

Comparative Analysis of Growth Responses in Indian Mustard to Azotobacter Priming and Vermicompost for Sustainable Cultivation

A study investigating the responses of Indian mustard (*Brassica juncea*) to biopriming with *Azotobacter*, incorporation of vermicompost and their comparative analysis was performed in 2024 Rabi season in new alluvial zone at Bidhan Chandra Krishi Viswavidyalaya, Nadia district, West Bengal. The research involved ten genotypes of Indian mustard and four treatments in completely randomized design. Treatments were designed as seeds sown in field soil, *Azotobacter*-primed seeds in field soil, seeds sown in vermicompost mixed with field soil, and *Azotobacter*-primed seeds in vermicompost with field soil. Eight key parameters were considered i.e. germination rate, seedling fresh and dry weight, seedling length, vigour index I and II, proline, and chlorophyll content. Results established improvements in germination and other growth parameters, particularly with the treatment combining *Azotobacter* priming and vermicompost, which presented the highest values across most genotypes. The performance of TM 306 - 1 and TBM 143 genotypes had produced the best results. The results emphasize the potential for utilizing biofertilizers and organic amendments in sustainable mustard cultivation, providing an effective substitute for chemical fertilizers.

ABSTRACT

Keywords: Azotobacter, Vermicompost, Seed Priming, Indian Mustard

1. INTRODUCTION

Among the oilseeds, Indian mustard, from the Brassicaceae family, is a significant oilseed crop as it occupies a very vital place in the Indian agriculture scenario. It ranks second to groundnut in both area and production and is responsible for about 80% of the total rapeseed-mustard production. The mustard seeds are rich in nutrients, having an oil content that ranges from 38 to 50% and comprising erucic acid, linoleic acid, and oleic acid (Bater Dabi *et al.*, 2001; Gantait *et al.*, 2024; Janaki *et al.*, 2022; Kaushik *et al.*, 2024). The adverse effects of chemical fertilizers on Indian agriculture, both on soil and human health are manifold. The long-term effects of synthetic fertilizer usage have resulted in soil degradation, including reduced fertility and increased soil pH, so that at one point in time, it might turn unproductive land (Bhokare P. R. & Wankhade R. R., 2024; Dube *et al.*, 2024). Excessive use of chemical fertilizers also leads to the contamination of the soil with metals, like cadmium and lead, which impose extensive environmental and health hazards (Dash *et al.*, 2022). Farmers are complaining that synthetic fertilizers not only detract nutritional quality from the crops but also taste bad, in addition to causing health problems such as hemoglobin disorders and chronic health issues because of high nitrate levels (Nichols, 2023). To solve the problem, biopriming with bio-fertilizers like *Azotobacter* and some organic amendments like vermicompost are good areas to explore as these provide sustainable and eco-friendly alternatives. Biopriming, a sort of seed treatment, refers to soaking seeds in a solution containing a beneficial microorganism. Microorganisms like bacteria or fungi colonize and, in some cases, penetrate the seed coat (Gantait *et al.*, 2024; Govind *et al.*, 2024). Freelifving nitrogen fixing bacteria i.e. *Azotobacter* can convert atmospheric nitrogen into an available form from which plants can derive. Seed germination and seedling vigor are enhanced by growth-promoting chemical compounds produced by *Azotobacter* (Bater Dabi *et al.*, 2001; Janaki *et al.*, 2022; Kaushik *et al.*, 2024). Vermicompost refers to organic fertilizer formed from the digestion of organic waste materials by earthworms. It is a nutrient-rich semi-bulky organic fertilizer containing high concentrations of macro and micronutrients. Vermicompost is

31 a good additive to soil because it improves soil quality, soil fertility and microbial activity (M. Kumar *et al.*,
32 2023; Singh *et al.*, 2018). The present experiment has been conducted to compare the germination,
33 seedling vigor, chlorophyll and proline content of different varieties when primed and exposed to
34 vermicompost.

35 2. MATERIALS AND METHODS

36 The research investigated the impacts of ten genotypes (G) and four treatments (T) on several
37 seedling growth and physiological metrics in new alluvial zone at Bidhan Chandra Krishi Viswavidyalaya,
38 Nadia district, West Bengal. The experiment was arranged in a completely randomized design with three
39 replicates. Parameters including germination, seedling fresh and dry weight, seedling length, vigor index I
40 and II, proline content, and chlorophyll content were evaluated to determine the growth potential and
41 resilience of the crop under various treatments. The genotypes were BRM4(G₁), BRM13(G₂), BRM 14(G₃),
42 Varuna(G₄), JD 6(G₅), PM 25(G₆), PM 29(G₇), TM 306-1(G₈), TBM 204(G₉) and TBM 143(G₁₀). The four
43 treatments are as follows- Seeds sown in field soil(T₁), Azotobacter primed seeds sown in field soil(T₂),
44 Seeds sown in vermicompost + field soil(T₃), Azotobacter primed seeds sown in vermicompost + field
45 soil(T₄). 50 seeds were placed in each sterilized plastic container and left in open condition. In case of
46 vermicompost treatment 50% of vermicompost and 50% of field soil was used. For Azotobacter seed
47 priming a 5:1 ratio of Azotobacter to seed was maintained and seeds were soaked for one hour then
48 dried. During the time of experiment maximum and minimum temperatures were 31.8°C and 12.4°C
49 respectively, maximum, and minimum relative humidity were 78% and 54% respectively with 8.1 hours of
50 average bright sunshine hours. After the seventh day seedlings from the container were counted and
51 germination(%) was calculated by dividing the number of seeds germinated by the total seeds planted,
52 then multiplied by 100. Ten seedlings were picked gently from container after 7th day and seedling length
53 was measured using a centimeter scale. Average data was presented in centimeter(cm). For seedling
54 fresh weight five random seedlings were taken out from each container and their weight was measured in
55 a weighing balance and the average was calculated. To obtain seedling dry weight they were put in hot air
56 oven till constant temperature was achieved. After that weights of five dry seedlings were observed using
57 a weighing balance and average was calculated. The vigor index I was assessed to evaluate the overall
58 vigor and health of the seedlings under controlled conditions. The vigor index I was calculated using the
59 formula given by Abdul-Baki and Anderson (1973) [Vigor Index I = Germination Percentage × Average
60 seedling length(cm)]. The vigor index II is a critical parameter for evaluating seedling vigor, providing
61 insights into the overall health and growth potential of plants. The vigor index II was calculated using the
62 formula given by Abdul-Baki and Anderson (1973) [Vigor Index II = Germination Percentage ×
63 Mean dry weight of seedlings(mg)]. Proline content was determined spectrophotometrically by adopting
64 the ninhydrin method of Bates *et al.* (1973). Total chlorophyll was estimated following Arnon's method
65 (Arnon, 1949). Statistical analysis was done using OPSTAT (Sheoran *et al.*, 1998).

66 3. RESULTS AND DISCUSSION

67 3.1 Results

68 3.1.1 Germination(%)

69 Among the genotypes highest mean germination was shown by G₃ (99.43%), G₆ (98.87%)
70 followed by G₅ (98.68%), while the lowest mean germination was recorded for G₈ (94.31%). The
71 treatment with the highest germination was T₄ (99.02%) followed by T₃ (97.07%), and the lowest was T₁
72 (95.00%). The interaction of genotypes and treatments showed the three highest germination rates for G₃
73 × T₄ (100.00%), G₆ × T₄ (100.00%), and G₅ × T₄ (100.00%). The difference between T₄ and T₃ was not
74 statistically significant.

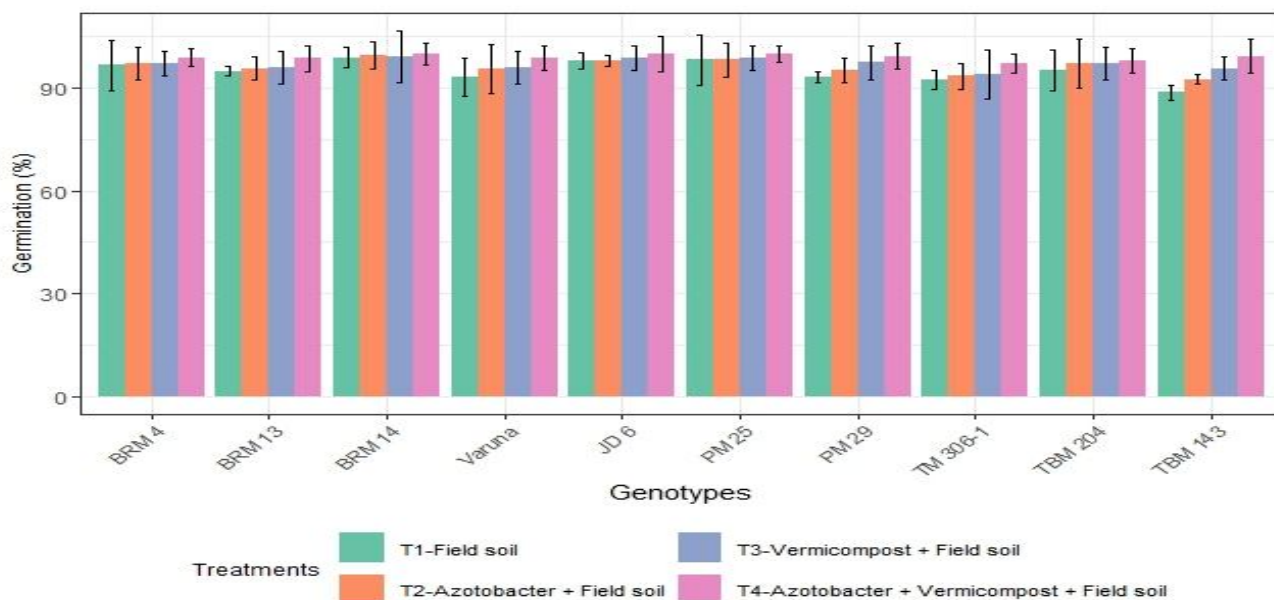


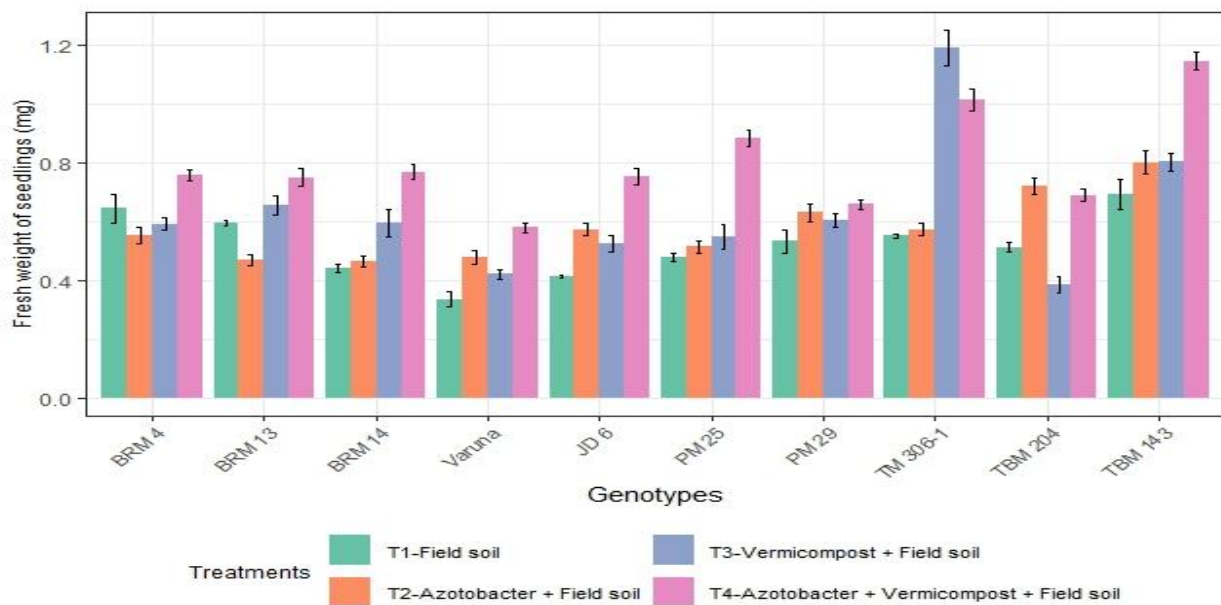
Figure 1. Comparative analysis of germination (%) across different Indian mustard genotypes (G_1 - G_{10}) under four treatments (T_1 - T_4). Error bars represent Standard Error of Means [SEM(\pm)]. Values are means of three replicates.

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77 3.1.2 Fresh weight of seedlings(mg)

78 Genotypes G_{10} (0.861mg), G_8 (0.833 mg), and G_7 (0.607 mg) had the highest seedling fresh
 79 weight while G_4 (0.453 mg) had the lowest. Among treatments, T_4 (0.801 mg) and T_3 (0.632 mg) had the
 80 highest fresh weight of seedlings with T_1 (0.520 mg) being the lowest. Among interaction effects
 81 $G_{10} \times T_4$ (1.147 mg), $G_8 \times T_4$ (1.015 mg), and $G_8 \times T_3$ (1.192 mg) had the most seedling fresh weight. G_{10} and G_8
 82 were significantly different from G_7 , but there was no significant difference between G_{10} and G_8 . The
 83 difference between T_4 and T_3 was significant, suggesting that T_4 provides a notable improvement in fresh
 84 weight.



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Fig 2: Comparative analysis of fresh weight of seedlings (mg) across different Indian mustard genotypes (G_1 - G_{10}) under four treatments (T_1 - T_4). Error bars represent Standard Error of Means [SEm(\pm)]. Values are means of three replicates.

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Table 1: Effect of Azotobacter priming and vermicompost treatments on germination (%) and fresh weight of seedlings (mg) in different Indian mustard genotypes

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Table 1	Germination(%)					Fresh weight of seedlings(mg)				
	T ₁	T ₂	T ₃	T ₄	MeanG	T ₁	T ₂	T ₃	T ₄	MeanG
G₁	96.75	97.25	97.25	99.00	97.56	0.646	0.555	0.592	0.760	0.638
G₂	95.00	95.75	96.00	98.70	96.36	0.594	0.470	0.657	0.751	0.618
G₃	99.00	99.50	99.25	100.00	99.43	0.440	0.465	0.595	0.770	0.568
G₄	93.25	95.75	96.00	98.75	95.93	0.335	0.478	0.421	0.579	0.453
G₅	98.00	98.00	98.75	100.00	98.68	0.414	0.574	0.525	0.752	0.566
G₆	98.25	98.25	99.00	100.00	98.87	0.478	0.514	0.548	0.884	0.606
G₇	93.25	95.25	97.50	99.25	96.31	0.534	0.631	0.604	0.660	0.607
G₈	92.50	93.50	94.00	97.25	94.31	0.551	0.574	1.192	1.015	0.833
G₉	95.25	97.25	97.25	98.00	96.93	0.513	0.722	0.386	0.689	0.578
G₁₀	88.75	92.50	95.75	99.25	94.06	0.692	0.802	0.803	1.147	0.861

Mean T	95.00	96.30	97.07	99.02		0.520	0.579	0.632	0.801	
	Factor G	Factor T	Factor G X T			Factor G	Factor T	Factor G X T		
C.D(5%)	NS	NS	NS			0.040	0.025	0.079		
SEm(±)	2.112	1.336	4.224			0.014	0.009	0.028		

^a G₁: BRM 4; G₂: BRM13; G₃: BRM 14; G₄: Varuna; G₅: JD 6; G₆: PM 25; G₇: PM 29; G₈: TM 306-1; G₉: TBM 204; G₁₀: TBM 143

^b T₁: Seeds sown in field soil; T₂: Azotobacter primed seeds in field soil; T₃: Seeds in vermicompost + field soil; T₄: Azotobacter primed seeds in vermicompost + field soil

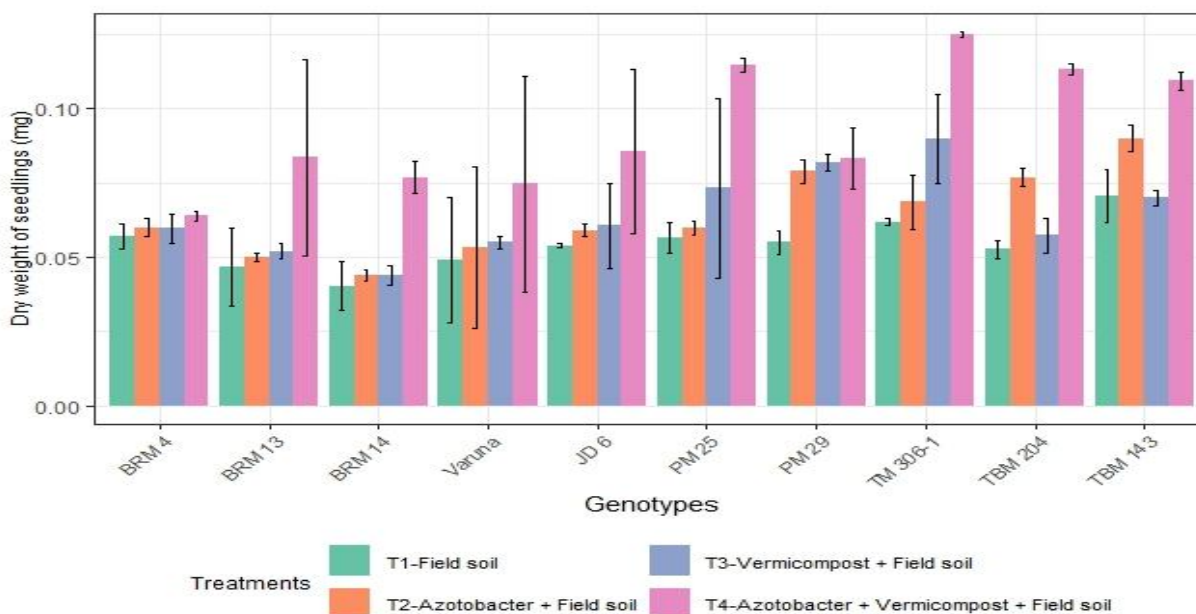
^c CD: Critical Difference

^dSEm(±): Standard Error of Mean e NS: Non-significant

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90 3.1.3 Dry weight of seedlings(mg)

91 In the case of seedling dry weight best genotypes were G₈ (0.086 mg), G₁₀ (0.085 mg), and G₆
 92 (0.076 mg), while the worst one was G₃ (0.051 mg). Among treatments, T₄ (0.093 mg) and T₃ (0.064 mg)
 93 had the highest, with T₁ (0.054 mg) having the lowest value. The best three interactions were G₈ × T₄
 94 (0.125 mg), G₆ × T₄ (0.115 mg), and G₁₀ × T₄ (0.110 mg). Both genotypes and treatments showed
 95 significant difference but G₈, G₁₀, and G₆ exhibited non-significant difference.



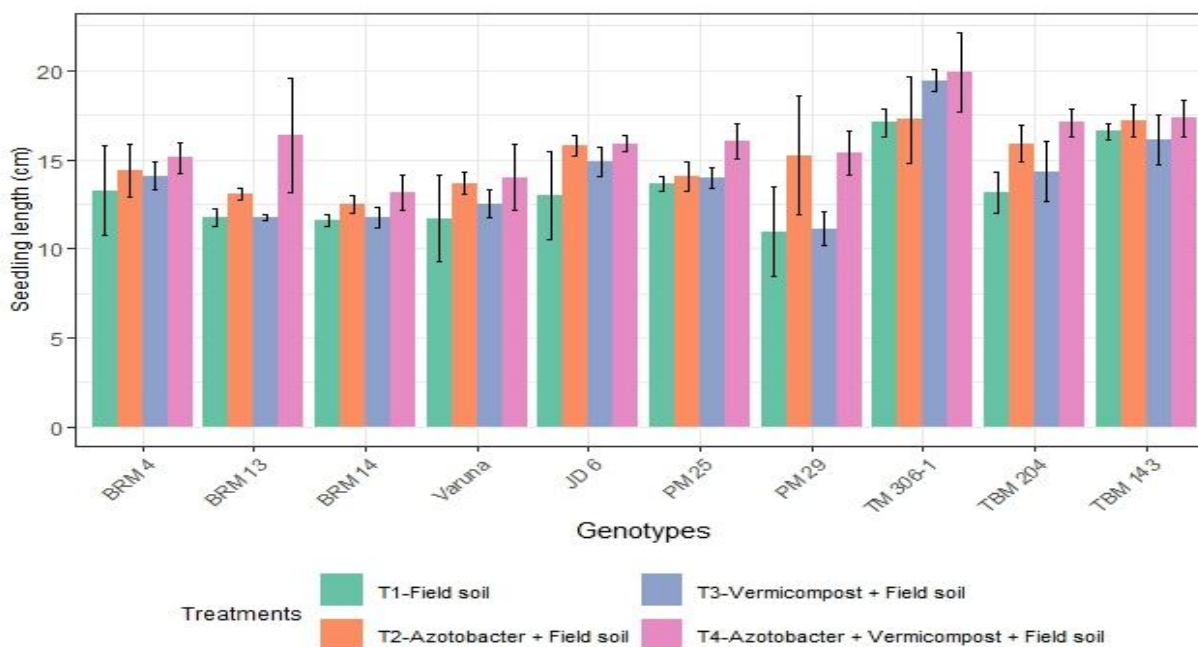
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Fig 3: Comparative analysis of dry weight of seedlings (mg) across different Indian mustard genotypes (G₁-G₁₀) under four treatments (T₁-T₄). Error bars represent Standard Error of Means [SEm(±)]. Values are means of three replicates.

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98 **3.1.4 Seedling length(cm)**

99 G₈ (18.408 cm), G₁₀ (16.804 cm), and G₅ (14.892 cm) recorded maximum seedling length but G₃
 100 (12.233 cm) was the lowest. In the case of treatments T₄ (16.023 cm) and T₂ (14.900 cm) had the highest,
 101 with T₁ (13.270 cm) having the lowest seedling length after 7 days. Both genotypes and treatments were
 102 significant and G₈ was significantly different from G₁₀ and G₅, while the latter two did not differ significantly
 103 from each other. The difference between T₄ and T₂ is significant, highlighting that T₄ strongly enhances
 104 seedling length.



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Fig 4: Comparative analysis of seedling length (cm) across different Indian mustard genotypes (G₁-G₁₀) under four treatments (T₁-T₄). Error bars represent Standard Error of Means [SEm(±)]. Values are means of three replicates.

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107 **Table 2: Effect of Azotobacter priming and vermicompost treatments on dry weight of seedlings**
 108 **(mg) and seedling length (cm) in different Indian mustard genotypes**

Table 2	Dry weight of seedlings(mg)					Seedling length(cm)				
	T ₁	T ₂	T ₃	T ₄	Mean G	T ₁	T ₂	T ₃	T ₄	Mean G
G ₁	0.057	0.060	0.060	0.064	0.060	13.26	14.40	14.06	15.10	14.20

G₂	0.047	0.050	0.052	0.084	0.058	11.73	13.06	11.76	16.36	13.23
G₃	0.040	0.044	0.044	0.077	0.051	11.60	12.46	11.73	13.13	12.23
G₄	0.049	0.053	0.055	0.075	0.058	11.70	13.66	12.53	14.00	12.97
G₅	0.054	0.059	0.061	0.086	0.065	12.96	15.80	14.90	15.90	14.89
G₆	0.057	0.060	0.073	0.115	0.076	13.63	14.03	13.96	16.06	14.42
G₇	0.055	0.079	0.082	0.083	0.075	10.96	15.23	11.13	15.36	13.17
G₈	0.062	0.069	0.090	0.125	0.086	17.06	17.23	19.43	19.90	18.40
G₉	0.053	0.077	0.057	0.113	0.075	13.16	15.90	14.33	17.06	15.11
G₁₀	0.071	0.090	0.070	0.110	0.085	16.60	17.20	16.08	17.33	16.80
Mean T	0.054	0.064	0.064	0.093		13.27	14.90	13.99	16.02	
	Factor G	Factor T	Factor G X T			Factor G	Factor T	Factor G X T		
C.D(5%)	0.018	0.011	NS			2.058	1.302	NS		
SEm(±)	0.006	0.004	0.012			0.730	0.461	1.459		
^a G ₁ : BRM 4; G ₂ : BRM13; G ₃ : BRM 14; G ₄ : Varuna; G ₅ : JD 6; G ₆ : PM 25; G ₇ : PM 29; G ₈ : TM 306-1; G ₉ : TBM 204; G ₁₀ : TBM 143 ^b T ₁ : Seeds sown in field soil; T ₂ : Azotobacter primed seeds in field soil; T ₃ : Seeds in vermicompost + field soil; T ₄ : Azotobacter primed seeds in vermicompost + field soil ^c CD: Critical Difference ^d SEm(±): Standard Error of Mean e NS: Non-significant										

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110 3.1.5 Vigour Index I

111 The best vigour index I was presented by T₄ (1,585) and T₂ (1,429), and T₁ (1,255) was the
112 lowest. The best genotypes for high vigour index I was G₈ (1,736), G₁₀ (1,580), and G₅ (1,471), while the
113 worst was G₇ (1,270). The highest interactions were G₈ × T₄ (1,927), G₈ × T₃ (1,819), and G₁₀ × T₄
114 (1,724). both treatments and genotypes were significant and G₈, G₁₀, and G₅ also showed significant
115 differences among each other. The difference between T₄ and T₂ is highly significant, indicating a
116 substantial effect of T₄ on vigour index I.

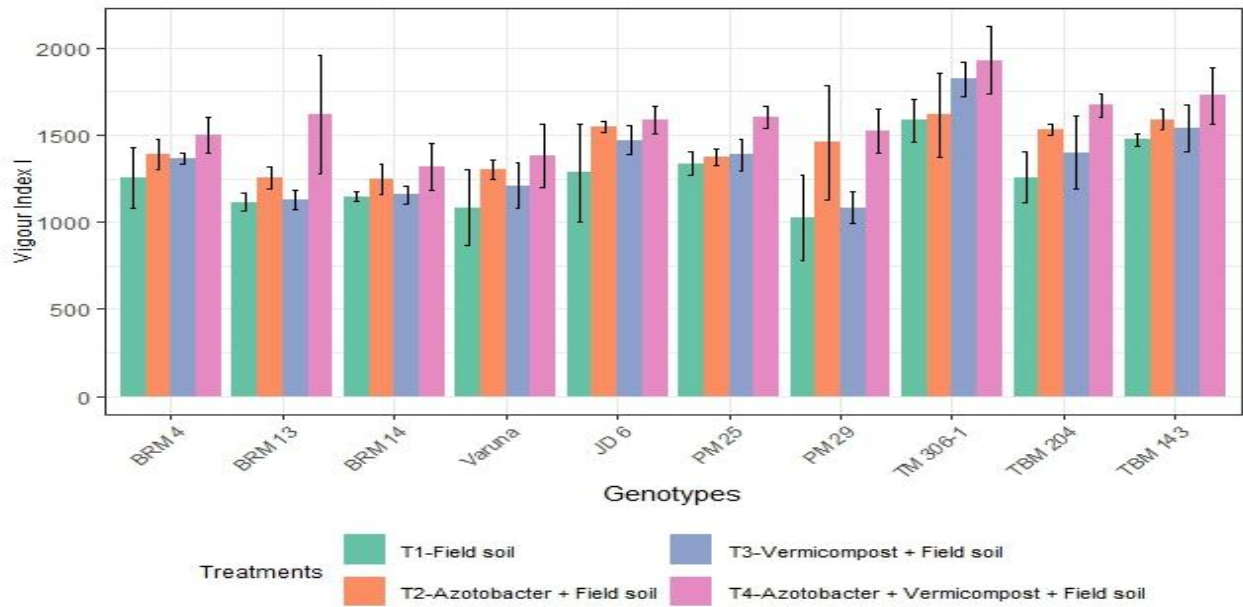
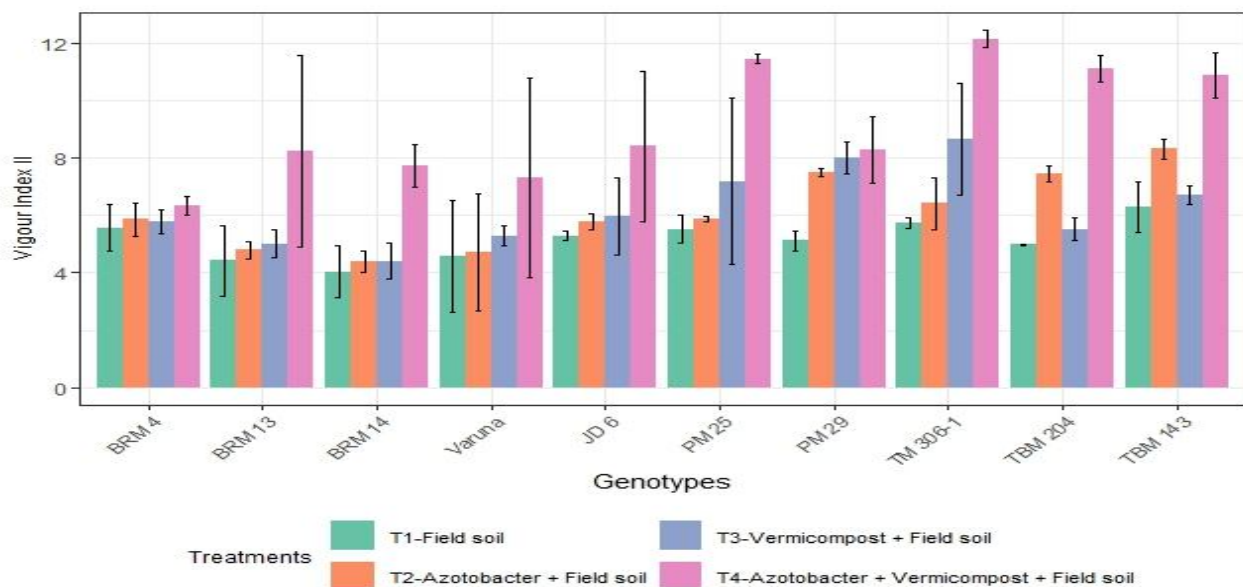


Fig 5: Comparative analysis of vigour index I across different Indian mustard genotypes (G₁-G₁₀) under four treatments (T₁-T₄). Error bars represent Standard Error of Means [SEm(±)]. Values are means of three replicates.

3.1.6 Vigour Index II

The genotypes for vigour index II were G₈ (8.247), G₁₀ (8.053), and G₆ (7.514), with G₃ (5.146) being the lowest. For treatments T₄ (9.198) and T₃ (6.257) were the highest, and T₁ (5.156) was the lowest. The three highest interactions were G₈ × T₄ (12.152), G₆ × T₄ (11.457), and G₁₀ × T₄ (10.896). Both genotypes and treatments were significant, with G₈ being significantly different from G₁₀ and G₆, while G₁₀ and G₆ did not differ significantly. The difference between T₄ and T₃ was significant, reinforcing T₄'s superior performance in vigor improvement.



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Fig 6: Comparative analysis of vigour index II across different Indian mustard genotypes (G₁-G₁₀) under four treatments (T₁-T₄). Error bars represent Standard Error of Means [SEm(±)]. Values are means of three replicates.

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128 **Table 3 :** Effect of Azotobacter priming and vermicompost treatments on Vigour Index I and Vigour

129 **Index II in different Indian mustard genotypes**

Table 3	Vigour Index I					Vigour Index II				
	T ₁	T ₂	T ₃	T ₄	Mean G	T ₁	T ₂	T ₃	T ₄	Mean G
G ₁	1,253	1,386	1,362	1,496	1,374	5.577	5.865	5.784	6.344	5.892
G ₂	1,114	1,252.0	1,129	1,618	1,278	4.423	4.797	5.017	8.258	5.624
G ₃	1,146	1,242	1,157	1,319	1,216	4.041	4.392	4.416	7.733	5.146
G ₄	1,083	1,300	1,209	1,381	1,243	4.571	4.714	5.283	7.335	5.476
G ₅	1,282	1,546	1,469	1,588	1,471	5.293	5.785	5.966	8.412	6.364
G ₆	1,333	1,371	1,384	1,601	1,423	5.528	5.872	7.199	11.457	7.514
G ₇	1,024	1,455	1,081	1,522	1,270	5.121	7.498	8.003	8.281	7.226
G ₈	1,583	1,614	1,819	1,927	1,736	5.734	6.429	8.672	12.152	8.247
G ₉	1,257	1,531	1,399	1,669	1,464	4.980	7.443	5.530	11.109	7.266
G ₁₀	1,472	1,588	1,538	1,724	1,580	6.295	8.318	6.702	10.896	8.053
Mean T	1,255	1,429	1,355	1,585		5.156	6.111	6.257	9.198	
	Factor G	Factor T	Factor G X T			Factor G	Factor T	Factor G X T		
C.D(5%)	206.239	130.437	NS			1.747	1.105	NS		
SEm(±)	73.110	46.239	146.220			0.619	0.392	1.239		

^a G₁: BRM 4; G₂: BRM13; G₃: BRM 14; G₄: Varuna; G₅: JD 6; G₆: PM 25; G₇: PM 29; G₈: TM 306-1; G₉:

TBM 204; G₁₀: TBM 143

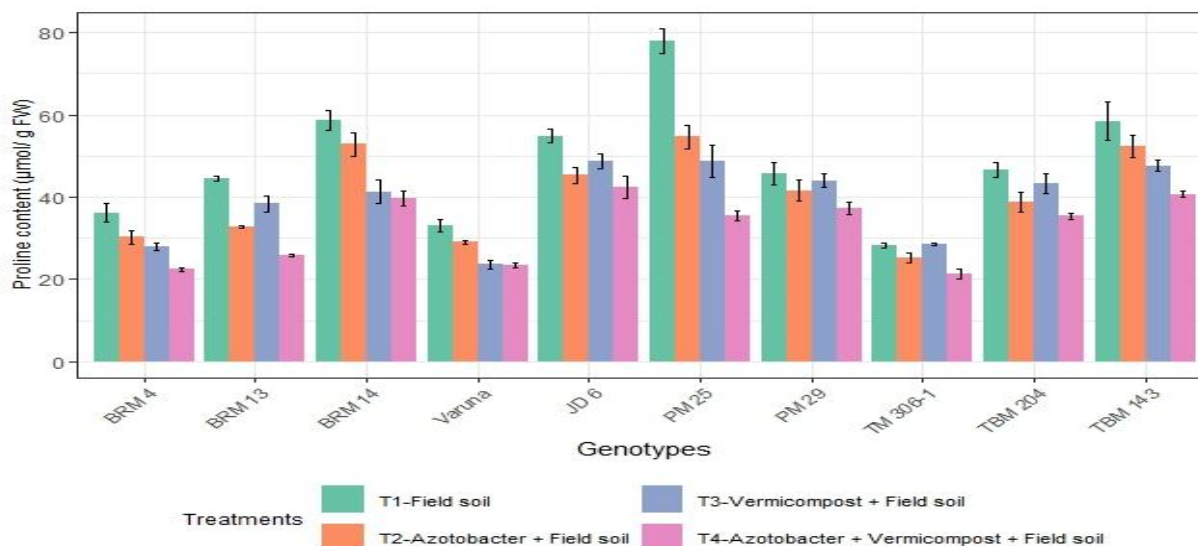
^b T₁: Seeds sown in field soil; T₂: Azotobacter primed seeds in field soil; T₃: Seeds in vermicompost + field soil; T₄: Azotobacter primed seeds in vermicompost + field soil

^c CD: Critical Difference

^oSEm(±): Standard Error of Mean e NS: Non-significant

130 3.1.7 Proline content(μmol/ g FW)

131 In the case of proline content of seedlings, highest values were observed in G₇ (56.311 μmol/ g
 132 FW), G₈ (46.200 μmol/ g FW), and G₄ (44.600 μmol/ g FW), with G₅ (28.567 μmol/ g FW) being the
 133 lowest. Also, in treatments, T₄ (40.801 μmol/ g FW) and T₃ (40.071 μmol/ g FW) were the highest, and T₂
 134 (38.511 μmol/ g FW) was the lowest. The three highest interactions were G₇ × T₄ (77.656 μmol/ g FW), G₆
 135 × T₄ (45.744 μmol/ g FW), and G₈ × T₄ (34.878 μmol/ g FW). The results showed significant differences
 136 between G₇ and the other groups (G₈ and G₄), while G₈ and G₄ were not significantly different. However,
 137 treatments were not significant.



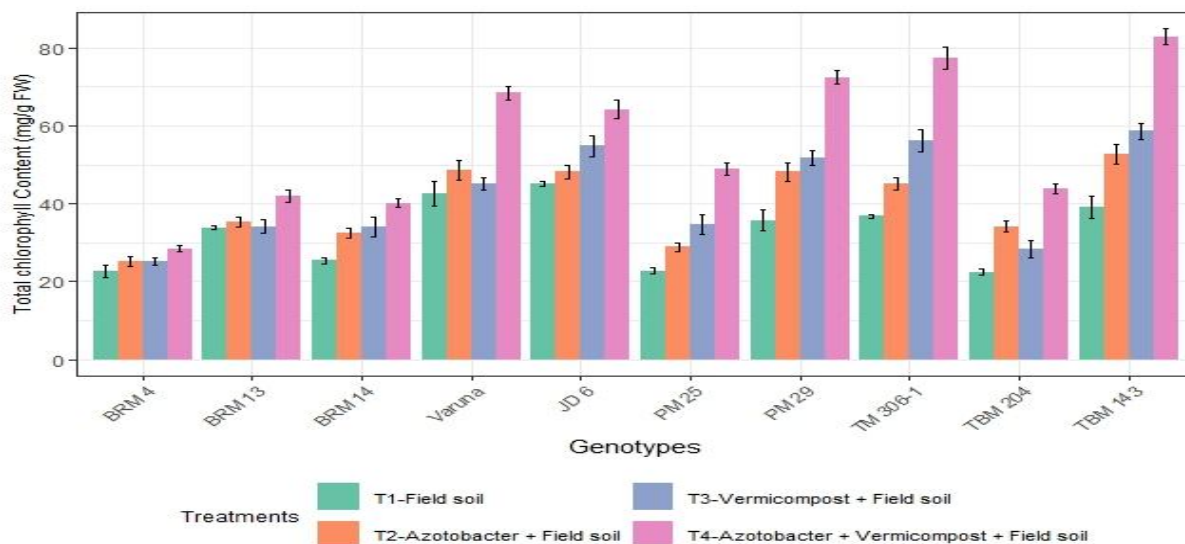
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Fig 7: Comparative analysis of proline content (μmol/ g FW) across different Indian mustard genotypes (G₁-G₁₀) under four treatments (T₁-T₄). Error bars represent Standard Error of Means [SEm(±)]. Values are means of three replicates.

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140 3.1.8 Total chlorophyll Content(mg/g FW)

141 T₄ (56.840 mg/g FW) and T₃ (42.290 mg/g FW) showed the highest chlorophyll content with T₁
 142 (32.640 mg/g FW) being the lowest. For genotypes, G₁₀ (58.300 mg/g FW), G₈ (53.900 mg/g FW), and G₇
 143 (52.000 mg/g FW) had the highest total chlorophyll content values, while G₁ (25.375 mg/g FW) was the
 144 lowest. The three highest interactions were G₁₀ × T₄ (82.800 mg/g FW), G₈ × T₄ (77.400 mg/g FW), and
 145 G₇ × T₄ (72.400 mg/g FW). Factors Genotype, treatment, and their interaction were significant. Significant
 146 differences were observed among G₁₀, G₈, and G₇. The difference between T₄ and T₃ was significant,
 147 indicating that T₄ greatly enhances chlorophyll content.



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Fig 8: Comparative analysis of total chlorophyll Content (mg/g FW) across different Indian mustard genotypes (G₁-G₁₀) under four treatments (T₁-T₄). Error bars represent Standard Error of Means [SEM(±)]. Values are means of three replicates.

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Table 4 : Effect of Azotobacter priming and vermicompost treatments on proline content (µmol/ g FW) and total chlorophyll Content (mg/g FW) in different Indian mustard genotypes

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Table 4	Proline content(µmol/ g FW)					Total chlorophyll Content(mg/g FW)				
	T ₁	T ₂	T ₃	T ₄	Mean G	T ₁	T ₂	T ₃	T ₄	Mean G
G₁	36.20	31.66	30.53	27.28	31.42	22.70	25.20	25.10	28.50	25.37
G₂	22.36	36.92	41.11	32.62	33.25	33.80	35.40	34.20	41.80	36.30
G₃	38.33	30.11	36.97	58.35	40.94	25.40	32.40	34.10	40.20	33.02
G₄	52.80	44.00	41.97	39.62	44.60	42.50	48.60	45.10	68.40	51.15
G₅	33.06	30.15	28.06	22.97	28.56	45.10	48.20	54.80	64.20	53.07
G₆	23.43	44.54	51.16	45.74	41.22	22.80	28.90	34.70	48.90	33.82
G₇	48.66	44.82	54.10	77.65	56.31	35.70	48.20	51.70	72.40	52.00
G₈	54.63	49.57	45.71	34.87	46.20	36.80	45.20	56.20	77.40	53.90
G₉	45.63	42.01	44.07	43.30	43.75	22.50	34.20	28.40	43.800	32.22
G₁₀	37.30	31.30	26.98	25.56	30.28	39.10	52.70	58.60	82.80	58.30
Mean T	39.24	38.51	40.07	40.80		32.64	39.90	42.29	56.84	

	Factor G	Factor T	Factor G X T	Factor G	Factor T	Factor G X T
C.D(5%)	5.773	NS	11.545	2.627	1.661	5.253
SEm(±)	2.046	1.294	4.093	0.931	0.589	1.862
^a G ₁ : BRM 4; G ₂ : BRM13; G ₃ : BRM 14; G ₄ : Varuna; G ₅ : JD 6; G ₆ : PM 25; G ₇ : PM 29; G ₈ : TM 306-1; G ₉ : TBM 204; G ₁₀ : TBM 143						
^b T ₁ : Seeds sown in field soil; T ₂ : Azotobacter primed seeds in field soil; T ₃ : Seeds in vermicompost + field soil; T ₄ : Azotobacter primed seeds in vermicompost + field soil						
^c CD: Critical Difference						
^d SEm(±): Standard Error of Mean e NS: Non-significant						

152

153 3.2 Discussion

154 **The** germination study indicated higher mean values for treatments G₃, G₆, and G₅, especially under
155 T₄, which consistently demonstrated greater germination rates. **However**, the interactions between
156 genotype and treatment **were** non-significant which means the treatments did have an effect but their
157 influence was largely consistent among genotypes. Many studies reported the positive effect of
158 vermicompost on the germination and growth of mustard seedlings and plants (Merta, 2023; Haque & Ali,
159 2020; Reza, 2023; Reza *et al.*, 2022). The improved germination upon vermicompost application could be
160 attributed to several factors. Vermicompost has many available mineral nutrients, humic substances, and
161 plant growth-promoting agents such as auxins, which are known to improve seed germination and
162 seedling growth (Bhattacharya *et al.*, 2019; Pathma & Sakthivel, 2012). Vermicompost helps in increasing
163 the porosity, aeration, and water retention capabilities which enhance the germination and growth of
164 mustard plants (Sarma & Gogoi, 2015; Merta, 2023). Azotobacter further increases seed germination in
165 crops by ensuring plant health through nitrogen fixation, phosphate solubilization, and growth hormone
166 production. Together, these factors lead to optimum growth, increased vigor, and effective germination
167 (Abbas *et al.*, 2024). Azotobacter has been found effective in promoting seed germination in several crops
168 of paddy (Chennappa *et al.*, 2017a), wheat (Siliniet *al.*, 2012), buckwheat, winter wheat (Roi *et al.*, 2022),
169 and beetroot (Kurdish *et al.*, 2008).

170 The fresh and dry weight of the seedlings indicated that genotypes G₁₀, G₈, and G₇ performed to
171 the best under T₄ conditions. The genotypes that showed a significant improvement in growth, based on
172 fresh weight measurements under T₄ **conditions** were G₁₀ and G₈. In addition, the treatments together with
173 the genotypes significantly affected seedling length; under T₄ conditions, G₈ was the best combination.
174 Vermicompost significantly increases the fresh weight and dry weight of seedlings, especially for
175 tomatoes and pepper plants (Brace, 2017). Riwandiet *al.* (2023) showed that vermicomposting **had** a
176 great influence on both fresh and dry weights in maize seedlings. Shoot fresh weights were increased
177 **by** over 23% for wheat inoculated with Azotobacter strain Azo-8, and increases **by** over 23% in shoot dry
178 weights along with marked improvements in root biomass have also been reported (Singh *et al.*, 2013). In
179 addition to the above, *Vigna radiata* seedlings have shown a 20.07% increase in fresh weight and **a** little
180 over 62% increase in dry weight through Azotobacter inoculation (Munnaza *et al.*, 2012). Scientists show
181 that the addition of vermicompost to growth medium can bring a change in seedling height to cucumbers

182 from 1.9% to about 18.6%, related to leaf area and fresh weight increase (Jankauskienė and others,
183 2022).

184 The vigour indices (I and II) reconfirmed the superiority of G₈, G₁₀, and G₆. Most importantly, G₈ under
185 T₄ recorded the maximum values. By enhancing seed germination, promoting disease resistance, and
186 enhancing the overall health of the plants, vermicompost improves vigour index of crops (Mohite *et al.*,
187 2024). Research conducted by Bajaj (2023) disclosed that the tomato plants' vigour index has increased
188 with altered levels of vermicompost, indicating a positive effect of vermicompost on the crop's growth. The
189 culture filtrate of *Azotobacter salinestrus* (GVT-1) has improved the vigour index of paddy seeds, thus
190 enhancing growth and seedling germination rates in crops (Chennappa *et al.*, 2017b).

191 The proline content was variable from one genotype to another, with a maximum content of G₇ and G₈
192 met under T₄ treatment. This indicates an increased possibility for genotypes to develop physiological
193 resistance toward such stresses. In high correlation, some genotypes recorded very high chlorophyll
194 contents which are vital for photosynthesis, that is, G₁₀, G₈, G₇ under T₄ conditions. These results
195 confirmed those genotypes as most suitable for maximizing treatment benefits towards better
196 physiological output. Mixed inoculation with different *Azotobacter* strains on wheat seedlings has been
197 studied, which increased proline level and growth parameters under osmotic stress, shown to be
198 significantly related to drought resistance (Liu *et al.*, 2013). Various *Azotobacter* strains have
199 been reported to improve the physiological attributes such as proline synthesis in maize grown on saline
200 soils and hence its usefulness toward osmotic adjustment and alleviation of stress in plants (Abdel Latef *et al.*,
201 2020). Research suggests that the addition of vermicompost resulted in an increase in proline
202 concentration, which is the major osmotic regulator that helps plants in overcoming abiotic problems like
203 drought and salinity (Bokobana *et al.*, 2020; Hosseinzadeh *et al.*, 2017). During water stress conditions,
204 30% vermicompost induced a 39% increase in the proline content of chickpea seedlings (Hosseinzadeh *et al.*,
205 2017). In fact, when tomato seedlings are subjected to vermicompost-leachate, especially during heat
206 and moisture stresses, there are increased proline levels (Chinsamy *et al.*, 2014). Researchers have well
207 put vermicompost as an important source of macro- and micronutrients which henceforth augments plant
208 nutrition and at the same time improves chlorophyll concentration, as commonly exhibited by *Capsicum*
209 *annuum* and other vegetable crop seedlings (Kamalkant Yadav *et al.*, 2014; Theunissen, 2010). Kumar *et al.*
210 (2016) observed improvement in the photosynthetic pigments of *Jatropha* by *Azotobacter* and
211 arbuscular mycorrhizal fungus. A treatment of *Azotobacter* in wheat plants indicated a very high increase
212 in the total chlorophyll content (mg g⁻¹) (El-zawawy *et al.*, 2023). The chlorophyll content is increased with
213 the inoculation of *Azotobacter*, either alone or with *Rhizobium*, in black gram (*Vigna mungo*) as compared
214 to the control (Tiwari *et al.*, 2017).

215 The combination of vermicompost with *Azotobacter*, produced positive improvements in crop growth,
216 yield, and nutrient uptake in many crops. This is a blend of the benefits of vermicompost from organic
217 matter and nitrogen-fixing *Azotobacter*, which enhances plant growth parameters in crops such as chili,
218 strawberry, and maize, more than that by either one of the components, or chemical fertilizers alone
219 (Kalpana, 2019; Shirkhani & Nasrolahzadeh, 2016; Tripathi *et al.*, 2015). In *Amaranthus*, this combination of
220 vermicompost and *Azotobacter* was conducive to early emergence and higher germination percentages,
221 which led to the development of more vigorous seedlings (Yadav *et al.*, 2024). Furthermore, the combined
222 application of *Azotobacter* and plant-based composts, such as *Moringa*, has been demonstrated to
223 enhance growth parameters and nutrient levels in various crops, further supporting the synergistic
224 benefits of *Azotobacter* when utilized in conjunction with organic soil amendments (Albureikan, 2024). It
225 also helps in increasing the availability and uptake of essential nutrients such as nitrogen, phosphorus,
226 and potassium in plants and much better nutrient content in crops such as rice and wheat (Ghadimi *et al.*,
227 2021; V. Kumar & Singh, 2001; Rather & Sharma, 2009) reducing the use of chemical fertilizer, being
228 among the sustainable agricultural practices towards environmental sustainability (Rather & Sharma,
229 2009; Shirkhani & Nasrolahzadeh, 2016). The presence of *Azotobacter* in vermicompost, therefore, helps
230 improve the microbial activity of the soil, which translates to a better soil structure and health, thus
231 benefitting long-term crop productivity (Ghadimi *et al.*, 2021; Mal *et al.*, 2021).

232 **4. LIMITATIONS**

233 Although the study establishes the benefits of Azotobacter seed priming and vermicompost
234 integration in Indian mustard cultivation, it is limited because it focused only on seedling stages in a
235 controlled setup. Exactly similar results may not be replicated in a field trial also without exploring long
236 term effects of treatments on seed yield and plant health. So further field trials and cost-benefit analysis
237 are needed.

238 **5. CONCLUSION**

239 The study highlights the importance of combining genotype selection with advanced treatment
240 techniques for improving mustard cultivation. Genotypes TM 306-1(G₈) and TBM 143(G₁₀) are the top
241 performers, especially when paired with treatment Azotobacter primed seeds sown in vermicompost +
242 field soil(T₄), which provides favorable conditions for nutrient uptake, growth, and stress resilience.
243 Farmers can improve germination rates, seedling vigor, and stress tolerance by selecting these
244 genotypes and applying T₄. These genotypes are adaptable to varying conditions, making them ideal for
245 cultivation in diverse agro-climatic regions.

246 **COMPETING INTERESTS**

247 Authors have declared that no competing interests exist.

248 **AUTHORS' CONTRIBUTIONS**

249 Aritra Mukherjee designed the study, performed the statistical analysis, wrote the protocol, and wrote the
250 first draft of the manuscript. Ananya Baidya and Achyuta Basak managed the analyses of the study.
251 Md.Sabir Ahmed Mondol managed the literature search. All authors read and approved the final
252 manuscript.

253

254 **Disclaimer (Artificial intelligence)**

255 **Option 1:**

256 Author(s) hereby declare that NO generative AI technologies such as Large Language Models (ChatGPT,
257 COPILOT, etc.) and text-to-image generators have been used during the writing or editing of this
258 manuscript.

259 **Option 2:**

260 Author(s) hereby declare that generative AI technologies such as Large Language Models, etc. have been
261 used during the writing or editing of manuscripts. This explanation will include the name, version,
262 model, and source of the generative AI technology and as well as all input prompts provided to the
263 generative AI technology

264 Details of the AI usage are given below:

265 1.

266 2.

267 3.

268

269

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