**Assessment of Genetic Variability and Heritability in Indian Mustard (*Brassica juncea* L.) Genotypes**

**ABSTRACT**

The present study was undertaken to assess genetic variability and divergence among 42 genotypes of Indian mustard (*Brassica juncea* L.) under field conditions at the Zonal Agricultural Research Station, Morena, Madhya Pradesh, during 2020–21. Using a randomized block design with three replications, twelve agronomic traits were evaluated, including phenological, morphological, and yield-related characters. Significant genotypic differences were observed for all traits, highlighting substantial variability within the studied genotypes. Notably, genotypic and phenotypic coefficients of variation were moderate to high for key traits such as secondary branches per plant, seed yield per plant, test weight, and siliqua length. High heritability (>60%) coupled with high genetic advance was recorded for several traits, including seed yield per plant (99.85%, 32.61%), indicating the predominance of additive gene action and scope for effective selection. Traits such as plant height, test weight, number of secondary branches, and siliqua length also exhibited promising genetic parameters. These findings suggest that seed yield and its contributing traits could serve as reliable selection indices in mustard breeding programs, providing valuable insights for the development of improved cultivars with higher productivity and desirable agronomic characteristics.

**Keywords:** Genetic variability, Heritability, *Brassica juncea* L., Genotypic diversity, Morphological traits, Genetic advance

**1. INTRODUCTION**

Indian mustard (*Brassica juncea* L.) is a naturally autogamous plant, outcrossing occurs frequently in this crop, ranging from 5 to 30 percent depending on climatic condition and the random diversity of pollinating insects. Indian mustard is an amphidiploid (2n=36) developed from an interspecific cross of *Brassica campestris* (2n=20) and *Brassica nigra* (2n=16), followed by natural chromosomal doubling. The seeds are used as medicine, spices, and as ingredients in salads, juices, curries, and pickles. The mustard plant is primarily cultivated for oil, and the cake produced after oil extraction is frequently utilised for cow feed. Mustard seeds contain 39 to 44 percent oil and 38 to 41 percent protein (Tomar *et al.,* 2015).Indian mustard is a major winter oil seed crop in India. Rapeseed mustard (*Brassica* spp.) is one of the seven edible oil seeds grown in India, accounting for 28.6 percent of total oil seed output. It is India's second most significant edible oilseed after groundnut, accounting for 27.8 percent of the country's oilseed sector. In terms of acreage (17.19 percent), India remains fourth behind Canada, China, and the European Union, and fourth in terms of output after the European Union, Canada, and China (8.54 percent). Rapeseed mustard production has been expanding rapidly in a number of nations throughout the world. The global demand for edible oils and their products is consistently growing. Globally, rapeseed-mustard was cultivated on 36.59 million hectares, producing 72.37 million tonnes with an average productivity of 1980 kg/ha in 2018–19. In India, the area under mustard increased from 5.98 mha in 2017–18 to 6.12 mha in 2018–19, with production rising from 8.43 to 9.26 million tonnes and productivity from 1410 to 1511 kg/ha. In Madhya Pradesh, mustard was grown on 0.78 mha, yielding 1.11 million tonnes at a productivity of 1422 kg/ha(Anonymous, 2019). Despite available yield-enhancing technologies, the area under cultivation remains static.

Understanding the factors responsible for high yield is challenging, as yield is a complex trait influenced by multiple interacting components. To enhance yield effectively, breeders must break down this complexity by focusing on individual yield components, even when they are negatively correlated. This requires a clear understanding of the nature and strength of relationships among these components, as well as insight into how both qualitative and quantitative traits contribute to overall yieldParoda and Joshi (1970)and Lush (1949).The research of genetic progress is also essential since it measures genetic gain based on selection in a certain character. As a result, the study of genetic variability and heritability, as well as genetic advance, is required for every crop development programme based on selection and hence, the present study was conducted with the objective to assess the Genetic Variability and Heritability in different genotypes of Indian Mustard (*Brassica juncea* L.).

**2. MATERIALS AND METHODS**

The present investigation was carried in the assigned experimental field of Mustard at Zonal Agriculture Research Station, Morena, Madhya Pradesh, during 2020-21. A total of 42 mustard genotypes were evaluated using a Randomized Block Design (RBD) with three replications.All the genotypes were raised with good plant health by following recommended package of practices. The net plot area of the replicated plots was 0.6 m x 4.0 m maintaining the spacing of 30 cm (row to row) and 10 cm (plant to plant).

**2.1 Observations recorded:**

Evaluation of genetic variability and divergence of different traits of mustardwas carried out.Observations were recorded on the ten plants of each of the plots for twelve characters *viz.,* days to 50% flowering, days to maturity, plant hight, number of primary branches per plant, number of secondary branches per plant, length of main raceme, number of siliquae on main raceme, number of seeds per siliquae, siliquae length, test weight, biological yield and seed yield per plant were worked out and the data was tabulated.

**2.2 Statistical analysis:**

The mean values of varieties in each replication were used for statistical analysis. The data were analysed for a randomized block design to test the significance of differences between the genotypes for various characters by calculating the Critical Difference (C.D.) and Standard Error of means (S.E.m) as described by Panse and Sukhatme (1967). Further, Genotypic, Phenotypic and Error variance were calculated as per the given formulae:

Genotypic variance (σ²g) = (MSt/MSe)/r

Phenotypic variance (σ²p) = σ²g + σ²e Error variance (σ²e) = MSe

Where,MSe = Mean sum square of error; MSt = Mean sum square of treatments; r = No. of replications.

For further calculations, Heritability in broad sense h² (b) was computed as a ratio of genotypic variance to phenotypic variance as described by (Allard, 1960).

h2 (b) = x 100

Where,h² (b) = Heritability in broad sense; σ²g = Genotypic variance; σ²p = Phenotypic variance

The expected Genetic Advance (G.A.) under selection for the different characters was estimated as suggested by Allard (1960).

G.A. = h² (b) × σp × k

Where, h² (b) = Heritability in broad sense; σp = Phenotypic standard deviation (); k = Intensity of selection, the value of which is 2.06 when 5 percent of the individual are selected from the population as given by Lush (1949).

Genetic advance as percent of means for each character was calculated as suggested by Johnson *et al.,* (1955).

G.A.(%) = x 100

**3. RESULTS AND DISSCUSION**

The mean performance of forty-two mustard varieties observed for twelve different characters studied along with their range and critical differences are presented in Table 1.The present investigation of 42 mustard varieties revealed significant genotypic variation across twelve agronomic traits. Days to 50% flowering varied from 42.48 (L-4) to 62.79 (Varuna), and days to maturity ranged from 120.67 (PM-25) to 141.33 (RVM-1). Plant height ranged between 158.14 cm (PM-27) and 225.89 cm (PDZM-31), while the number of primary and secondary branches per plant ranged from 3.74 (RVM-3) to 7.42 (Albeli) and 4.47 (RGN-73) to 15.37 (RH-749), respectively. The main raceme length varied from 55.59 cm (Pusa Jagannath) to 74.56 cm (L-4), and siliquae per raceme ranged from 46.56 (RVM-12-1-18) to 56.15 (L-4). Seeds per siliqua ranged from 12.91 (Maya) to 17.59 (RGN-73), with siliqua length ranging from 4.25 cm (L-4) to 6.49 cm (Albeli). Test weight ranged from 3.66 g (RMM-12-2-18) to 7.24 g (RH-749). Biological yield ranged from 62.23 g (RMM-12-1-18) to 71.43 g (Krishna), while seed yield per plant varied from 14.42 g (PM-27) to 28.31 g (Pusa Bold). These results underscore considerable genetic variability among the mustard genotypes, offering valuable insights for future selection and improvement programs.High amount of genetic variability for many of these traits has also been reported earlier by Uddin *et al.* (1995) revealed high genotypic and phenotypic coefficients of variation for seed weight, seeds per siliquae, biological yield, oil content, secondary branches per plant and seed yield per plant. Yadav *et al.* (2011), Singh *et al.* (2013) and Shekhawat *et al.* (2014) also reported the similar results as described in the results.

Results from the present study (Table2) in this context indicated that no genotypic coefficient of variance (GCV) and phenotypic coefficient of variance (PCV) were high (>25%). Further GCV and PCV were moderate (10-25%) for number of secondary branches per plant (23.709 and 23.758), seed yield per plant (15.846 and 15.857), test weight (15.539 and 15.635), number of primary branches per plant (14.273 and 14.433), siliqua length (11.198 and 11.582), and low (<10%) for days to 50% flowering (8.688 and 8.738), plant height (7.723 and 7.726), number of seeds per siliqua (6.846 and 6.991), length of main raceme (6.259 and 6.266), number of siliqua on main raceme (4.531 and 4.537), biological yield (3.977 and 4.011) and number of maturity days (3.884 and 3.910). Further, the present findings showed that estimates of PCV were generally higher than their corresponding GCV for all the character studied. Furthermore, the current data revealed that PCV estimations were significantly greater than the corresponding GCV for all of the characters tested. These findings are similar in agreement with earlier reported by Kumari and Kumari (2018) reported highly significant genotypic differences were found in the combined analysis of variance for number of days to flower initiation, crop plant height, number of days to 50 per cent flowering, length of the siliquae, number of days to 75% maturity, test weight and harvest index, to authenticate the probability of building up these characters through breeding program. Yadav *et al*. (2012), Yohannes and Belete (2013), Singh *et al*. (2013), Shekhawat *et al*. (2014) and Bibi *et al*. (2016) also reported the similar results as described in the results.

Most of the character under studied observed high heritability (>60%) *viz.,* for plant height (99.93%), seed yield per plant (99.85%), length of main raceme (99.78%), number of siliquae on main raceme (99.71%), number of secondary branches per plant (99.59%), days to 50% flowering (98.85%), test weight (98.77%), number of seed per siliquae (98.69%), number of maturity days (98.64%), biological yield (98.31%), number of primary branches per plant (97.79%) and siliqua length (93.48%)(Table2).In line with the present results,Singh *et al*. (2011) also reported the relationship between heritability and GA. Goyat*et al.* (2012) and Singh *et al.* (2013) described this interaction in relation to Indian mustard. Shekhawat *et al.* (2014) also reported the similar results as described in the results. Gadi *et al*. (2020) investigated how variation, heritability in a wide sense, and anticipated genetic progress in germplasm will aid in determining the genotype's true potential. Hence, the current investigation was intended to determine the variability, heritability in broad sense and estimated genetic advance for crop output and other crop traits among the varieties. The high heritability indicates a significant share of genetic factors in the determination of these characteristics, which may be used to improve seed production. Further studies conducted at various parts by Upadhyay and Kumar (2009), Sohanram and Verma (2010).

Expected genetic advance expressed as percentage of mean (Table2)was exhibited high (>20%) for number of secondary branches per plant (48.73), seed yield per plant (32.61), test weight (31.81), number of primary branches per plant (29.07) and siliquae length (22.30). Moderate GA (10-20%) was observed for days to 50% flowering (17.79), plant height (15.90), number of seed per siliquae (14.00) and length of main raceme (12.87). Low GA (<10%) was noted for number of siliquae on main raceme (9.31), biological yield (8.12) and number of maturity days (7.94). In the present investigation, high heritability coupled with high genetic advance observed for plant height, days to maturity, length of main raceme, number of siliquae on main raceme and seed yield per plant. Sarvesh *et al.* (2018) showed moderate to high level of genotypic coefficient of variance (GCV) and phenotypic coefficient of variance (PCV). In general, phenotypic coefficient of variance was found to be higher than their genotypic coefficient of variance but the extent was quite small, indicating very less environmental influence on the expression of the characters. Genotypic coefficient of variation (GCV) was found to be higher for 1000-seed weight (24.33), biological yield per plant (22.47), secondary branches plant (22.41), seed yield per plant (19.75). Heritability and genetic advance are important selection parameters. Gadi *et al.* (2020) reported that variability in different varieties, heritability within the genotypes and estimated genetic advance in germplasm will help to investigate the actual potential of the varieties. Singh *et al.* (2012), Yohannes and Belete (2013) and Shekhawat *et al.* (2014) also reported the similar results as described in the results.

**4. CONCLUSION**

Genetic variability of the forty-two varieties among the twelve different characters suggests that all characters are *viz.,* days to 50% flowering, number of maturity days, height of plant, primary branches plant-1, secondary branches plant-1, main raceme length, siliquae on major raceme, seeds siliquae-1, length of siliquae, test weight, biological yield per plant and seed yield per plant can be relied upon direct selection for further improvement.High heritability coupled with high genetic advance were observed for the characters under study revealed that the characters undergo the influence of additive gene action. Seed yield per plant exhibiting significant stable and positive correlation with the characters. Biological yield per plant displayed high order of direct effect on seed yield per plant. All these results indicates that they can be considered as guidelines for further selection programmes.

**5. REFERENCES**

Allard, R.W. (1960). Principles of Plant Breeding Edn. First. John Wiley and Sons Inc., New York. Pp. 115- 128.

Anonymous. 2019. Agricultural Statistics at a Glance. <https://eands.dacnet.nic.in/PDF/At%20a%20Glance%202019%20Eng.pdf> (accessed on 08/12/2021).

Bibi T, Rauf S, Mahmood T, Haider Z, Salah D (2016). Genetic Variability and Heritability Studies in Relation to Seed Yield and its Component Traits in Mustard (*Brassica Juncea* L.). Acad. J. Agric. Res. 4(8): 478-482.

Gadi. J., Ranjan, N,, Chakraborty and Imam, Z., (2020). To study the genetic variability, heritability and genetic advance for different quantitative characters in Indian mustard (*Brassica juncea* L. Czern&Coss). *Int. J. Curr. Microbiol. App. Sci.* 9(10): 1557-1563.

Goyat,B.;Singh,D.;Avatar,R.andSingh,A. (2012). EstimationofselectionparametersintheelitegenepoolofIndian mustard[*Brassicajuncea*(L.) Czern&Coss.]. *Res.on Crops*, 13(3): 1098-1101.

Johnson, H.W.; H. F. Robinson and R.E. Comstock (1955). Estimates of genetics and environmental variability in soybean. *Journal of Agronomy,* 47: 314-318.

Kumari, A., Kumari, V., (2018). Studies on genetic diversity in Indian mustard (*Brassica Juncea*Czern&Coss) for morphological characters under changed climate in the mid-hills of Himalayas. *Pharma Innovation*;**7**(7):290-296.

Lush, J.L. (1949). Animal breeding plants (2nd Ed). IOWA State college press, Ames*,* IOWA.

Panse, V.G. and Sukhatme, P.V. (1967). Statistical Method for Agricultural Workers 2nd Ed. ICAR, New Delhi: 381

Paroda, R.S. and Joshi, A.B. (1970). Genetic architecture of yield and components of yield in wheat. *Indian J. Genet.,* 30: 289-314.

Sarvesh Kr. Maurya, K.N. Maurya, Kanhaiya Lal, Yogendra Singh, Sujit Singh, Brijesh Dixit and Soni Singh. (2018). Assessment of Genetic Variability, Heritability and Genetic Advance in Indian Mustard [*Brassica juncea* L. Czern&Coss.]. *Int. J. Curr. Microbiol. App. Sci.* 7(11): 13- 18

Shekhawat, N. Jadeja, G.C. and Singh, J. (2014). Genetic variability for yield and itscomponentsinIndian mustard(*Brassicajuncea*L.Czern&Coss).*Electronic JournalofPlantBreeding,* 5(1):117-119.

Singh, A. Avtar, R. Singh, D. Sangwan, O. and Balyan, P. (2013). Geneticvariability, character association and path analysis for seed yield and componenttraitsundertwoenvironmentsin Indianmustard.*J.ofOilseedBrassica,* 4(1):43-48.

Singh, A. Ram Krishna. Singh, L. and Ram Jeet (2011). Heritability and genetic advance in Indian mustard (*Brassica juncea* L. Czern&Coss.). *Curr. Adva. Agric. Sci.*, 3(1): 52-53.

Singh, D.K.; Kumar, K. and Singh, P. (2012). Heterosis and heritability analysis fordifferentcrossesin*Brassicajuncea*withinheritanceofwhiterustresistance. *Journalof Oilseed,* 3(1): 18-26.

Sohan ram and Verma, N. (2010). Genetic variability for yield and yield components in Indian mustard (*Brassica juncea*). *J. of Oilseeds Research*, 27(2): 170-171.

Tomar,A.;Singh,M.andSingh,S.K. (2015). Geneticanalysisofyieldanditscomponentsbasedonheteroticresponseandcombiningabilityparametersin Indianmustard(*Brassicajuncea*L.CzernandCoss). *ProgressiveAgriculture* 15(1):85-91.

Uddin, M.J. Chaudhary, M.A.Z. and Mai, M.F.U. (1995). Genetic variability, character association and path analysis in Indian mustard *(Brassica juncea*L.). *Ann. Bangladesh Agric.,* 5(1): 51-54.

Upadhyay, D.K. and Kumar, K. (2009). Analysis of heritability and genetic advance inrelation to yield and its components in Indian mustard (*Brassica juncea* L.Czern and Coss) under normal and late sown conditions. *Inter. J.Pl. Sci*., 4(1):12-14.

Yadav, D.K. Giri, S.C.; Vignesh, M.; Sujata Vasudev; Yadav, A.K.; Dass, B.; Singh, R; Singh, N.; Mohapatra, T. and Prabhu, K.V. (2011). Genetic variability and trait association studies in Indian mustard (*Brassica juncea*). *Indian J. of Agric. Sci.,* 81(8): 712-716.

Yadav, R.K. Chakravarty, A. Kumar, N. Yadav, J.R. and Ram Krishna (2012). Genetic variability and component analysis of some metric traits under sodic soil condition in Indian mustard (*Brassica juncea* L. Czern and Coss.). *Progressive Agriculture*, 12(2): 396-401.

Yohannes, M.T.W. and Belete, Y.S. (2013). Genetic variability of Ethiopian mustard(*Brassicacarinata*A.Brun) accessions basedon some morphological characters.International*J.ofPlantBreedingandGenetics,* 7(3):169-175.

**Table 1: Mean performance of forty-two mustard genotypes for twelve characters.**

|  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| **S. No.** | **Genotypes** | **Days to 50% Flowering** | **Days to maturity** | **Plant height (cm)** | **Number of primary branches/ plant** | **Number of secondary branches/ plant** | **Length of main raceme (cm)** | **Number of siliquae on main raceme** | **Number of seed per siliquae** | **Siliquae length (cm)** | **Test weight (g)** | **Biological yield** | **Seed yield per plant (g)** |
| 1 | **Pusa Bold** | 54.820 | 123.333 | 186.130 | 5.920 | 13.350 | 59.230 | 51.120 | 14.340 | 4.340 | 4.530 | 68.520 | 28.360 |
| 2 | **Varuna** | 62.787 | 125.000 | 208.610 | 5.640 | 8.540 | 63.560 | 53.230 | 15.420 | 5.700 | 5.030 | 64.530 | 25.060 |
| 3 | **Rohini** | 51.050 | 121.333 | 204.970 | 4.780 | 7.230 | 55.850 | 51.430 | 14.520 | 6.480 | 4.650 | 63.450 | 17.900 |
| 4 | **Kranti** | 47.350 | 125.333 | 215.390 | 5.690 | 13.350 | 58.740 | 50.240 | 15.440 | 5.270 | 4.220 | 64.230 | 15.450 |
| 5 | **Krishna** | 53.550 | 135.000 | 213.770 | 6.710 | 14.650 | 63.530 | 52.330 | 15.630 | 4.860 | 4.020 | 71.430 | 23.700 |
| 6 | **Maya** | 48.907 | 132.667 | 210.387 | 6.267 | 9.910 | 64.580 | 52.920 | 12.907 | 4.340 | 6.163 | 66.580 | 24.260 |
| 7 | **Vardan** | 47.653 | 130.000 | 208.640 | 6.463 | 11.687 | 72.433 | 55.300 | 15.557 | 5.273 | 5.430 | 65.247 | 26.303 |
| 8 | **Vasundhara** | 47.467 | 134.333 | 216.187 | 5.420 | 10.613 | 68.263 | 52.303 | 14.440 | 4.417 | 4.340 | 68.593 | 21.427 |
| 9 | **Swarnjyoti** | 45.500 | 130.667 | 194.077 | 4.883 | 8.770 | 63.477 | 50.693 | 17.270 | 4.597 | 6.230 | 65.010 | 19.310 |
| 10 | **Pusa Jagannath** | 53.533 | 132.000 | 209.540 | 5.410 | 7.650 | 55.587 | 47.687 | 16.383 | 5.633 | 6.233 | 64.857 | 18.743 |
| 11 | **Pusa Mahak** | 52.517 | 130.000 | 212.773 | 5.587 | 11.480 | 60.617 | 52.370 | 14.127 | 5.540 | 4.820 | 63.627 | 20.287 |
| 12 | **Pusa Jaikisan** | 50.630 | 126.000 | 217.960 | 5.680 | 14.767 | 62.117 | 51.440 | 15.400 | 4.433 | 6.250 | 64.560 | 21.640 |
| 13 | **Albeli** | 49.860 | 123.333 | 191.870 | 7.417 | 11.797 | 59.823 | 51.553 | 15.280 | 6.493 | 4.313 | 68.267 | 21.440 |
| 14 | **Sej 2** | 49.837 | 130.667 | 217.880 | 5.303 | 11.567 | 63.767 | 51.113 | 15.323 | 5.617 | 4.693 | 71.070 | 21.350 |
| 15 | **Shraddha** | 54.227 | 135.667 | 192.280 | 6.543 | 14.610 | 69.797 | 53.330 | 16.297 | 4.490 | 5.460 | 68.420 | 26.607 |
| 16 | **L-4** | 42.483 | 140.000 | 217.470 | 5.787 | 14.877 | 74.557 | 56.153 | 16.273 | 4.250 | 7.050 | 64.023 | 27.507 |
| 17 | **L-6** | 49.597 | 125.000 | 207.677 | 5.620 | 14.670 | 68.653 | 53.127 | 14.617 | 4.727 | 6.233 | 68.953 | 24.310 |
| 18 | **JMWR-908-1** | 43.743 | 130.333 | 212.837 | 5.697 | 14.647 | 60.427 | 52.417 | 15.643 | 5.613 | 6.227 | 68.423 | 23.097 |
| 19 | **RGN-73** | 43.993 | 129.333 | 210.970 | 5.723 | 4.473 | 62.510 | 53.497 | 17.593 | 4.830 | 5.457 | 68.550 | 23.373 |
| 20 | **NRCHB-101** | 47.477 | 125.000 | 225.390 | 5.450 | 10.633 | 58.483 | 51.377 | 14.293 | 4.297 | 5.650 | 62.960 | 21.493 |
| 21 | **RVM-3** | 48.790 | 129.333 | 205.750 | 3.740 | 7.163 | 61.457 | 51.307 | 15.230 | 4.530 | 5.243 | 63.440 | 20.683 |
| 22 | **RH-406** | 43.977 | 132.000 | 168.397 | 4.387 | 6.640 | 62.667 | 53.423 | 16.337 | 5.690 | 5.693 | 62.413 | 23.483 |
| 23 | **RH-749** | 48.007 | 133.667 | 202.813 | 5.440 | 15.367 | 60.717 | 50.407 | 16.410 | 5.293 | 7.237 | 63.217 | 20.403 |
| 24 | **Giriraj** | 49.033 | 131.333 | 218.663 | 5.563 | 11.450 | 60.297 | 50.537 | 16.303 | 4.517 | 5.170 | 64.700 | 19.517 |
| 25 | **JM-1** | 45.290 | 131.000 | 202.517 | 5.373 | 9.620 | 63.733 | 53.337 | 14.423 | 5.150 | 4.657 | 68.653 | 24.503 |
| 26 | **JM-2** | 44.727 | 134.000 | 185.947 | 7.330 | 13.250 | 62.403 | 52.247 | 15.423 | 5.503 | 5.347 | 69.687 | 22.490 |
| 27 | **JM-3** | 44.907 | 125.333 | 199.713 | 5.313 | 11.433 | 64.280 | 53.177 | 14.323 | 5.123 | 5.967 | 62.693 | 23.613 |
| 28 | **RVM-1** | 48.160 | 141.333 | 205.530 | 5.293 | 9.593 | 62.417 | 50.230 | 15.393 | 5.417 | 6.250 | 69.997 | 20.580 |
| 29 | **RVM-2** | 44.207 | 140.667 | 166.357 | 5.427 | 14.940 | 63.570 | 54.407 | 13.707 | 5.333 | 5.323 | 65.233 | 25.577 |
| 30 | **PM-25** | 45.833 | 120.667 | 208.797 | 6.393 | 9.567 | 64.020 | 54.367 | 15.437 | 5.670 | 5.430 | 67.730 | 24.530 |
| 31 | **PM-26** | 49.017 | 132.000 | 196.570 | 5.590 | 14.747 | 61.690 | 52.313 | 17.260 | 5.383 | 5.340 | 69.497 | 22.427 |
| 32 | **PM-27** | 43.700 | 131.333 | 158.143 | 4.353 | 9.517 | 59.313 | 46.993 | 16.367 | 4.490 | 5.003 | 68.500 | 14.417 |
| 33 | **PM-28** | 48.317 | 137.333 | 186.113 | 4.597 | 8.437 | 60.263 | 52.777 | 16.183 | 4.510 | 5.177 | 69.710 | 18.330 |
| 34 | **PM-30** | 45.237 | 131.000 | 209.280 | 6.583 | 10.743 | 58.680 | 48.353 | 16.327 | 4.637 | 5.243 | 65.683 | 17.350 |
| 35 | **Pusa Vijay** | 46.523 | 129.000 | 210.500 | 5.720 | 10.530 | 58.660 | 47.187 | 14.433 | 5.683 | 5.120 | 65.643 | 14.493 |
| 36 | **PDZM-31** | 48.543 | 133.000 | 225.893 | 5.660 | 12.700 | 59.447 | 48.337 | 14.337 | 5.313 | 4.357 | 62.983 | 16.333 |
| 37 | **JMM-927** | 43.770 | 132.000 | 205.203 | 5.560 | 13.883 | 65.290 | 55.290 | 14.220 | 4.643 | 5.817 | 67.317 | 24.303 |
| 38 | **JMM-991** | 56.760 | 135.000 | 220.130 | 5.230 | 12.750 | 64.360 | 50.270 | 15.360 | 5.427 | 4.030 | 68.560 | 20.520 |
| 39 | **RMM-10-1-1** | 55.720 | 134.333 | 203.190 | 4.430 | 14.030 | 59.630 | 48.330 | 16.220 | 5.480 | 4.520 | 65.330 | 26.650 |
| 40 | **RMM-12-1-18** | 52.010 | 122.333 | 173.810 | 4.260 | 11.530 | 62.940 | 46.560 | 14.720 | 4.590 | 4.240 | 62.230 | 24.760 |
| 41 | **RMM-12-2-18** | 50.280 | 126.333 | 217.650 | 4.520 | 10.550 | 64.960 | 53.480 | 15.240 | 4.280 | 3.660 | 64.230 | 18.460 |
| 42 | **RMM-12-3-18** | 53.140 | 125.000 | 197.550 | 5.450 | 9.560 | 65.760 | 54.670 | 17.560 | 4.840 | 5.160 | 63.180 | 21.330 |
| **Mean** | | **48.879** | **130.310** | **203.412** | **5.529** | **11.364** | **62.680** | **51.705** | **15.428** | **5.064** | **5.262** | **66.236** | **21.818** |
| **Range** | | **42.483-62.787** | **120.66-141.333** | **158.143- 225.893** | **3.740-7.417** | **4.473-15.367** | **55.587-74.557** | **46.560-56.153** | **12.907- 17.593** | **4.250-6.493** | **3.660- 7.237** | **62.230- 71.430** | **14.417- 28.360** |
| **SEm ±** | | **0.456** | **0.595** | **0.415** | **0.118** | **0.173** | **0.181** | **0.123** | **0.122** | **0.150** | **0.091** | **0.345** | **0.132** |
| **C.D. at 5%** | | **1.286** | **1.673** | **1.188** | **0.332** | **0.487** | **0.514** | **0.354** | **0.342** | **0.421** | **0.256** | **0.972** | **0.371** |

**Table 2: Estimates of variability parameters for twelve characters in mustard (*Brassica juncea*) varieties.**

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Sr. No.** | **Characters** | **GCV (%)** | **PCV (%)** | **Heritability (%) (Broad sense)** | **Genetic Advance** | **Genetic Advance**  **(% of Mean)** |
| **1** | Days to 50% Flowering | 8.688 | 8.738 | 98.85 | 8.697 | 17.79 |
| **2** | Days to maturity | 3.884 | 3.910 | 98.64 | 10.353 | 7.94 |
| **3** | Plant height (cm) | 7.723 | 7.726 | 99.93 | 32.350 | 15.90 |
| **4** | Number of primary branches/ plants | 14.273 | 14.433 | 97.79 | 1.600 | 29.07 |
| **5** | Number of secondary branches/ plants | 23.709 | 23.758 | 99.59 | 5.538 | 48.73 |
| **6** | Length of main raceme (cm) | 6.259 | 6.266 | 99.78 | 8.072 | 12.87 |
| **7** | Number of siliquae on main raceme | 4.531 | 4.537 | 99.71 | 4.818 | 9.31 |
| **8** | Number of seed per siliquae | 6.846 | 6.891 | 98.69 | 2.161 | 14.00 |
| **9** | Siliquae length (cm) | 11.198 | 11.582 | 93.48 | 1.129 | 22.30 |
| **10** | Test weight (g) | 15.539 | 15.635 | 98.77 | 1.673 | 31.81 |
| **11** | Biological yield | 3.977 | 4.011 | 98.31 | 5.380 | 8.12 |
| **12** | Seed yield per plant (g) | 15.846 | 15.857 | 99.85 | 7.116 | 32.61 |