

Original Research Article

Genetic variability, correlation and path analysis in F₃ and F₄ generation of bottle gourd [*Lagenaria siceraria* (Mol.) Standl.]

ABSTRACT

The bottle gourd is an important summer vegetable in India, but its productivity is hindered by biotic and abiotic stresses, challenging growing conditions and the lack of high-yielding varieties. Therefore, it is essential to understand its genetics to develop high-yielding cultivars. A thorough understanding of genotypic and phenotypic variability, heritability and genetic advance is necessary for a successful crop improvement program. Research conducted during the spring-summer and rainy seasons of 2021–22 at CCS HAU, Hisar, on the F₃ and F₄ generations of the bottle gourd cross GH-28 × Pusa Santushti, showed variability in various yield and yield-contributing traits. The fruit yield per hectare showed high values for genotypic coefficient of variation, phenotypic coefficient of variation, heritability and genetic advance in both generations, with progeny 4 performing best in yield-related traits. The observed changes in genetic variability across generations suggested heterozygosity, which may stabilize once homozygosity is achieved in later generations. Additionally, correlation and path analysis indicated that the number of primary branches, days to first harvest, average fruit weight and fruit length directly affected fruit yield per hectare. Thus, selecting for these traits could significantly enhance a breeding program's success.

Keywords: bottle gourd, heritability, progenies, segregating generation, variability

1. INTRODUCTION

Bottle gourd, an important vegetable in the Cucurbitaceae family, is extensively grown in India during the summer for its nutritious fruits. With a water content of 96.3 percent (Thamburaj and Singh, 2015), it has a cooling effect, enhancing its popularity in the country. Beyond its culinary uses, the pulp of the fruit provides therapeutic benefits for issues like constipation, biliousness, and indigestion. The monoecious nature of plant, which encourages cross-pollination while avoiding inbreeding depression, can be exploited to achieve stable homozygosity for desired traits. Sirohi and Sivakami (1991) have documented the variation in bottle gourd fruit characteristics, and utilizing this diversity is essential for developing cultivars that offer early fruiting, a higher number of female flowers, more primary branches, greater yields, and improved fruit shape and size. The main goal of any breeding program is to enhance crop yield and productivity.

Direct selection may not be effective for complex traits like fruit yield, which are controlled by multiple genes and influenced by environmental factors. Therefore, analyzing variability parameters is crucial for understanding the inheritance of traits and enabling effective selection. Burton (1952) suggested that genetic variation combined with heritability estimates can predict genetic advance through selection, a concept supported by Johnson *et al.* (1955). Heritability and genetic advance help determine the potential for improvement post-selection and the environmental influence on trait expression (Robinson *et al.*, 1949).

Moreover, examining correlation coefficients and the magnitude of directional influence among various fruit yield and yield-contributing traits is important for selecting specific traits. Genetic studies in segregating generations offer insights into traits valuable for breeding programs. These generations introduce variation due to random allele recombination, promoting heterozygosity for these traits. With this in mind, the current study aims to investigate genetic variability parameters, correlation coefficients, and path analysis within the F₃ and F₄ generations of bottle gourd.

2. MATERIALS AND METHODS

2.1 Experimental details and observations recorded

The research took place in the Department of Vegetable Science at CCS Haryana Agricultural University, Hisar. The experimental materials included bottle gourd cross between GH-28 and Pusa Santushti, set up in a Randomized Block Design with three replications. Five progenies from each of the F₃ and F₄ generations of the cross were cultivated during the rainy season of 2021 and the spring-summer season of 2022, respectively. Each progeny consisted of ten plants, amounting to a total of 200 plants per generation per replication. Standard agronomic practices were maintained throughout the crop period. Observations on 17 quantitative traits were made on five randomly selected plants in each generation and analyzed statistically.

2.2 Statistical Analysis

The observed data for various traits were statistically analyzed to determine the genetic variability parameters as proposed by Panse and Sukhatme (1985). The phenotypic coefficient of variance (PCV) and genotypic coefficient of variance (GCV) were estimated according to Burton and Devane (1952). Heritability was calculated based on the method by Lush (1940), and genetic advance was determined as described by Johnson *et al.* (1955). Correlation coefficients were calculated using the approach by Al-Jibouri *et al.* (1958), and path coefficients were computed as explained by Dewey and Lu (1959).

2.3 Categorization of variability parameters

GCV and PCV was classified as low (0 -10%), moderate (10 - 20%) and high (>20%) as suggested by Shivasubramanian and Madhavamenonenon (1973). Johnson *et al.* (1955) categorized heritability in broad sense values as low (Less than 50 %), moderate (50 - 75 %) and high (More than 75 %). Johnson *et al.* (1955) categorized the range of genetic advance as per cent of mean values as low (Less than 10 %), moderate (10 -20%) and high (More than 20 %).

3. RESULTS AND DISCUSSION

The examination of variability indices, such as the genotypic coefficient of variation (GCV), phenotypic coefficient of variation (PCV), heritability, and genetic advance as a percent of the mean, revealed the presence of variability in the F₃ and F₄ generations of the GH-28 × Pusa Santushti cross. Comparable mean and range values for corresponding traits were observed in these generations (Table 1). However, there was a decline in traits contributing to fruit yield, such as average fruit weight, number of fruits per vine, fruit yield per plot, and fruit yield per hectare from the F₃ to F₄ generation. This decline can be attributed to seasonal fluctuations, with higher summer temperatures adversely affecting flowering and fruit behaviour in the F₄ generation. Similar ranges for several traits in bottle gourd segregating generations were reported by Vaidya (2018) and Chandramouli *et al.* (2021).

3.1 Phenotypic coefficient of variance (PCV) and genotypic coefficient of variance (GCV)

For ease of understanding, coefficients of variation are categorized as genotypic and phenotypic. The minor discrepancy between phenotypic and genotypic coefficients of variation in this study is due to a minimal environmental impact, indicating that the phenotype closely represents the genotype. Higher GCV and PCV values (Figure 1) were observed for fruit length (22.53% and 23.23%, respectively), average fruit weight (20.09% and 22.79%, respectively), fruit yield per plot (28.75% and 33.45%, respectively), and fruit yield per hectare (28.75% and 33.45%, respectively) in the F₃ generation. In the F₄ generation, higher values were reported for fruit yield per plot (24.63% and 29.42%, respectively) and fruit yield per hectare (24.63% and 29.42%, respectively). These findings align with results from Alekar *et al.* (2019) in the F₄ population of bitter gourd and Kanimozhi *et al.* (2015) in wax gourd. Due to the significant variability caused by heterozygosity, these traits can be directly exploited in improvement programs through simple selection. The presence of moderate variability indicates that selection for these traits is feasible to some extent for enhancement. Similar findings were reported by Chandramouli *et al.* (2021) in the F₂ generation of bottle gourd and Kumar *et al.* (2011) in the F₂ generation of cucumber.

3.2 Heritability and genetic advance as per cent of mean

Understanding heritability is crucial to determine whether phenotypic differences among individuals are due to genetic differences or environmental factors. In the F₃ and F₄ generations (Figure 2), the average fruit weight demonstrated high heritability estimates of 77.67% and 74.02%, respectively, with substantial genetic advance as a percentage of the mean (36.47% and 32.09%, respectively). For fruit length, heritability estimates were exceptionally high at 94.06% and 90.51%, respectively, with significant genetic advances of 45.00% and 31.53% in the F₃ and F₄ generations. Fruit yield per hectare also showed high heritability estimates of 73.89% and 70.10%, along with considerable genetic advance percentages of 50.91% and 42.49% in F₃ and F₄ generations, respectively. Vine length had heritability values of 84.13% and 94.91%, with high genetic advances (34.34% and 32.51%) in the respective generations. Comparable results were reported by Alekar *et al.* (2019) in the F₄ generation of bitter gourd and by Chinthalapudi *et al.* (2021) in the F₃ generation of ridge gourd. Kumar *et al.* (2011) also observed similar findings in the F₂ population of cucumber regarding flowering traits. Direct selection for these traits is feasible due to their control by additive genes with minimal environmental influence. Traits such as fruit diameter, which exhibit high heritability but moderate genetic advance, suggest limited potential for further improvement due to the

presence of both additive and non-additive gene activity. Pradhan *et al.* (2021) reported similar findings for fruit traits in the F₄ generation of bitter gourd, while Sravani *et al.* (2021) and Kannan *et al.* (2019) found comparable heritability values in the F₂ and F₄ generations of ridge gourd, respectively.

Table 1 Estimation of mean values (GH-28 × Pusa Santushti, F₃ and F₄ generations)

Trait	Mean		Minimum		Maximum	
	F ₃	F ₄	F ₃	F ₄	F ₃	F ₄
DFG	8.40	8.53	7.33	6.67	9.67	9.67
NPB	15.38	14.30	13.00	11.27	17.57	17.13
DMF	43.12	43.43	41.00	41.00	46.00	46.00
DFF	46.96	46.87	43.00	42.00	50.00	50.00
NMF	7.62	6.91	6.83	5.61	8.78	8.11
NFF	11.54	11.15	9.94	9.33	12.79	12.55
LL	20.35	20.33	18.15	18.03	22.96	22.50
LW	25.18	27.06	22.85	24.11	28.85	29.55
DFH	58.67	56.33	56.00	52.00	61.00	60.00
FL	31.98	34.58	25.73	28.33	44.33	41.17
FD	7.28	6.75	5.93	5.48	8.63	8.13
VL	6.38	7.11	5.04	5.65	7.91	8.43
AFW	653.87	628.47	531.33	514.33	882.00	832.67
SW	18.26	17.87	16.57	15.74	21.52	21.36
FPV	6.46	5.81	5.67	4.83	7.20	7.07
FYP	42.91	36.71	32.11	29.02	63.65	50.37
FYH	257.47	220.28	192.68	174.14	381.90	302.20

DFG: Days to 50% germination;
NPB: Number of

primary branches ; DMF: Days to first male flower opening; DFF: Days to first female flower opening; NMF: Node to first male flower; NFF: Node to first female flower; LL: Leaf length (cm); LW: Leaf width (cm); DFH: Days to first harvest; FL: Fruit length (cm); FD: Fruit diameter (cm); VL: Vine length at the time of final harvest (m); AFW: average fruit weight (g); SW: 100 seed weight (g); FPV: Number of fruits per vine; FYP: Fruit yield per plot (kg); FYH: fruit yield per hectare (q/ha).

The variations in GCV, PCV, and heritability from the F₃ to F₄ generations showed no clear pattern. The increase in GCV, PCV, and heritability values from F₃ to F₄ suggests increasing homozygosity, whereas selection in later generations is necessary to establish homozygosity in traits with decreasing values. Similar trends were observed in faba beans by Ahmad (2016) and in ridge gourd by Suresh and Balamohan (2018).

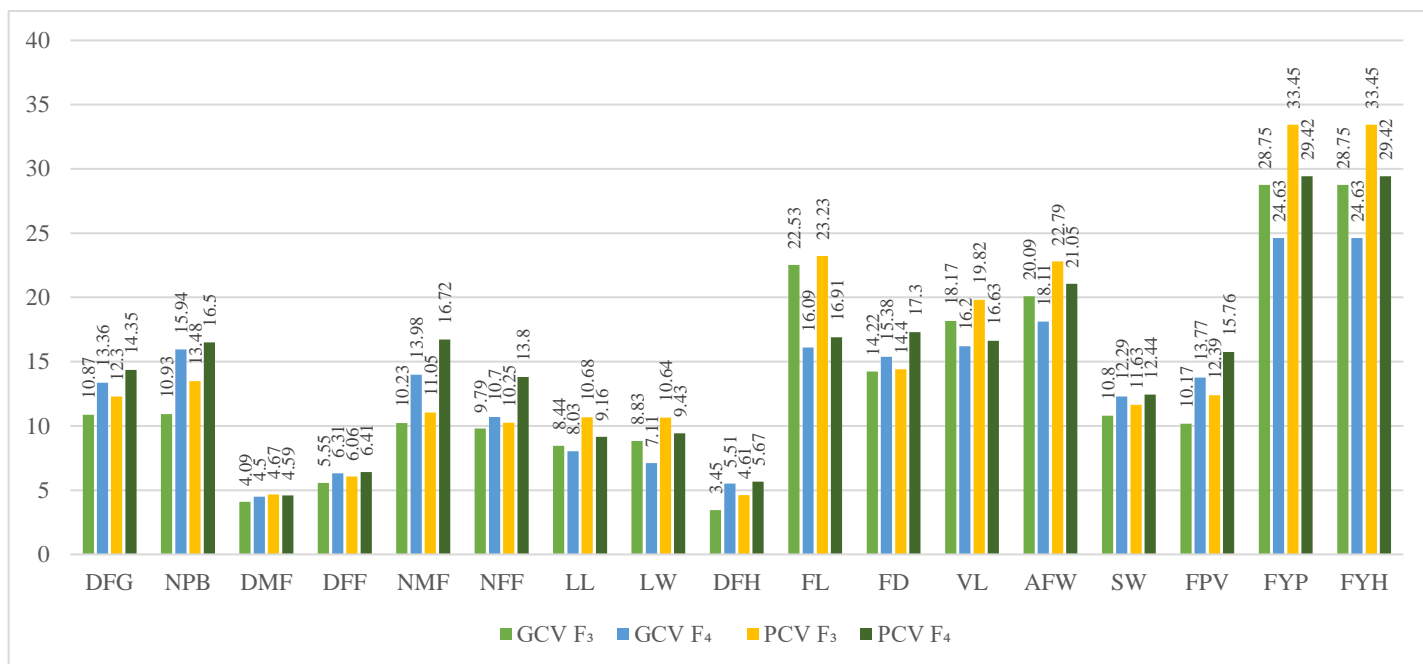


Figure 1. Estimation of GCV and PCV, (GH-28 × Pusa Santushti, F₃ and F₄ generations) : DFG: Days to 50% germination; NPB: Number of primary branches ; DMF: Days to first male flower opening; DFF: Days to first female flower opening; NMF: Node to first male flower; NFF: Node to first female flower; LL: Leaf length (cm); LW: Leaf width (cm); DFH: Days to first harvest; FL: Fruit length (cm); FD: Fruit diameter (cm): VL: Vine length at the time of final harvest (m); AFW: average fruit weight (g); SW: 100 seed weight (g); FPV: Number of fruits per vine; FYP: Fruit yield per plot (kg); FYH: fruit yield per hectare (q/ha).

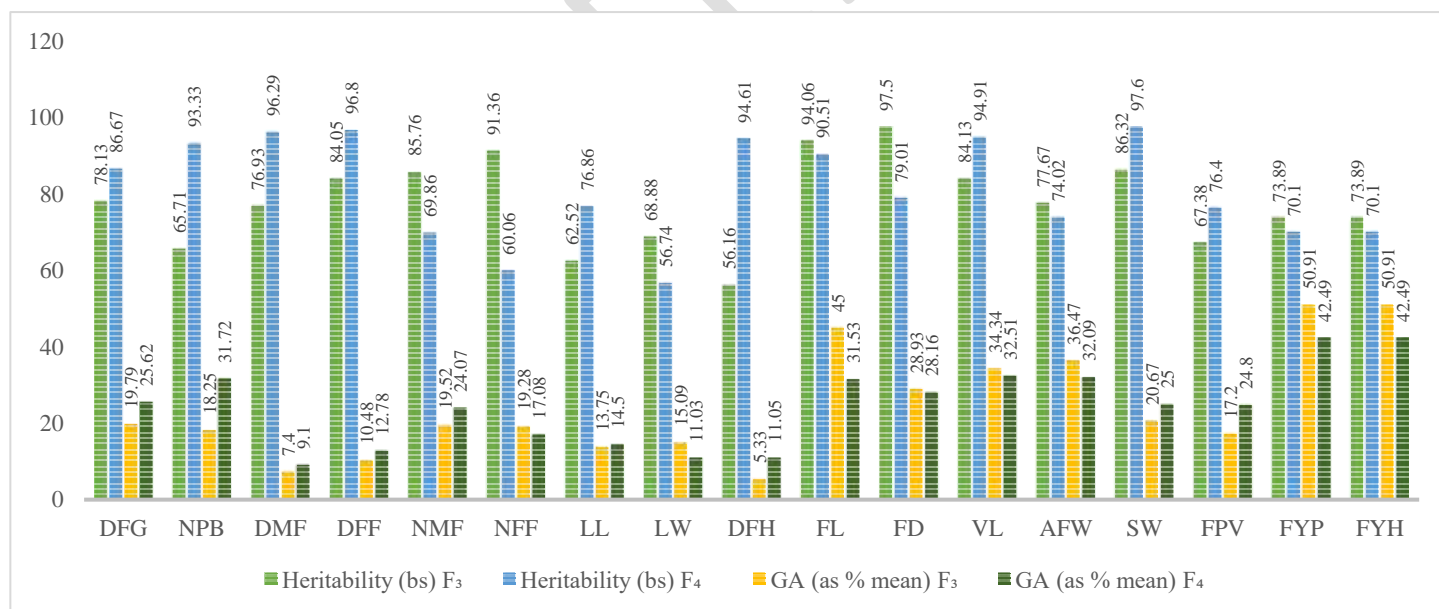


Figure 2. Estimation of heritability (b.s.) and genetic advance (GH-28 × Pusa Santushti, F₃ and F₄ generations) : DFG: Days to 50% germination; NPB: Number of primary branches ; DMF: Days to first male flower opening; DFF: Days to first female flower opening; NMF: Node to first male flower; NFF: Node to first female flower; LL: Leaf length (cm); LW: Leaf width (cm); DFH: Days to first harvest; FL: Fruit length (cm); FD: Fruit diameter (cm): VL: Vine length at the time of final harvest (m); AFW: average fruit weight (g); SW: 100 seed weight (g); FPV: Number of fruits per vine; FYP: Fruit yield per plot (kg); FYH: fruit yield per hectare (q/ha).

3.2 Correlation coefficients and path coefficient analysis

The analysis of correlation values revealed that, for most traits, the genotypic correlation coefficients were generally higher than the phenotypic correlation coefficients. This indicates that the environmental factors tend to diminish the phenotypic expression of traits even when there is a strong inherent association between them.

Fruit yield per hectare exhibited significant positive genotypic correlations with the number of primary branches (0.778 and 0.904), leaf width (0.943 and 0.836), number of fruits per vine (0.995 and 0.682), and average fruit weight (0.974 and 0.856). Conversely, there were negative genotypic correlations with days to first harvest (-0.624 and -0.784), days to first female flower opening (-0.926 and -0.661), node to first female flower (-0.524 and -0.650), and days to first male flower opening (-0.858 and -0.510) in both the F₃ and F₄ generations (Tables 2 and 3). This pattern suggests that earlier blooming, indicated by fewer nodes and fewer days until the first flower opens, leads to higher yields. Similar findings were reported by Rani *et al.* (2014) in the F₆ generation of ridge gourd for flowering traits. The observed correlations among fruit traits align with the results of Kannan *et al.* (2019) and Muttur *et al.* (2016) in the F₄ generation of pumpkin. Additionally, Vaidya (2018) noted a significant positive genotypic correlation between fruit yield and traits such as the number of primary branches, fruit length, fruit diameter, number of fruits per vine, and average fruit weight. The data suggest that fruit size, influenced by fruit length and diameter, positively correlates with average fruit weight.

To evaluate the direct and indirect effects of contributing traits on yield per hectare, path coefficient analysis was conducted using the genotypic correlation coefficients, with yield per hectare as the dependent variable and other traits as independent variables. The results of the path coefficient analysis are presented in Table 4. The findings indicate that in the F₃ generation, fruit yield per plot (1.078) had the highest direct positive effect on yield per hectare, while average fruit weight (-0.133) and node to first female flower (-0.063) had direct negative effects. Similarly, in the F₄ generation, fruit yield per plot (0.581) had the highest direct positive effect, whereas vine length (-0.6617) had a direct negative effect on fruit yield per hectare.

The results for the F₃ generation were consistent with the findings of Bhoomika *et al.* (2021). Similarly, in the F₂ generation of pumpkin, Gupta *et al.* (2018) identified a direct effect of average fruit weight and the number of fruits per vine on fruit yield. Kannan *et al.* (2019) observed a direct positive effect on fruit yield from fruit length, fruit diameter, average fruit weight, and the number of fruits per vine in the F₄ generation of ridge gourd. In the F₃ generation of pumpkin, Krishnamoorthy and Avinashgupta (2021) found a negative direct effect on fruit yield per hectare from the days to the first female bloom and the length of the fruit. Additionally, Das (2017) recorded a positive indirect effect on fruit yield via fruit length through average fruit weight in the F₃ generation of bottle gourd.

Table 2. Genotypic and phenotypic correlation co-efficient for yield and yield contributing characters (GH-28 × P. Santushti, F₃ generation)

	DFG	NPB	DMF	DFF	NMF	NFF	LL	LW	DFH	FL	FD	VL	AFW	SW	FPV	FYP	FYH
DFG	1.000	-0.135	-0.340	-0.548*	-0.429	0.520*	0.549*	0.259	0.030	-0.274	0.252	-0.445	-0.028	-0.075	0.474	0.149	0.149
NPB	-0.186	1.000	-0.671**	-0.178	-0.532*	-0.417	0.096	0.392	0.093	0.541*	0.601*	0.743**	0.476	0.447	0.409	0.431	0.431
DMF	-0.364	-0.889**	1.000	0.375	0.928**	0.081	-0.485*	-0.467	0.036	-0.463	-0.490*	-0.332	-0.595*	-0.170	-0.655**	-0.639**	-0.639**
DFF	-0.647**	-0.308	0.572*	1.000	0.550*	0.197	0.005	-0.797**	0.553*	-0.530*	-0.214	0.274	-0.547*	-0.628**	-0.738**	-0.618**	-0.618**
NMF	-0.453	-0.743**	0.990**	0.632**	1.000	0.054	-0.483*	-0.507*	0.188	-0.501*	-0.311	-0.154	-0.680**	-0.190	-0.772**	-0.732**	-0.732**
NFF	0.654**	-0.412	-0.017	0.301	0.005	1.000	0.606**	-0.246	0.581*	-0.807**	0.070	-0.296	-0.613**	-0.794**	-0.006	-0.462	-0.462
LL	0.552*	0.034	-0.579*	0.087	-0.621**	0.857**	1.000	-0.162	0.547*	-0.329	0.092	-0.042	-0.175	-0.529*	0.352	-0.073	-0.073
LW	0.421	0.896**	-0.815**	-0.876**	-0.730**	-0.495*	-0.109	1.000	-0.367	0.656**	0.546*	0.179	0.593*	0.597*	0.512*	0.612**	0.612**
DFH	-0.158	-0.074	0.010	0.903**	0.171	0.760**	0.521*	-0.637**	1.000	-0.577*	0.237	0.328	-0.685**	-0.624**	-0.327	-0.678**	-0.678**
FL	-0.296	0.788**	-0.522*	-0.539*	-0.521*	-0.902**	-0.412	0.698**	-0.747**	1.000	0.107	0.312	0.845**	0.734**	0.379	0.737**	0.737**
FD	0.230	0.726**	-0.541*	-0.240	-0.328	0.090	0.078	0.707**	0.304	0.123	1.000	0.577*	0.114	0.252	0.202	0.156	0.156
VL	-0.570*	0.883**	-0.470	0.317	-0.218	-0.293	-0.052	0.283	0.465	0.384	0.650**	1.000	0.225	0.178	-0.176	0.106	0.106
AFW	0.024	0.836**	-0.772**	-0.778**	-0.801**	-0.692**	-0.022	0.857**	-0.634**	0.998**	0.146	0.231	1.000	0.590*	0.527*	0.972**	0.972**
SW	-0.110	0.415	-0.175	-0.716**	-0.165	-0.820**	-0.810**	0.904**	-0.933**	0.833**	0.283	0.098	0.743**	1.000	0.326	0.528*	0.528*
FPV	0.849**	0.397	-0.910**	-0.940**	-0.989**	0.017	0.489*	0.894**	-0.524*	0.487*	0.282	-0.154	0.836**	0.324	1.000	0.648**	0.648**
FYP	0.289	0.778**	-0.858**	-0.926**	-0.911**	-0.524*	0.145	0.943**	-0.624**	0.915**	0.202	0.105	0.974**	0.703**	0.995**	1.000	1.000**
FYH	0.289	0.778**	-0.858**	-0.926**	-0.911**	-0.524*	0.146	0.943**	-0.624**	0.915**	0.201	0.105	0.974**	0.703**	0.995**	1.000**	1.000

DFG: Days to 50% germination; NPB: Number of primary branches ; DMF: Days to first male flower opening; DFF: Days to first female flower opening; NMF: Node to first male flower; NFF: Node to first female flower; LL: Leaf length (cm); LW: Leaf width (cm); DFH: Days to first harvest; FL: Fruit length (cm); FD: Fruit diameter (cm); VL: Vine length at the time of final harvest (m); AFW: average fruit weight (g); SW: 100 seed weight (g); FPV: Number of fruits per vine; FYP: Fruit yield per plot (kg); FYH: fruit yield per hectare (q/ha), *significant at 5 % level of significance, **significant at 1 % level of significance.

Table 3. Genotypic and phenotypic correlation co-efficient for yield and yield contributing characters (GH-28 × P. Santushti, F₄ generation)

	DFG	NPB	DMF	DFF	NMF	NFF	LL	LW	DFH	FL	FD	VL	AFW	SW	FPV	FYP	FYH
DFG	1.000	0.090	0.346	-0.152	-0.202	-0.632**	0.563*	0.032	-0.017	0.566*	0.618**	0.434	0.322	0.288	0.360	0.415	0.415
NPB	0.118	1.000	-0.754**	-0.753**	0.780**	-0.147	0.365	0.494*	-0.895**	0.465	0.681**	0.817**	0.791**	-0.728**	0.402	0.791**	0.791**
DMF	0.363	-0.804**	1.000	0.316	-0.773**	-0.312	-0.205	-0.258	0.660**	-0.027	-0.226	-0.647**	-0.348	0.923**	-0.243	-0.379	-0.379
DFF	-0.132	-0.764**	0.338	1.000	-0.332	0.472	-0.291	-0.324	0.900**	-0.703**	-0.641**	-0.419	-0.879**	0.445	0.081	-0.600*	-0.600*
NMF	-0.117	0.874**	-0.871**	-0.422	1.000	0.198	0.058	0.603*	-0.571*	0.013	0.291	0.676**	0.423	-0.540*	0.515*	0.573*	0.573*
NFF	-0.923**	-0.356	-0.397	0.715**	0.089	1.000	-0.363	-0.136	0.274	-0.657**	-0.616**	-0.211	-0.458	-0.201	-0.037	-0.360	-0.360
LL	0.752**	0.423	-0.177	-0.316	-0.07	-0.632**	1.000	-0.136	-0.285	0.745**	0.726**	0.654**	0.383	-0.311	0.120	0.332	0.332
LW	0.172	0.653**	-0.334	-0.444	0.914**	-0.293	-0.347	1.000	-0.358	-0.071	0.157	0.397	0.616**	-0.055	0.598*	0.752**	0.752**
DFH	0.021	-0.952**	0.675**	0.932**	-0.718**	0.407	-0.358	-0.527*	1.000	-0.557*	-0.601*	-0.612**	-0.823**	0.725**	-0.015	-0.610**	-0.610**
FL	0.637**	0.478	-0.005	-0.750**	-0.128	-0.946**	0.833**	-0.090	-0.604*	1.000	0.779**	0.454	0.640**	-0.233	-0.130	0.399	0.399
FD	0.793**	0.746**	-0.282	-0.702**	0.362	-0.880**	0.875**	0.405	-0.699**	0.910**	1.000	0.757**	0.660**	-0.275	0.342	0.674**	0.674**
VL	0.499*	0.856**	-0.645**	-0.430	0.739**	-0.387	0.669**	0.527*	-0.633**	0.466	0.864**	1.000	0.594*	-0.561*	0.601*	0.737**	0.737**
AFW	0.348	0.912**	-0.413	-0.982**	0.649**	-0.833**	0.405	0.568*	-0.968**	0.758**	0.927**	0.701**	1.000	-0.401	0.246	0.856**	0.856**
SW	0.347	-0.781**	0.948**	0.463	-0.695**	-0.270	-0.391	-0.055	0.743**	-0.259	-0.385	-0.592*	-0.453	1.000	-0.021	-0.304	-0.304
FPV	0.463	0.412	-0.357	0.134	0.747**	-0.074	0.161	0.756**	-0.108	-0.164	0.375	0.764**	0.201	-0.054	1.000	0.707**	0.707**
FYP	0.507*	0.904**	-0.510*	-0.661**	0.894**	-0.650**	0.397	0.836**	-0.784**	0.477	0.884**	0.943**	0.856**	-0.377	0.682**	1.000	1.000**
FYH	0.507*	0.904**	-0.510*	-0.661**	0.894**	-0.650**	0.397	0.836**	-0.784**	0.477	0.884**	0.943**	0.856**	-0.377	0.682**	1.000**	1.000

DFG: Days to 50% germination; NPB: Number of primary branches ; DMF: Days to first male flower opening; DFF: Days to first female flower opening; NMF: Node to first male flower; NFF: Node to first female flower; LL: Leaf length (cm); LW: Leaf width (cm); DFH: Days to first harvest; FL: Fruit length (cm); FD: Fruit diameter (cm); VL: Vine length at the time of final harvest (m); AFW: average fruit weight (g); SW: 100 seed weight (g); FPV: Number of fruits per vine; FYP: Fruit yield per plot (kg); FYH: fruit yield per hectare (q/ha), *significant at 5 % level of significance, **significant at 1 % level of significance

		DFG	NPB	DMF	DFF	NMF	NFF	LL	LW	DFH	FL	FD	VL	AFW	SW	FPV	FYP
DFG	F ₃	0.027	0.001	-0.014	-0.034	0.019	-0.041	0.003	0.016	0.004	-0.003	-0.004	-0.020	-0.003	0.003	0.013	0.311
	F ₄	0.377	-0.012	-0.035	0.011	-0.025	-0.041	0.012	0.038	0.005	-0.084	0.318	-0.308	0.060	-0.142	0.029	0.295
NPB	F ₃	-0.005	-0.005	-0.033	-0.016	0.031	0.026	0.0001	0.033	0.002	0.009	-0.012	0.031	-0.111	-0.012	0.006	0.839
	F ₄	0.045	-0.105	0.077	0.066	0.184	-0.016	0.007	0.143	-0.246	-0.063	0.299	-0.528	0.157	0.320	0.026	0.526
DMF	F ₃	-0.010	0.004	0.037	0.030	-0.042	0.001	-0.003	-0.030	0.0001	-0.006	0.009	-0.017	0.103	0.005	-0.014	-0.925
	F ₄	0.137	0.085	-0.096	-0.029	-0.184	-0.018	-0.003	-0.073	0.175	0.001	-0.113	0.398	-0.071	-0.389	-0.023	-0.297
DFF	F ₃	-0.018	0.002	0.021	0.053	-0.027	-0.019	0.0001	-0.033	-0.020	-0.006	0.004	0.011	0.103	0.021	-0.015	-0.999
	F ₄	-0.050	0.081	-0.032	-0.086	-0.089	0.032	-0.005	-0.097	0.241	0.099	-0.281	0.266	-0.169	-0.190	0.009	-0.384
NMF	F ₃	-0.012	0.004	0.037	0.033	-0.042	0.0001	-0.003	-0.027	-0.004	-0.006	0.005	-0.008	0.106	0.005	-0.015	-0.982
	F ₄	-0.044	-0.092	0.084	0.036	0.211	0.004	-0.001	0.200	-0.186	0.017	0.145	-0.456	0.112	0.285	0.047	0.520
NFF	F ₃	0.018	0.002	-0.001	0.016	0.0001	-0.063	0.004	-0.018	-0.017	-0.011	-0.001	-0.010	0.092	0.024	0.0001	-0.565
	F ₄	-0.348	0.038	0.038	-0.061	0.019	0.045	-0.010	-0.064	0.105	0.125	-0.352	0.239	-0.143	0.111	-0.005	-0.378
LL	F ₃	0.015	0.0001	-0.021	0.005	0.026	-0.054	0.005	-0.004	-0.012	-0.005	-0.001	-0.002	0.003	0.024	0.008	0.157
	F ₄	0.283	-0.045	0.017	0.027	-0.015	-0.028	0.016	-0.076	-0.093	-0.110	0.350	-0.413	0.070	0.160	0.010	0.231
LW	F ₃	0.012	-0.004	-0.030	-0.046	0.031	0.031	-0.001	0.037	0.014	0.008	-0.012	0.010	-0.114	-0.026	0.014	1.017
	F ₄	0.065	-0.069	0.032	0.038	0.193	-0.013	-0.005	0.219	-0.136	0.012	0.162	-0.325	0.098	0.023	0.048	0.486
DFH	F ₃	-0.004	0.0001	0.0001	0.048	-0.007	-0.047	0.003	-0.024	-0.022	-0.009	-0.005	0.016	0.084	0.027	-0.008	-0.673
	F ₄	0.008	0.100	-0.065	-0.080	-0.151	0.018	-0.006	-0.115	0.259	0.080	-0.280	0.391	-0.166	-0.305	-0.007	-0.455
FL	F ₃	-0.008	-0.004	-0.019	-0.028	0.022	0.056	-0.002	0.026	0.017	0.012	-0.002	0.013	-0.133	-0.024	0.008	0.987
	F ₄	0.240	-0.050	0.0001	0.064	-0.027	-0.042	0.013	-0.020	-0.156	-0.132	0.365	-0.288	0.130	0.106	-0.010	0.277
FD	F ₃	0.006	-0.004	-0.020	-0.013	0.014	-0.006	0.000	0.026	-0.007	0.001	-0.017	0.023	-0.019	-0.008	0.004	0.217
	F ₄	0.299	-0.079	0.027	0.060	0.076	-0.040	0.014	0.089	-0.181	-0.120	0.400	-0.533	0.159	0.158	0.024	0.513
VL	F ₃	-0.016	-0.004	-0.017	0.017	0.009	0.018	0.0001	0.011	-0.010	0.005	-0.011	0.035	-0.031	-0.003	-0.002	0.113
	F ₄	0.188	-0.090	0.062	0.037	0.156	-0.017	0.010	0.115	-0.164	-0.061	0.346	-0.617	0.121	0.243	0.048	0.548

AFW	F ₃	0.001	-0.004	-0.029	-0.041	0.034	0.043	0.0001	0.032	0.014	0.012	-0.002	0.008	-0.133	-0.022	0.013	1.050
	F ₄	0.131	-0.096	0.040	0.084	0.137	-0.037	0.006	0.124	-0.250	-0.100	0.371	-0.433	0.172	0.186	0.013	0.497
SW	F ₃	-0.003	-0.002	-0.007	-0.038	0.007	0.051	-0.004	0.034	0.021	0.010	-0.005	0.003	-0.099	-0.029	0.005	0.758
	F ₄	0.131	0.082	-0.091	-0.040	-0.147	-0.012	-0.006	-0.012	0.192	0.034	-0.154	0.365	-0.078	-0.410	-0.003	-0.219
FPV	F ₃	0.023	-0.002	-0.034	-0.050	0.042	-0.001	0.003	0.033	0.012	0.006	-0.005	-0.005	-0.111	-0.009	0.016	1.073
	F ₄	0.174	-0.044	0.034	-0.012	0.157	-0.003	0.003	0.165	-0.028	0.022	0.150	-0.471	0.034	0.022	0.063	0.396
FYP	F ₃	0.008	-0.004	-0.032	-0.049	0.038	0.033	0.001	0.035	0.014	0.011	-0.003	0.004	-0.129	-0.021	0.016	1.078
	F ₄	0.191	-0.095	0.049	0.057	0.189	-0.029	0.006	0.183	-0.203	-0.063	0.354	-0.582	0.147	0.154	0.043	0.581

Table 4. Estimates of direct (diagonal values) and indirect effects of various characters over yield per hectare cross (GH-28 × P. Santushti)

Residual are 0.00184 in F₃ and 0.03534 in F₄; DFG: Days to 50% germination; NPB: Number of primary branches ; DMF: Days to first male flower opening; DFF: Days to first female flower opening; NMF: Node to first male flower; NFF: Node to first female flower; LL: Leaf length (cm); LW: Leaf width (cm); DFH: Days to first harvest; FL: Fruit length (cm); FD: Fruit diameter (cm); VL: Vine length at the time of final harvest (m); AFW: average fruit weight (g); SW: 100 seed weight (g); FPV: Number of fruits per vine; FYP: Fruit yield per plot (kg)

4. CONCLUSION

Although greater amount of variability is present in bottle gourd, this is not utilized in crop improvement program efficiently. Therefore, there is a need of hybrid development keeping in mind the results of genetic parameters studied. The variability analysis indicated that the traits, number of primary branches, fruit length and average fruit weight showed considerable variabilities in both the generations. Correlation and path analysis studies revealed that number of primary branches, average fruit weight, days to first harvest and days to first female flower opening had significant effect on fruit yield per hectare. And therefore, these characteristics can be successfully employed in breeding program as their contribution to fruit yield per hectare is significant. Along with the abovementioned information, emphasis on the traits with stabilized homozygosity will make the selection efficient. Further, the traits in heterozygous condition can be improved in subsequent generations.

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