply of nitrogen to crop plants. This therefore indicates the role of nitrogen towards biosynthesis and accumulation of sugar in crop plants. Similar findings were reported by Setua *et al.* (2005) in mulberry leaves. The results also reveals that dual application of biofertilizers and chemical fertilizer has pronounced influence on biosynthesis of carbohydrates in the leaves in terms of soil application over spray at different phases of crop growth (Rao *et al.*, 2007).

Significant variation in the level of amino acid content in leaves among the different varieties may be attributed towards the differential rate of inorganic nitrogen assimilation towards nitrogen transporting amino acid which are used to transfer nitrogen from source organs to sink tissues and to build up reserves during periods of nitrogen availability for subsequent use in growth, defence and reproductive processes. The results suggest that higher reduction of recommended dose of nitrogen fertilizer may be inhibitive towards biosynthesis

of amino acid in leaves due to lesser assimilation of inorganic nitrogen transporting amino acid which are used to transfer nitrogen from source organs to sink tissues and to build up reserves during periods of nitrogen availability for subsequent use in growth, defence and reproductive processes. Soil application of biofertilizer was found to be stimulatory over spray application towards biosynthesis of amino acid which indicates lesser availability of nitrogen under non-rhizospheric (spray) application.

Significant level of variation in the level of total soluble protein content in leaves of different varieties of mustard under the field trial of 2003 reveals differential rate of nitrogen assimilation by crop plants. The protein content in leaves significantly reduced from 40% reduction of recommended dose of nitrogen fertilizer ( $T_3$ ) onwards due to lesser supply of nitrogen towards protein synthesis in leaves. Soil application of biofertilizer was found to be superior over spray application in terms of protein content in leaves at different stages of crop growth which might be due to optimum supply of precursor molecule for protein synthesis in crop plants under rhizospheric application (soil application) of biofertilizer.

Significant level of variation in total soluble sugar content in seeds of mungbean varieties may be attributed towards the variable rate of photosynthesis leading to variable rate of accumulation of sugar amount in seeds of six mungbean varieties. During 2004, soluble sugar in seeds increased up to  $T_2$  treatment and then decreased gradually which might be attributed towards utilization of sugar molecules in TCA cycle for amino acid protein synthesis in plant cell. Our findings were similar with earlier findings of Almodares et al., (2009). During 2005, the soil application of biofertilizer at different stages of crop growth was found to be stimulatory over spray application which might be due to higher availability of nitrogen under soil application. This may have increased the vigour of mungbean plants and therefore promoted increased leaf area with higher synthesis of assimilates due to enhance rate of photosynthesis. Such effects have attributed to increased rate of translocation of photosynthetic products from leaves to developing fruits, there by increasing total sugar. Such enhancement of sugar was noted by Magge, (1963) in strawberry plants.

During 2003, the protein content in seeds showed significant variation among the six studied mungbean varieties which might be due to variable rate of protein synthesis in developing seeds under the influence of variable rate of nutrient uptake from the soil by the mungbean varieties. Protein content in seeds reduced significantly with gradual enhanced rate of reduced dose of nitrogen fertilizer as nitrogen is the precursor molecule for protein synthesis. Soil application of biofertilizer promoted greater availability of nitrogen to

mungbean crop plants under the influence of symbiotic nitrogen fixation by the applied *Rhizobium* strain resulting into higher accumulation of protein in mungbean seeds. Our findings were similar with the earlier findings of Selvakumar *et al.*, (2009). Higher availability of nitrogen under soil application of biofertilizer increases amino acid synthesis in the leaves and this stimulates the accumulation of protein in the seed (Patilet *et al.*, 1997).

Amino acid content in seeds showed significant variation among the six studied mungbean varieties indicating towards variable rate of nitrogen uptake by the mungbean varieties. The reduced accumulation of amino acid content in seeds under different reduced dose of nitrogen fertilizer may be due to lesser availability of nitrogen leading to reduced biosynthesis of amino acids in crop plants. Soil application of biofertilizer significantly improved nitrogen availability to crop plants and therefore promoting increased amino acid synthesis in leaves leading to higher accumulation of amino acid in seeds through translocation (Ram Rao *et al.*, 2007).

Ascorbic acid content in seeds of six studied mungbean varieties varied significantly due to variability in the antioxidant property of different varieties under oxidative stress condition. Low dose of nitrogen means low nitrogen mineralization in soil and decrease vitamin-C content in seeds (Tuncay *et al.*, 2004). Elevated level of ascorbic acid content in seeds of mungbean under different rate and time of soil application of biofertilizer may be due to higher availability of nitrogen which promoted the ascorbic acid content in seeds. Our findings are with the line of earlier findings of Bhat, (1999) in strawberry.

The variation in the level of phenolics in seeds of six studied mungbean varieties may be due to growing environment, season and cultivar with cultivar differences being most significant (Zhao *et al.*, 2007). With reduction dose of chemical fertilizer phenol content decreased significantly. The similar observation was noted by Alizadeh *et al.*, (2010). They are found that fertilizer can effect on total phenolic compound in *Saturejahortensish* (Lamiaceae). Higher phenol content in mungbean seeds under the influence of soil application of biofertilizer over spray application might be attributed towards the use of accumulated nitrate in plant thus enabling it to use more carbohydrates for structural growth. (Hanafy Ahmed *et al.*, 2000).

The variation of DNA content in all the six varieties is due to genetic make up of the crop varieties. The nucleic acid content (both DNA and RNA) reduced significantly with higher rate of nitrogen fertilizer reduction dose as biosynthesis of basic nitrogenous compounds such as purines, pyrimidines were inhibited under reduced supply of nitrogen (Devlin, 1972). The soil application of biofertilizer were found to be stimulatory towards availability of nitrogen to crop plants and biosynthesis of purines and pyrimidines leading to higher

nucleic acid (both DNA and RNA) content in mungbean seeds (Devlin, 1972). Nucleic acid (DNA and RNA) contents were much higher obtained from plants treated with biofertilizer. This may be due the active synthesis of nucleic acids concurrently with decreasing the hydrolytic and oxidative enzyme activities. Furthermore, it is known that auxins, gibberellins, and cytokinins stimulate the synthesis of nucleic acids and inhibit the activities of RNase, DNase and 3-nucleotidase (Steward, 1991). RNA content in seeds was found to be higher than DNA content.

Correlation study revealed that only protein and sugar in 2004-05 and amino acid and chlorophyll in 2005-06 in mungbean leaves showed a positive correlation at 5% level of significance. Moreover in 2004, mungbean seeds sugar shows positive relationship with phenol and DNA at 1% level of significance. Ascorbic acid showed a positive correlation with amino acid; protein at 0.05 level of significance and with RNA at 1% level of significance. Phenol level in seeds showed positive correlation with protein and DNA at 5% and 1% level of significance respectively. Ascorbic acid showed a positive relationship with protein and RNA at 5% level and 1% level of significance respectively. In 2004, under subsequent reduction of nitrogen fertilizer dose amino acid was found to be positively correlated with protein, ascorbic acid, DNA and RNA at 1% level of significance. Similarly ascorbic acid and protein individually showed positive correlation at 1% level of significance with RNA and DNA. During 2005, a positive correlation exists between sugar and DNA, RNA and amino acid with protein and RNA at 1% level of significance (Table-5).

**Table 1.**Biochemical contents of leaves of six mungbean varieties during experimental field trial of 2003

Variety	Chlorophyll (mg.g <sup>-1</sup> f.w)	Soluble sugar (g/100g)	Amino acid (g/100g)	Protein (g/100g)
V <sub>1</sub>	0.206 <sup>bc</sup>	22.91 <sup>b</sup>	35.49 <sup>a</sup>	36.52 <sup>bc</sup>
V <sub>2</sub>	0.272 <sup>ab</sup>	23.71 <sup>d</sup>	35.49 <sup>ab</sup>	36.96 <sup>b</sup>
V <sub>3</sub>	0.326 <sup>a</sup>	24.91 <sup>bc</sup>	40.19 <sup>a</sup>	41.68 <sup>a</sup>
V <sub>4</sub>	0.130 <sup>c</sup>	22.42 <sup>a</sup>	32.66 <sup>bc</sup>	34.23 <sup>d</sup>
V <sub>5</sub>	0.074 <sup>de</sup>	21.40 <sup>e</sup>	24.46 <sup>d</sup>	30.87 <sup>e</sup>
V <sub>6</sub>	0.120 <sup>de</sup>	17.91 <sup>f</sup>	23.40 <sup>e</sup>	30.25 <sup>e</sup>

Note: Means followed by the same letter within a treatment were not significantly different at 5% level using Duncan's multiple range test (DMRT). Means of three replicates were taken.

**Table 2.** Biochemical contents of leaves of mungbean (PDM-54) during experimental field trials of 2004 and 2005

Treatments	Chlorophyll (mg.g <sup>-1</sup> f.w)		Soluble sugar (g/100g)		Amino acid (g/100g)		Protein (g/100g)	
	2004	2005	2004	2005	2004	2005	2004	2005
T <sub>1</sub>	1.094 <sup>e</sup>	0.761 <sup>ad</sup>	18.96 <sup>l</sup>	30.46 <sup>e</sup>	20.85 <sup>e</sup>	33.62 <sup>ce</sup>	36.69 <sup>de</sup>	40.26 <sup>e</sup>
T <sub>2</sub>	1.810 <sup>a</sup>	0.673 <sup>ac</sup>	44.00 <sup>a</sup>	39.36 <sup>d</sup>	25.78 <sup>a</sup>	35.63 <sup>cd</sup>	40.17 <sup>a</sup>	40.84 <sup>bcd</sup>
T <sub>3</sub>	1.326 <sup>cd</sup>	0.916 <sup>ab</sup>	38.40 <sup>bc</sup>	42.12 <sup>ab</sup>	24.84 <sup>bc</sup>	40.65 <sup>b</sup>	38.12 <sup>bc</sup>	42.99 <sup>ab</sup>
T <sub>4</sub>	1.254 <sup>cde</sup>	0.867 <sup>ac</sup>	23.56 <sup>d</sup>	40.23 <sup>c</sup>	24.18 <sup>cd</sup>	36.03 <sup>c</sup>	37.59 <sup>cd</sup>	42.64 <sup>abc</sup>
T <sub>5</sub>	1.476 <sup>bc</sup>	1.062 <sup>a</sup>	17.02 <sup>e</sup>	42.82 <sup>a</sup>	19.38 <sup>f</sup>	43.38 <sup>a</sup>	33.22 <sup>f</sup>	43.68 <sup>a</sup>
T <sub>6</sub>	1.583 <sup>ab</sup>		38.41 <sup>bc</sup>		25.64 <sup>ab</sup>		38.96 <sup>b</sup>	

Note: Means followed by the same letter within a treatment were not significantly different at 5% level using Duncan's multiple range test (DMRT). Means of three replicates were taken.

**Table 3.** Biochemical contents in freshly harvested seeds of six mungbean varieties during experimental field trial of 2003

Varieties	Soluble sugar (g/100g)	Amino acid (g/100g)	Protein (g/100g)	Phenol (µg/g)	Ascorbic acid (mg/100g)	DNA (mg/100g)	RNA (mg/100g)
V <sub>1</sub>	7.61 <sup>bc</sup>	24.60 <sup>bc</sup>	28.22 <sup>bc</sup>	751 <sup>c</sup>	8.61 <sup>c</sup>	482.35 <sup>c</sup>	766.92 <sup>c</sup>
V <sub>2</sub>	8.22 <sup>b</sup>	25.30 <sup>b</sup>	29.54 <sup>ab</sup>	906 <sup>b</sup>	11.20 <sup>b</sup>	510.97 <sup>b</sup>	888.57 <sup>ab</sup>
V <sub>3</sub>	9.30 <sup>a</sup>	31.80 <sup>a</sup>	29.66 <sup>a</sup>	947 <sup>a</sup>	13.13 <sup>a</sup>	547.47 <sup>a</sup>	895.01 <sup>a</sup>
V <sub>4</sub>	6.60 <sup>d</sup>	22.90 <sup>d</sup>	28.09 <sup>cd</sup>	623 <sup>d</sup>	8.03 <sup>cd</sup>	397.15 <sup>de</sup>	744.97 <sup>d</sup>
V <sub>5</sub>	6.33 <sup>e</sup>	18.50 <sup>e</sup>	26.51 <sup>e</sup>	620 <sup>de</sup>	7.65 <sup>df</sup>	394.66 <sup>de</sup>	724.74 <sup>e</sup>
V <sub>6</sub>	5.01 <sup>f</sup>	15.30 <sup>f</sup>	17.54 <sup>f</sup>	605 <sup>f</sup>	7.51 <sup>f</sup>	243.32 <sup>f</sup>	984.73 <sup>e</sup>

Note: Means followed by the same letter within a treatment were not significantly different at 5% level using Duncan's multiple range test (DMRT). Means of three replicates were taken.

**Table 5.** Correlation between different biochemical variables in leaves of mungbean plants during 2003, 2004 and 2005

Year of Experiment	Variables	r-value	Levels of significance
2003(leaf)			No significance correlation was found
2004(leaf)	Protein and sugar	0.762	0.05
	Sugar and chlorophyll	-0.897	0.01
	Amino acid and chlorophyll	0.756	0.05
	Amino acid and sugar	-0.822	0.01
2003(seed)	Phenol and sugar	0.793	0.01
	Sugar and DNA	0.925	0.01
	Ascorbic acid and amino acid	0.476	0.05
	DNA and amino acid	-0.565	0.05

	Phenol and protein	0.461	0.05
	Ascorbic acid and protein	0.572	0.05
	RNA and protein	0.549	0.05
	Phenol and DNA	0.668	0.01
	Ascorbic acid RNA	0.917	0.01
2004(seed)	Protein and sugar	-0.463	0.05
	Ascorbic acid and sugar	-0.844	0.01
	Protein and amino acid	0.865	0.01
	Ascorbic acid and amino acid	0.655	0.01
	DNA and amino acid	0.955	0.01
	RNA and amino acid	0.865	0.01
	Ascorbic acid and protein	0.796	0.01
	DNA and protein	0.681	0.01
	RNA and protein	0.695	0.01
	DNA and ascorbic acid	0.527	0.05
	RNA and ascorbic acid	0.679	0.01
2005(seed)	DNA and Sugar	0.609	0.01
	RNA and sugar	0.638	0.01
	Protein and amino acid	0.909	0.01
	Ascorbic acid amino acid	-0.887	0.01
	RNA and amino acid	0.556	0.01
	Ascorbic acid and protein	-0.946	0.01
	Ascorbic acid and phenol	-0.447	0.05
	DNA and phenol	-0.756	0.01
	RNA and phenol	0.675	0.01

## Conclusion

This study concludes that reduced dose of chemical fertilizer and different method and rate of application of biofertilizer reflects the variable biochemical response of mungbean plants towards synthesis and accumulation of simple organic molecules such as sugar, free amino acid, chlorophyll and protein in leaves and sugar, free amino acid, phenol, ascorbic acid, nucleic acid in seeds which reflects crop plant metabolism leading to better crop growth and yield. Higher reduction in the recommended dose of nitrogen fertilizer lead to inhibition of biosynthesis and accumulation of chlorophyll, sugar, protein,

ascorbic acid, phenol and nucleic acid (both DNA and RNA) in leaves and seeds respectively. Although 30% reduction of nitrogen fertilizer dose and soil application of biofertilizer (T<sub>3</sub> and T<sub>5</sub>) were found to be stimulatory towards accumulation of biological macromolecules in leaves and seeds as well as accumulation of secondary plant metabolites such as phenolics. From this it can be interpreted that chemical fertilizer reduction upto a certain level and soil application of biofertilizer is a suitable strategy in regulating plant metabolism leading to accumulation of certain metabolites for optimum plant growth along with enhanced plant defence system against insect and disease attack as well as through antioxidant defence mechanism which may adversely affect the growth and yield attributes of crop plants under agroclimatic condition of old alluvial soil, Burdwan, India. Nonetheless, more research efforts and participatory evaluation of the effects of reduction of chemical fertilizer dose and soil application of biofertilizer on biochemical response of crop plants are needed in this area.

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**Table 4.** Biochemical contents in seeds of mungbean (PDM-54) during experimental field trials of 2004 and 2005

Treatments	Soluble sugar (g/100g)		Amino acid (g/100g)		Protein(g/100g)		Phenol(µg/g)		Ascorbic acid (mg/100g)		DNA(mg/100g)		RNA (mg/100g)	
	2004	2005	2004	2005	2004	2005	2004	2005	2004	2005	2004	2005	2004	2005
T <sub>1</sub>	6.74 <sup>d</sup>	6.53 <sup>bcd</sup>	25.30 <sup>e</sup>	30.32 <sup>e</sup>	30.32 <sup>cde</sup>	28.65 <sup>cd</sup>	889 <sup>e</sup>	663 <sup>c</sup>	8.75 <sup>de</sup>	6.93 <sup>abcd</sup>	480.62 <sup>abcde</sup>	398.69 <sup>de</sup>	876.43 <sup>abcde</sup>	788.43 <sup>cde</sup>
T <sub>2</sub>	9.76 <sup>a</sup>	6.71 <sup>bcd</sup>	44.16 <sup>a</sup>	34.52 <sup>d</sup>	37.00 <sup>a</sup>	28.93 <sup>cd</sup>	946 <sup>a</sup>	723 <sup>d</sup>	10.86 <sup>a</sup>	7.26 <sup>abcd</sup>	486.94 <sup>a</sup>	401.11 <sup>d</sup>	887.66 <sup>ab</sup>	810.77 <sup>cd</sup>
T <sub>3</sub>	7.61 <sup>bc</sup>	7.53 <sup>ab</sup>	39.76 <sup>bc</sup>	40.12 <sup>ab</sup>	32.50 <sup>bc</sup>	33.31 <sup>ab</sup>	916 <sup>c</sup>	763 <sup>b</sup>	9.77 <sup>bc</sup>	8.13 <sup>ab</sup>	483.77 <sup>abc</sup>	489.36 <sup>ab</sup>	885.57 <sup>abc</sup>	854.56 <sup>ab</sup>
T <sub>4</sub>	7.61 <sup>bc</sup>	7.25 <sup>abc</sup>	33.21 <sup>d</sup>	36.32 <sup>c</sup>	30.50 <sup>cd</sup>	31.54 <sup>abc</sup>	910 <sup>cd</sup>	748 <sup>c</sup>	9.31 <sup>cd</sup>	7.55 <sup>abc</sup>	480.76 <sup>abcd</sup>	411.37 <sup>c</sup>	885.41 <sup>abcd</sup>	827.09 <sup>abc</sup>
T <sub>5</sub>	5.03 <sup>f</sup>	8.71 <sup>a</sup>	13.73 <sup>e</sup>	40.32 <sup>a</sup>	29.29 <sup>f</sup>	34.21 <sup>a</sup>	857 <sup>f</sup>	892 <sup>a</sup>	8.13 <sup>e</sup>	8.26 <sup>a</sup>	478.11 <sup>abcde</sup>	495.23 <sup>a</sup>	872.63 <sup>abcdef</sup>	881.78 <sup>a</sup>
T <sub>6</sub>	8.42 <sup>ab</sup>		40.30 <sup>b</sup>		33.10 <sup>b</sup>		923 <sup>b</sup>		10.46 <sup>ab</sup>		486.64 <sup>ab</sup>		886.32 <sup>ab</sup>	

Note: Means followed by the same letter within a treatment were not significantly different at 5% level using Duncan's multiple range test (DMRT). Means of three replicates were taken.