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Correlation and path analysis of yield contributing characters in fifteen strawberry genotypes under Mid-Hill conditions of Himachal Pradesh

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Abstract

A comprehensive evaluation of fifteen strawberry genotypes was conducted over two consecutive growing seasons (2015-16 and 2016-17) at the experimental field, Department of Fruit Science, Dr. YS Parmar University of Horticulture and Forestry, Nauni, Solan, Himachal Pradesh. The experiment was laid out in a Randomized Complete Block Design with three replications. Correlation and path analysis studies in fifteen strawberry cultivars namely Addie, Belrubi, Brighton, Chandler, Confectura, Douglas, Etna, Fern, Gorella, Elyana, Selva, Shasta, Sweet Charlie, Tioga, and Torrey were carried out. Number of fruits/plant, average berry weight and leaf number/plant had high positive direct effect on yield/plant during first year of study while during 2016-17, days to flowering had maximum positive direct effect on fruit yield.

Keywords: Strawberry genotypes, yield attributes, correlation analysis, path analysis

Introduction

The cultivated strawberry ($Fragaria \times ananassa$ Duch.) stands as one of the most economically significant horticultural crops worldwide, belonging to the family Rosaceae. This octaploid species (2n=8x=56) represents a remarkable example of interspecific hybridization. The maternal parent, Fragaria chiloensis L., contributed large fruit size characteristics from its Chilean origin, while the paternal parent, Fragaria virginiana Duch., from eastern North America, provided essential vigor and adaptability traits that established the foundation of modern strawberry cultivation.

Contemporary strawberry breeding programs face the perpetual challenge of enhancing fruit yield while maintaining quality attributes. Yield, being a complex quantitative trait, results from the intricate interaction of multiple morphological, physiological, and biochemical components, further modulated by environmental factors. This complexity renders direct selection for yield improvement challenging and often inefficient. Consequently, understanding the genetic architecture underlying yield components and their interrelationships becomes paramount for developing effective breeding strategies.

Given the complex genetic architecture of strawberry and the multifaceted nature of yield components, the present investigation was undertaken to evaluate the extent and magnitude of variability in vegetative and reproductive traits within strawberry germplasm. The primary objectives include assessing phenotypic and genotypic variations, determining correlation coefficients among various morphological and yield-related characters, and conducting path coefficient analysis to elucidate the direct and indirect effects of component traits on fruit yield. This comprehensive approach will provide valuable insights for developing efficient selection criteria and breeding strategies for strawberry improvement programs.

Methodology

The present investigation entitled "Evaluation of strawberry genotypes through morphological and molecular markers" was conducted at the Department of Fruit Science, Dr. Y.S. Parmar University of Horticulture and Forestry, Nauni, Solan, Himachal Pradesh, India during 2015-16 and 2016-17 for fifteen genotypes. The statistical analysis was performed using MS-Excel and OPSTAT software packages. The significance of differences between cultivars was tested at 5% level of probability (P≤0.05). Critical Difference (CD)

values were calculated at 0.05 % level of significance to compare the means of different treatments. The pooled analysis was conducted by combining the data from both experimental years to obtain overall performance estimates for each cultivar. Standard statistical procedures were followed to ensure the validity and reliability of the results obtained from the morphological evaluation of strawberry genotypes.

Results and Discussion Correlation Studies

The simple correlations among fruit yield per plot and other vegetative, flowering, fruit yield characters were worked out and are presented in Table 1 (2015-16) and in Table 2 for (2016-17). During the first year yield per plant had positive and significant correlation with characters like leaf area (0.23), flower size (0.07), fruit breadth (0.24), number of flowers per plant (0.47), average berry weight (0.08), number of achene per fruit (0.54) during the first year of evaluation whereas trait namely plant height (-0.21), number of leaves(-0.51), number of runners (-0.10) days to flowering (-0.10) duration of flowering (-0.11), days to maturity (-0.23), and fruit length (-0.01) days to flowering (-0.57) showed negative significant correlation with yield per plant. During the second year number of leaves (0.01), number of runners (0.01), days to flowering (0.45), duration of flowering (0.14), days to maturity (0.21), fruit length (0.23) number of flowers (0.56) and average berry weight (0.13) showed positive and significant correlation whereas plant height (-0.11), leaf area (-0.06), flower size (-0.23), fruit breadth (-0.22) and number of achene per fruit (-0.29) showed negative and significant correlation with yield per plot.

Genotypic and Phenotypic Correlation

The correlation coefficient among the different characters was worked out at genotypic and phenotypic levels. In general, the genotypic correlation coefficient was higher in magnitude than phenotypic correlation coefficients. The association was found to be positive and significant between yield per plot and, leaf area (0.73 and 0.69), fruit size (0.72 and 0.52), days to flower (0.56 and 0.49) fruit length (0.52

and 0.49), fruit breadth (0.61 and 0.58), number of fruits per plant (0.92 and (0.89), and berry weight (0.76 and 0.66) achene number (0.84 and 0.80) except leaf number (0.16 and 0.17), number of runners per plant (0.14 and 0.13) which shows positive and non-significant association both at genotypic and phenotypic level during the year 2015-16. Plant height shows positive and significant association at genotypic level (0.483) and positive and non significant association at phenotypic level (0.52). While significant and negative correlations at both genotypic and phenotypic levels was observed with days maturity (-0.75 and -0.68). Non-significant and negative association was observed for days to flowering after planting (-0.21 and -0.19).

In the year 2016-17 plant height (0.58 and 0.51), leaf number (0.62 and 0.59), leaf area (0.69 and 0.66), number of runners per plant (0.53 and 0.47), flower size (0.86 and (0.54), days to flower (0.45 and 0.39), fruit length (0.60 and 0.52), fruit breadth (0.57 and (0.52), number of fruits per plant (0.96 and 0.92) and berry weight (0.81 and 0.79) had significant and positive correlation with yield per plant both at genotypic and phenotypic levels. The characters days to flower (-0.11 and -0.07) showed non-significant and negative whereas days to maturity (-0.75 and -0.68) shows significant and negative correlation both at genotypic and phenotypic level.

The genotypic correlation coefficients of fruit yield per plant and yield-contributing characters were higher than phenotypic correlation coefficients in most cases, indicating that the effects of environment suppressed the phenotypic relationship between these characters.

Similar correlations of yield with various other horticultural traits had also been reported by Sharma and Suman (2006) ^[10] and yield per plant was significantly and positively associated with achene number per berry, fruit number, fruit length, fruit breadth and plant spread. Mir *et al.* (2009) ^[6] Garg *et al.* (2014) ^[2] also observed positive significant genotypic and phenotypic correlation for yield with number of fruits followed by fruit length which is found in agreement with the present findings. Similar reports were also suggested by Chaubey and Singh (1994) ^[1] and Ojo *et al.* (2006) ^[7]. Fruit yield showed strong positive and significant correlations with most of the characters.

Table 1: Correlation matrix showing relationship at genotypic phenotypic levels with respect to vegetative, flowering and fruit yield characters (2015-16)

		2	3	4	5	6	7	8	9	10	11	12	13	14
1	G	0.35	0.59^{*}	0.03	0.75^{*}	0.57^{*}	0.01	-0.49*	0.64^{*}	0.68^{*}	0.38	0.61*	0.16	0.48^{*}
	P	0.28	0.38^{*}	-0.01	0.34	0.41^{*}	0.02	-0.36	0.43*	0.37	0.26	0.34	0.11	0.29
2	G		0.14	0.36	-0.02	-0.45*	-0.27	0.31	-0.01	-0.12	0.25	0.36	0.02	0.16
	P		0.14	0.32	0.01	-0.41*	-0.23	0.28	0.01	-0.09	0.23	0.33	0.02	0.17
3	G			-0.06	0.71^{*}	0.51^{*}	0.02	-0.71*	0.56^{*}	0.54^{*}	0.65^{*}	0.58^{*}	0.40^{*}	0.73^{*}
	P			-0.04	0.53^{*}	0.49^{*}	0.02	-0.65*	0.53^{*}	0.51^{*}	0.63^{*}	0.50^{*}	0.39^{*}	0.69^{*}
4	G				-0.35	-0.23	-0.47*	0.18	-0.17	-0.08	0.13	0.04	0.18	0.14
	P				-0.27	-0.21	-0.46*	0.20	-0.14	-0.05	0.14	0.02	0.17	0.13
5	G					0.65^{*}	0.047	-0.75*	0.83^{*}	0.81*	0.75^{*}	0.83*	0.68*	0.72^{*}
	P					0.47^{*}	0.02	-0.53*	0.56^{*}	0.54^{*}	0.52^{*}	0.61*	0.49^{*}	0.52^{*}
6	G						0.10	-0.79*	0.46^{*}	0.64^{*}	0.37	0.28	0.44*	0.56*
	P						0.06	-0.75*	0.42^{*}	0.59^{*}	0.35	0.19	0.43*	0.49^{*}
7	G							-0.11	0.16	0.05	-0.22	0.06	-0.53*	-0.21
	P							-0.12	0.14	0.05	-0.20	0.07	-0.50*	-0.19
8	G								-0.54*	-0.73*	-0.59*	-0.62*	-0.52*	-0.75*
	P				_				-0.51*	-0.68*	-0.56*	-0.48*	-0.51*	-0.68*
9	G									0.93*	0.40^{*}	0.76*	0.29	0.52^{*}
	P									0.88^{*}	0.38^{*}	0.62^{*}	0.29	0.49^{*}
10	G										0.45^{*}	0.69*	0.43*	0.61*

	P					0.43*	0.60^{*}	0.42^{*}	0.58^{*}
11	G						0.70^{*}	0.82^{*}	0.92^{*}
	P						0.59^{*}	0.81*	0.89^{*}
12	G							0.55^{*}	0.76^{*}
	P							0.46^{*}	0.66^{*}
13	G								0.84^{*}
	P								0.80^{*}

^{*} Significance at 5% level of significance.

Table 2: Correlation matrix showing relationship at genotypic phenotypic levels with respect to vegetative, flowering, fruit yield and quality characters (2016-17)

		2	3	4	5	6	7	8	9	10	11	12	13	14
1	G	0.39*	0.50^{*}	0.25	0.54*	0.57^{*}	-0.39*	-0.61*	0.41*	0.59*	0.64*	0.42*	0.80^{*}	0.58^{*}
	P	0.30	0.39*	0.16	0.29	0.38*	-0.25	-0.43*	0.24	0.41*	0.52*	0.39*	0.61*	0.51*
2	G		0.70^{*}	0.29	0.93*	0.59^{*}	0.03	-0.53*	0.49^{*}	0.44^{*}	0.51*	0.53*	0.46*	0.62^{*}
	P		0.68^{*}	0.28	0.65*	0.57*	0.02	-0.51*	0.41*	0.38*	0.50*	0.45*	0.46*	0.59*
3	G			0.47*	0.62^{*}	0.54^{*}	0.04	-0.74*	0.64^{*}	0.50^{*}	0.63*	0.55*	0.40^{*}	0.69^{*}
	P			0.44*	0.44*	0.51^{*}	0.05	-0.68*	0.56^{*}	0.44^{*}	0.62*	0.47^{*}	0.39*	0.66*
4	G				0.21	0.05	-0.04	-0.12	0.59^{*}	0.49*	0.47^{*}	0.58*	0.33	0.53*
	P				0.16	0.04	-0.03	-0.14	0.47^{*}	0.42*	0.45^{*}	0.41*	0.33	0.47^{*}
5	G					0.76^{*}	-0.09	-0.61*	0.71^{*}	0.67*	0.75^{*}	0.67*	0.69*	0.86^{*}
	P					0.48^{*}	-0.02	-0.42*	0.43*	0.44*	0.47^{*}	0.44*	0.46^{*}	0.54^{*}
6	G						0.14	-0.79*	0.47^{*}	0.72^{*}	0.40^{*}	0.29	0.46^{*}	0.45^{*}
	P						0.11	-0.74*	0.32	0.56^{*}	0.37	0.19	0.44*	0.39^{*}
7	G							-0.17	0.29	0.02	-0.19	0.02	-0.47*	-0.11
	P							-0.17	0.26	0.05	-0.16	0.05	-0.45*	-0.07
8	G								-0.59*	-0.73*	-0.60*	-0.56*	-0.52*	-0.67*
	P								-0.47*	-0.62*	-0.56*	-0.42*	-0.50*	-0.59*
9	G									0.78*	0.42^{*}	0.78^{*}	0.26	0.60^{*}
	P									0.75^{*}	0.35	0.65*	0.23	0.52^{*}
10	G										0.44^{*}	0.66^{*}	0.50^{*}	0.57^{*}
	P										0.39^{*}	0.59^{*}	0.45^{*}	0.52^{*}
11	G											0.64^{*}	0.83^{*}	0.96^{*}
	P											0.53*	0.82^{*}	0.92^{*}
12	G												0.57*	0.81*
	P												0.47*	0.79^{*}
13	G													0.81*
	P													0.76*

^{*}Significance at 5% level of significance

Thus, selection may be possible for these characters for improving yield. In earlier studies, fruit yield was significantly and positively associated with most of the characters (Lacey 1973, Webb *et al.*, 1974, Guttridge and Anderson 1981, Olsen *et al.*, 1985 and Strik and Proctor, 1988) ^[5, 12, 4, 8, 11], whereas findings of Gawronski and Hortynski (2011) are in disagreement with the present investigation who recorded negative correlation between the number of flowers per inflorescence and fruit weight. In a few cases, phenotypic correlation coefficients were the same as or higher than the genotypic correlation coefficients, indicating that both environmental and genotypic correlations in these cases acted in the same direction and finally maximized their expression at the phenotypic level.

Path Coefficient Analysis

Although correlation studies are helpful in determining the components of yield but it does not provide a clear picture of nature and extent of contributions made by number of independent traits. Path co-efficient analysis depicts the

effects of different independent characters individually and in combination with other characters on the expression of different characters on yield. The data on path co-efficient level showing the direct and indirect effects of significant characters over fruit yield per plant are presented in Table 18 and 19. The data revealed that leaf area (0.56) flower size (0.09) duration of flowering, fruit length (0.23), fruit length (0.03), fruit breadth (0.38), number of fruits (0.56 and 0.68), average berry weight (0.14) and number of achene per fruit (0.54) had positive direct effect, while negative direct effect of characters plant height (-0.26), leaf number (-0.11), number of runners (-0.30), days to flower (-0.23), days to maturity (-0.32) on yield per plot during first year of study. Maximum positive direct effect on yield per plot was shown by leaf area and average berry weight (0.56) whereas maximum positive indirect effects were observed for fruit length (0.48), average berry weight (0.45) via fruit breadth and number of achene respectively. Highest negative indirect effects were recorded by number of runners (-0.32) via days to maturity followed by fruit breadth (-0.28) via

^{1.} Plant height (cm) 2. Leaf number per plant 3. Leaf area (cm²) 4. Flower size (cm) 5. Days to flowering 6. Duration of flowering 7. Days to maturity (Duration) 8. Fruit length (mm) 9. Fruit breadth (mm) 10. Number of flowers per plant 11. Average berry weight (g) 12. Number of achenes/fruit 13. Yield per plot (kg)

^{1.} Plant height (cm) 2. Leaf number per plant 3. Leaf area (cm²) 4. Flower size (cm) 5. Days to flowering 6. Duration of flowering 7. Days to maturity (Duration) 8. Fruit length (mm) 9. Fruit breadth (mm) 10. Number of flowers per plant 11. Average berry weight (g) 12. Number of achenes /fruit 13. Yield per plot (kg)

number of achene during first year. During the second year the maximum direct effect on yield per plant was shown by duration of flowering followed by fruits breadth (0.80) average berry weight (0.68), leaf area (0.61), number of achenes per fruit (0.24) and flower size (0.04). The negative direct effect on yield per plant was observed by number of leaves (-0.69), runner number (-0.16), days to flowering (-0.33), days to maturity (-0.27), fruit length (-0.41) flower number (-0.50) achene number (-0.48) fruit breadth followed by (-0.18) and leaf area (-0.18) on yield per plot. The highest positive indirect effect on yield per plant were exhibited by plant height (0.50) via number of runners per plant followed by leaf area (0.50) via number of fruits per plant, plant spread (0.49) via number of runners per plant and fruit length (0.47), plant height (0.39), fruit weight (0.36) and leaf area (0.35) via number of fruits per plant.,

the characters number of leaves (-0.69), runner number (-0.16), days to flower (-0.33), days to maturity (-0.27), fruit length (-0.41) and flower number (-0.50) shows negative direct effect on yield per plot during second year of study. The indirect positive effects were maximum in case of flower number (0.75) followed by fruit length (0.67) via fruit length.

Rao and Lal (2010) [9] also recorded similar observations as that of present investigation that maximum positive significant correlation of yield recorded with berry width followed by number of achenes per fruit, berry length and number of flowers per plant. Path coefficient analysis revealed that fruit width had highest direct positive contribution towards fruit yield per plant. These important traits may be considered in selection programme for the further improvement of yield in strawberry.

Table 3: Estimates of direct and indirect effects on fruit yield per plant at genotypic level (2015-16)

	1	2	3	4	5	6	7	8	9	10	11	12	13	GCC
1	-0.69	0.21	-0.09	0.00	-0.25	0.57	-0.01	0.20	0.52	-0.34	0.26	0.14	-0.04	0.48*
2	-0.24	0.61	-0.02	0.01	0.01	-0.45	0.07	-0.12	-0.01	0.06	0.17	0.08	-0.01	0.16
3	-0.41	0.09	-0.16	-0.01	-0.23	0.52	-0.01	0.29	0.45	-0.27	0.44	0.14	-0.11	0.73*
4	-0.02	0.21	0.01	0.04	0.11	-0.23	0.13	-0.08	-0.14	0.04	0.09	0.01	-0.05	0.14
5	-0.52	-0.01	-0.11	-0.01	-0.33	0.66	-0.01	0.31	0.67	-0.41	0.51	0.20	-0.20	0.72*
6	-0.39	-0.27	-0.08	-0.01	-0.21	1.01	-0.02	0.33	0.37	-0.32	0.25	0.06	-0.13	0.56*
7	-0.01	-0.16	-0.01	-0.02	-0.01	0.10	-0.27	0.04	0.13	-0.02	-0.15	0.01	0.15	-0.21
8	0.34	0.18	0.11	0.01	0.25	-0.81	0.03	-0.41	-0.44	0.37	-0.40	-0.15	0.15	-0.75*
9	-0.44	-0.01	-0.09	-0.01	-0.27	0.47	-0.04	0.22	0.80	-0.47	0.27	0.18	-0.08	0.52*
10	-0.47	-0.07	-0.08	-0.01	-0.27	0.65	-0.01	0.30	0.75	-0.50	0.31	0.16	-0.12	0.61*
11	-0.26	0.15	-0.10	0.01	-0.25	0.37	0.06	0.24	0.32	-0.22	0.68	0.17	-0.24	0.92*
12	-0.43	0.22	-0.09	0.01	-0.27	0.28	-0.01	0.25	0.61	-0.35	0.48	0.24	-0.16	0.76*
13	-0.11	0.01	-0.06	0.01	-0.22	0.45	0.14	0.21	0.23	-0.22	0.56	0.13	-0.29	0.84*

^{*}significance at 5% level of significance.

Table 4: Estimates of direct and indirect effects on fruit yield per plant at genotypic level (2016-17)

	1	2	3	4	5	6	7	8	9	10	11	12	13	GCC
1	-0.69	0.21	-0.09	0.00	-0.25	0.57	-0.01	0.20	0.52	-0.34	0.26	0.14	-0.04	0.48*
2	-0.24	0.61	-0.02	0.01	0.01	-0.45	0.07	-0.12	-0.01	0.06	0.17	0.08	-0.01	0.16
3	-0.41	0.09	-0.16	-0.01	-0.23	0.52	-0.01	0.29	0.45	-0.27	0.44	0.14	-0.11	0.73*
4	-0.02	0.21	0.01	0.04	0.11	-0.23	0.13	-0.08	-0.14	0.04	0.09	0.01	-0.05	0.14
5	-0.52	-0.01	-0.11	-0.01	-0.33	0.66	-0.01	0.31	0.67	-0.41	0.51	0.20	-0.20	0.72*
6	-0.39	-0.27	-0.08	-0.01	-0.21	1.01	-0.02	0.33	0.37	-0.32	0.25	0.06	-0.13	0.56*
7	-0.01	-0.16	-0.01	-0.02	-0.01	0.10	-0.27	0.04	0.13	-0.02	-0.15	0.01	0.15	-0.21
8	0.34	0.18	0.11	0.01	0.25	-0.81	0.03	-0.41	-0.44	0.37	-0.40	-0.15	0.15	-0.75*
9	-0.44	-0.01	-0.09	-0.01	-0.27	0.47	-0.04	0.22	0.80	-0.47	0.27	0.18	-0.08	0.52*
10	-0.47	-0.07	-0.08	-0.01	-0.27	0.65	-0.01	0.30	0.75	-0.50	0.31	0.16	-0.12	0.61*
11	-0.26	0.15	-0.10	0.01	-0.25	0.37	0.06	0.24	0.32	-0.22	0.68	0.17	-0.24	0.92*
12	-0.43	0.22	-0.09	0.01	-0.27	0.28	-0.01	0.25	0.61	-0.35	0.48	0.24	-0.16	0.76*
13	-0.11	0.01	-0.06	0.01	-0.22	0.45	0.14	0.21	0.23	-0.22	0.56	0.13	-0.29	0.84*

^{*}Significance at 5% level of significance

Table 5: Estimates of direct and indirect effects on fruit yield per plant at genotypic level (2016-17)

	1	2	3	4	5	6	7	8	9	10	11	12	13	GCC
1	-0.26	-0.04	0.28	-0.08	0.05	-0.13	-0.09	-0.27	-0.02	0.28	0.36	0.06	0.44	0.58*
2	-0.10	-0.11	0.39	-0.09	0.09	-0.13	0.00	-0.23	-0.02	0.21	0.29	0.07	0.25	0.62*
3	-0.13	-0.07	0.56	-0.15	0.06	-0.12	0.01	-0.32	-0.03	0.24	0.36	0.08	0.22	0.69*
4	-0.06	-0.03	0.26	-0.32	0.02	-0.01	-0.01	-0.05	-0.03	0.24	0.26	0.08	0.18	0.53*
5	-0.14	-0.10	0.356	-0.07	0.09	-0.17	-0.02	-0.27	-0.04	0.32	0.43	0.10	0.37	0.86*
6	-0.15	-0.06	0.30	-0.01	0.07	-0.23	0.03	-0.34	-0.02	0.34	0.23	0.04	0.25	0.45*
7	0.10	-0.01	0.02	0.01	-0.01	-0.03	0.23	-0.07	-0.01	0.01	-0.10	0.01	-0.25	-0.11

^{1.} Plant height (cm) 2. Leaf number per plant 3. Leaf area (cm2) 4. Flower size (cm) 5. Days to flowering 6. Duration of flowering 7. Days to maturity (Duration) 8. Fruit length (mm) 9. Fruit breadth (mm) 10. No. Of flowers per plant 11. Average berry weight (g) 12. No. of achenes per fruit 13. Yield per plot (g)

^{1.} Plant height (cm) 2. Leaf number per plant 3. Leaf area (cm2) 4. Flower size (cm) 5. Days to flowering 6. Duration of flowering 7. Days to maturity (Duration) 8. Fruit length (mm) 9. Fruit breadth (mm) 10. No. Of flowers per plant 11. Average berry weight (g) 12. No. of achieves per fruit 13. Yield per plot (g)

8	-0.15	-0.04	0.28	-0.15	0.06	-0.16	0.01	-0.32	-0.04	0.48	0.25	0.09	0.27	-0.67
9	0.16	0.05	-0.42	0.04	-0.06	0.18	-0.04	0.44	0.03	-0.35	-0.34	-0.08	-0.28	-0.67*
10	-0.10	-0.05	0.36	-0.19	0.07	-0.11	0.07	-0.26	-0.05	0.38	0.23	0.11	0.14	0.60*
11	-0.16	-0.05	0.35	-0.15	0.07	-0.09	-0.04	-0.26	-0.02	0.21	0.56	0.09	0.45	0.96*
12	-0.11	-0.05	0.31	-0.18	0.06	-0.06	0.01	-0.25	-0.04	0.31	0.36	0.14	0.31	0.81*
13	-0.21	-0.05	0.23	-0.10	0.06	-0.10	-0.11	-0.23	-0.01	0.24	0.47	0.08	0.54	0.76*

^{*}Significance at 5% level of significance

1. Plant height (cm) 2. Leaf number per plant 3. Leaf area (cm2) 4. Flower size (cm) 5. Days to flowering 6. Duration of flowering 7. Days to maturity (Duration) 8. Fruit length (mm) 9. Fruit breadth (mm) 10. No. of flowers per plant 11. Average berry weight (g) 12. No. of achenes per fruit 13. Yield per plot (g)

Summary and Conclusion

A two-year correlation and path coefficient analysis revealed that genotypic correlations were consistently higher than phenotypic correlations, indicating strong genetic control over trait relationships. Most morphological and phenological traits showed positive significant correlations with yield per plot across both years, except duration of flowering and days to maturity in 2016-17. Path analysis demonstrated that leaf area, flower size, fruit dimensions, berry weight, and achene number had high positive direct effects on yield in 2015-16, while days to flowering showed maximum direct effect in 2016-17. These findings establish that genetic factors predominantly govern yield-related trait associations, and the year-to-year variation in trait importance suggests environmental influence on yield determinants. The results provide valuable selection criteria for breeding programs, validating these traits as reliable indices for developing high-vielding cultivars through genetic improvement strategies.

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