

# EFFECT OF PHOSPHATE SOLUBILIZERS AND FYM ON MICROBIAL POPULATION OF SOYBEAN FIELD [GLYCINE MAX (L.) MERRILL]

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## **KEYWORDS**

Microbial population Aspergillus awamori Bacillus polymixa FYM PSB Soybean

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## INTRODUCTION

Farmers are facing severe problem on availability of chemical fertilizers for soybean production. Growers generally use chemical fertilizers to increase soybean production. However, it gives hazardous effect as soil and water pollution. Biofertilizer (Rhizobium, PSB) and FYM (organic manure) compared with chemical fertilizers are an attractive and environmental safety method of soybean production as it helps to minimize the use of chemical fertilizer and proved environmental safe and ecological sustainable. Phosphorus is an essential major nutrient for the development of plants as it stimulates early development and promotes healthy growth of seedlings. It also enhances the formation of nodules and nitrogen fixation in legumes. Many scientists used various selected strains of phosphate solubilizers which increase the dry matter, grain yield and 'P' uptake (Ahmad and Jha, 1982), PSB increased grain yield and nodulation (Chandra et al., 1995). Application of fertilizer nutrients along with FYM, use of nitrogen fixers, phosphate solubilizers and VAM increased grain and straw yield (Saini et al., 2005). Sarawgi et al. (2012) reported that seed treatment with biofertilizers had their significant effect on microbial population in conjunction with P application in soybean field. The main objective of this study was to assess the effect of bio-fertilizers and FYM on microbial population for promoting better growth of soybean.

## MATERIALS AND METHODS

A field experiment was conducted at experimental farm of

ABSTRACT

Soybean [*Glycine max* (L.) Merrill] is an all famous oil seed as well as pulse crop which contains 40-44 % protein, 20 % oil and many other nutrients. A field experiment was conducted to know the effect of phosphate solubilizing bacteria and fungi and FYM on microbial population of soybean field during *Kharif* season 2009-10. Among treatments, *Aspergillus awamori and Bacillus polymixa* were used as seed treatment @ 20 g/kg seed and phosphorus levels were given through SSP (50 and 25 %  $P_2O_5$ ) and FYM levels (5 and 2.5 t/ha) as applied into the soil. Seed treatment with *A. awamori* increased fungal (25.25 and 29.06 cfu x 10<sup>3</sup>/g) and actinomycetes population (23.44 and 26.19 cfu x 10<sup>8</sup>/g) while *B. polymixa* increased phosphate solubilizing bacterial counts (21.37 and 23.31 cfu x 10<sup>6</sup>/g) significantly at 30 and 60 DAS, respectively. Application of FYM (5 t/ha) had significantly increased the fungi (22.21 and 27.25 cfu x 10<sup>3</sup>/g), actinomycetes (20.37 and 23.77 cfu x 10<sup>8</sup>/g), bacterial (30.55 and 36.02 cfu x 10<sup>6</sup>/g) and PSB population (18.42 and 21.30 cfu x 10<sup>6</sup>/g) in the soybean field at 30 and 60 DAS, respectively. Thus, it can be concluded that the application of *Aspergillus awamori* (20 g/kg seed) and FYM (5 t/ha) are better to increase microbial population as well as yield attributes of soybean in the field.

Department of Plant Pathology, College of Agriculture, Nagpur during kharif season 2009-10. The experiment was laid out in a Factorial Randomized Block Design (FRBD) with four replications. Among the treatments, the carrier based inoculants of Aspergillus awamori and Bacillus polymixa obtained from Plant Pathology Section, College of Agriculture, Nagpur were used as seed treatment @ 20 g/kg seed and phosphorus levels through SSP (50 and 25 % P2O5 ) and FYM (5 and 2.5 t/ha at 10 days before sowing) as applied into the soil. A common dose of 30 kg N/ha was applied in all the plots. The soybean cultivar JS-335 was drilled at geometry of 30 x 5 cm. Prior drilling, the seeds were treated with Aspergillus awamori and Bacillus polymixa @ 20 gm/kg seed. For population count, an isolation of fungi, bacteria, PSB, and actinomycetes were made from rhizospheric soils following serial dilution plate method of Vincent (1970) at the time of 30, 60 and 90 DAS, respectively.

The selected Potato Dextrose Agar medium was used for isolation of fungi, nutrient agar medium for bacterial isolation, Pikovaskaya's medium for PSB isolation and Kenknight medium used for actinomycetes. In each plate, 20 ml medium was poured and rotated with hands for even distribution of suspension and allowed to settle down. The plates were incubated at room temperature  $28 \pm 2^{\circ}$ C. After four days of incubation, the total numbers of colonies were counted and calculated the number of microorganism per ml of original suspension as follows:

Organism in 1 g of samples =

Amount of diluted suspension x dilution factor

Treatments	Fungal pop	Fungal population (cfu x 10 <sup>3</sup> / gm)	10³/ gm)	Bacterial p	Bacterial population (cfu x 10 <sup>6</sup> / gm)	ı x 10 <sup>6</sup> / gm)	PSB popula	PSB population (cfu x 10 <sup>6</sup> / gm)	(mg /ət	Actinomyc	Actinomycetes population (cfu x 10 <sup>8</sup> / gm)	1 (cfu x 10 <sup>8</sup> / gm)
	30 DAS	60 DAS	90 DAS	30 DAS	60 DAS	90 DAS	30 DAS	60 DAS	90 DAS	30 DAS	60 DAS	90 DAS
FYM levels												
5 t/ha	22.21	27.25	24.85	30.55	36.02	33.47	18.42	21.30	17.72	20.37	23.77	20.35
2.5 t/ha	18.10	22.30	21.04	27.72	32.87	29.5	14.82	18.20	14.80	18.02	19.77	17.90
S.Em+	0.13	0.08	0.08	0.05	0.18	0.09	0.07	0.10	0.09	0.10	0.10	0.06
CD (P = 0.05)	0.39	0.25	0.25	0.15	0.52	0.27	0.21	0.29	0.26	0.29	0.30	0.19
PSB												
A. awamori	25.25	29.06	26.93	30.00	36.13	33.00	18.18	21.18	17.31	23.44	26.19	23.25
B. polymixa	21.87	27.38	25.56	31.31	39.31	34.63	21.37	23.31	19.00	20.81	25.31	22.06
50% P,O,/ha	19.15	24.75	22.62	28.93	34.00	31.37	16.25	19.81	16.37	18.81	21.44	17.63
25% P <sub>2</sub> O <sub>2</sub> /ha	18.06	22.75	20.88	28.63	32.25	30.18	14.87	18.50	15.75	16.93	18.38	16.88
Control	16.43	19.93	18.73	26.81	30.56	29.13	12.43	15.94	12.88	16.00	17.56	15.81
S.Em+	0.97	0.62	0.63	0.38	1.29	0.67	0.51	0.70	0.65	0.72	0.75	0.47
CD (P = 0.05)	2.82	1.81	1.83	1.10	3.74	1.95	1.49	2.05	1.89	2.09	2.18	1.38
Interaction												
S.Em+	1.37	0.88	0.89	0.53	1.82	0.95	0.73	1.00	0.92	1.02	1.09	0.67
CD (P = 0.05)	ı		,	ı	,	,	,				,	

#### **RESULTS AND DISCUSSION**

#### Effect of phosphate solubilizers

It is cleared from the data (Table 1) that there were significant differences on fungal, bacteria, PSB and actinomycetes population at all the intervals. Maximum population of fungi  $(29.06 \text{ cfu x } 10^3/\text{g})$  and actinomycetes  $(26.19 \text{ cfu x } 10^8/\text{g})$ were attained at 60 DAS by treating the seeds with Aspergillus awamori and found significantly superior over all the treatments which were followed Bacillus polymixa (27.38 cfu x 10% g and 25.31 cfu x 108/g, respectively). The increase in fungal population from 30 to 60 DAS may be due to growth promoting substances secreted during crop growth period. At 90 DAS, there was a decrement in counts in all the treatments. Similar results were obtained by Gupta et al. (1992). Present results are also in accordance with the findings of Saini et al. (2005) where microbial biomass, C, N and P contents in the rhizosphere soil of soybean were maximum from 30-60 DAS and decreased from 60 DAS or harvest after applying fertilizer nutrients along with FYM, use of nitrogen fixers, phosphate solubilizers and VAM. Babana and Antoun (2006) reported that phosphate solubilizing fungal isolates of Aspergillus awamori Nakazawa C<sub>1</sub> and Penicillium chrysogenum Thom C<sub>13</sub> increased microbial population and root dry matter yield. Sarawgi et al. (2012) reported that seed treatment with biofertilizers had their significant effect on microbial population in conjunction with P application in soybean field. Significant differences were also observed on bacterial and PSB population at all the intervals. Maximum population of bacteria (39.31 cfu x 10<sup>6</sup>/g) and PSB (23.31cfu x 10<sup>6</sup>/g) were attained by seed inoculation with *B. polymixa* at 60 DAS and it was found significantly superior over all the treatments and was followed by seed treatment with A. awamori (36.13 cfu x 10<sup>6</sup>/g and 21.18 cfu x 10<sup>6</sup>/g, respectively). At 60 DAS, there was increased population counts in bacterial and PSB treatments and decreased at 90 days in all the treatments (Table 1). These results are in agreement with Kundu and Gaur (1980), Saini et al. (2005) and Qureshi et al. (2005).

#### Effect of FYM

The data (Table 1) showed that the application of FYM 5 t/ha had significantly increased the fungi, actinomycetes, bacterial and PSB population in soybean field up to 60 DAS. Maximum population of fungi (27.25 cfu x 10<sup>3</sup>/gm), actinomycetes (23.77 cfu x 10<sup>8</sup>/gm), bacterial (36.02 cfu x 10<sup>6</sup>/gm) and PSB (21.30 cfu x 10<sup>6</sup>/gm) was recorded by application of 5 t/ha FYM as compared to 2.5 t/ha. At 60 DAS, there was an increased viable count in all the treatments which are in agreement with Kundu and Gaur (1980), Gupta et al. (1992), Qureshi et al. (2005), Saini et al. (2005) and Chaturvedi et al. (2010). This may be due to fact that there might have been more amount of degradation of organic matter in soil which resulted in increased microbial populations (Table 1) while at 90 DAS, it was decreased in all the treatments. The decrease in population may be due to proceeding of crop to maturity stage.

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