

# GENETIC VARIABILITY AND CORRELATION STUDIES IN SOME GENOTYPES OF EGGPLANT 

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#### Abstract

Assessment of genetic variability is crucial in developing new genotypes with desired yield and quality traits. This evaluation allows breeders to identify superior germplasm, breeding genotypes with specific combinations, and enhance crop performance across diverse environmental conditions. Hence, this study aimed to study the performance of ten eggplant genotypes (Solanum melongena L.); i.e., Little Fingers (LF), Ping Tung (PT), Antigua (An), Aswad (As), Japanese White Egg (JWE), Apple Green (AG), Rotonda Bianca Stumata di Rosa (BSR), Korean Red (KR), Black Oblong (BO) and Black Very Long (BL). In addition, to explore genetic advance and correlation among studied traits under three intra-row plant spacings, ( 30,45 and 60 cm ). The study was performed at Zagazig District, Egypt during the two successive summer seasons of 2017 and 2018. The results indicated significant variation among the tested genotypes. The broad sense heritability $\left(h_{b}{ }^{2}\right)$ exhibited the highest percentage at narrow planting spaces and decreased with increased spaces in all studied traits. Broad sense heritabilities were moderate magnitude for number of branches ( $39 \%$ and $30.9 \%$ ) at 45 and 60 cm spaces, respectively. Genetic advance (GA \%) was high for all studied traits except number of branches. High heritability estimates and a high predicted genetic advance indicated that these qualities and genotypes could be effective in developing breeding programs to increase fruit quality and yield. Strong genetic correlations between morphological and yield-related variables were found via correlation studies. The path analysis results showed that the fruit weight had the largest direct effect on yield/ plant ( 2.984 and 3.11) followed by number of fruits ( 0.579 and 1.525 ), and branches number/ plant ( 0.161 and 0.304 ) at 30 and 45 cm planting spaces, respectively. While the branches number/ plant exhibited the largest direct effect on yield/ plant (5.341) followed by fruit length (3.328), and fruit diameter ( 0.625 ) at 60 cm planting space, indicating the effectiveness of direct selection for these traits for developing high yielding genotypes of eggplant under different planting spaces.


Key words: Eggplant, genotypes, heritability, genetic variability, correlation, yield traits.

## INTRODUCTION

The production potential of cultivated varieties of eggplant is lower and the choice of eggplant size, shape and skin color varies in in different locations according to the consumer. For the development of eggplant varieties with high productivity and environmental resistances, a systematic breeding plane is required. Evaluation of the most important traits helps to collect information for the development of the crop, and also provides breeders with a picture
of genotypes about genetic variance and diversity. Therefore, in order to determine the optimum genotypes for economic features, calculate the breeding value of the available germplasm, and comprehend genetic background, germplasm assessment at various planting locations is required. For any crop development to be ultimately successful, variability is desperately needed. Increased degree of population variability, greater chance for effective selection (Vavilov, 1951). The level of genetic variety that exists determines whether selection is

[^0]successful. Therefore, by estimating certain genetic factors as phenotypic (PCV), it is important to partition total variability into main components (non-heritable and heritable) and genotypic (GCV) coefficient of variability, heritability and expected genetic advance that gave complete indication of genetic variations of the studied traits. Heritability is important in helping the plant breeder plan and executes effective breeding strategy. Knowledge of correlation among characters is important in deciding on what selection strategy to use. Hence, there is the need to determine genetic parameters in order to design selection programmers aimed at improving the crop through selection among accessions or through hybridization. Correlation analysis measure the relationship between any pairs of traits and determines the component characters that selection can be based for improvement the economic traits. Plant height was positively correlated with average fruit weight, total yield/ plant, Fruit diameter and Fruit length (Muniappan et al., 2010; Shekar et al., 2014; Ullah et al., 2014; Prabakaran et al., 2015). Also, significant positive correlations were found between number of branches and total fruit weight (Tripathy et al., 2018). The objectives of this study were undertaken to assessment ten genotypes of eggplant for vegetative growth, yield and fruit quality, as well as estimate superior genotypes for future use.

## MATERIALS AND METHODS

The current study was completed in a private Farm at Bani Amer Village, Zagazig district, Sharqia Governorate, Egypt ( $30^{\circ} 35^{\prime} 51.3^{\prime \prime} \mathrm{N} 31^{\circ} 34^{\prime}$ $24.9^{\prime \prime} \mathrm{E}$ ) during the summers of 2017 and 2018 to assess the performance of various eggplant genotypes (Table 1) at various plant spacings of 30,45 , and 60 cm . In the summer seasons of 2017 and 2018, seeds of the genotypes were sown in speedling trays (209 sells) under greenhouses as a nursery at $1^{\text {st }} \mathrm{Feb}$. in the both seasons. This experiment was set up as a split plot with three replicates using a randomized complete block design. Each genotype was grown in a plot of 3 rows, 3 meter long and 80 cm apart. The entries in each experimental unit consisted of 30,20 and 15 plants, planted at a spacing of $30 \times 80 \mathrm{~cm} ; 45 \times 80 \mathrm{~cm}$ and $60 \times 80$
cm , respectively. Plants were thinned to one plants/hill; the plot area was $\left(7.2 \mathrm{~m}^{2}\right)$. The prescribed cultural measures for eggplant were implemented, including fertilization, irrigation, and insect control.

## Recorded Data

Ten plants at random from each plot were measured for the following traits:

## Plant growth traits

Four plants from every experimental unit were randomly selected to measure the following traits:

- Plant height ( $\mathrm{PH}, \mathrm{cm}$ ): it was measured at the season's end.
- Branches number / plant (BN): it was counted at the season's end.
- Leaves number / plant (LN): it was counted at the flowering stage.
- Leaf area (LA): leaf length $\times$ average leaf width.


## Yield traits

- Plant/ yield (yield, kg/ plant): it was determined by summing weight of all picked fruits during the productivity stage and presented as yield per plant (kg).
- Fruit number per plant (FN)
- Total yield/fed. (yield, ton/ fed.)


## Fruit quality traits:

- Fruit length (FL, cm)
- Fruit diameter (Dim, cm)
$\bullet$ Fruit weight (FW, g): (Fruit yield of each plant/Total number of fruits)
- Dry matter (DM)


## Statistical Analysis

Broad sense heritability $\left(\mathrm{h}_{\mathrm{b}}{ }^{2}\right)$ was estimated according to Allard (1960) and Falconer (1989). Heritability based on (Stansfield, 1983) $0 \leq \mathrm{x} \leq 0.2=$ low, $0.2 \leq \mathrm{x} \leq 0.5=$ medium and $\mathrm{x}>$ $0.50=$ high. Phenotypic (PCV\%) and genotypic (GCV \%) Coefficient of variability were calculated according to Singh and Chaudhary (1985). Genetic advance (GA) was calculated with

Table1. Variability among the cultivars under study and their sources

| Code | Cultivars | Abbriv. | Fruit shape | Fruit colour | Source |
| :---: | :---: | :---: | :---: | :---: | :---: |
| $\mathbf{1}$ | Little Fingers | L.F. | Long | Purple Black | BGHSC $^{1}$ |
| $\mathbf{2}$ | Ping Tung | P.T. | Long | Purple Rose | BGHSC $^{1}$ |
| $\mathbf{3}$ | Antigua | An. | Long | White Purple with | BGHSC $^{1}$ |
|  |  |  | Strips |  |  |
| $\mathbf{4}$ | Black Very Long | B.V.L. | Long | Dark Black | Metwally $^{2}$ |
| $\mathbf{5}$ | Aswad | As. | Like a squat teardrop | Black Purple | BGHSC $^{1}$ |
| $\mathbf{6}$ | Japanese White Egg | J.W.E. | Oval | Snow White | BGHSC $^{1}$ |
| $\mathbf{7}$ | Apple Green | A.G. | Oval | Light Green | BGHSC $^{1}$ |
| $\mathbf{8}$ | Rotonda Bianca Stumata | R.B.S. | Round | White - Rose - | BGHSC $^{1}$ |
|  | di Rosa | Korean Red | K.R. | Round | Red - Orange |
| $\mathbf{9}$ | Black Oblong | B.O. | Oval | BHSC ${ }^{1}$ |  |
| $\mathbf{1 0}$ | Black | Metwally $^{2}$ |  |  |  |

1: BGHSC: Baker Greek Heiriool Seed Company, 2278 Baker Greek Road Mansfield, MO-65704 World Wide Web: RareSeeds.com
2: Prof. Dr. E. I. Metwally, Fac, Agric,Kafr El-Sheikh Univ. Kafr El-Sheikh, Egypt
the method suggested by Johnson et al. (1955) as: $\mathrm{GA}=\mathrm{K} . \mathrm{h}^{2}{ }_{(\mathrm{b})} \cdot \sigma_{\mathrm{p}}$, where: $\mathrm{K}=1.76$, constant (On the basis of intensity of the selection $10 \%$ ), $h^{2}(b)=$ heritability in broad sense and $\sigma p=$ phenotypic standard deviation. Genetic advance as percent of mean (expected genetic advance) $\mathrm{GAM} \%=(\mathrm{GA} / \bar{X}) \times 100$. GAM\% based on Hadiati et al. (2003), $0-7 \%=$ low, $7-14 \%=$ medium and $>14=$ high.

The interrelationships among the 11 studied traits were studied overall genotypes using two statistical procedures which differ in their mathematical background, goals and final outputs. These used models are summarized as follows:

1. Using the formula provided by Al-Jibouri et al. (1958) estimates of the correlation between genotype and phenotype.
2. Following the instructions provided by Dewey and Lu (1959), the path coefficient analysis was conducted among seven traits: yield/plant and each of plant height, branch number/plant, fruit number per plant, fruit length, fruit diameter, and fruit weight. The statistical computer program PATHCA was used to calculate the correlation and path coefficients (Atia, 2007).

## RESULTS AND DISCUSSION

## Genetic Variability

Data information in Table 2 showed that for every trait, the size of the genotypic (GCV) and phenotypic (PCV) coefficients of variance were differed. The highest PCV and GCV were observed for all studied traits in the three spaces, indicating the high potential for effective selection (Burton, 1952; Hamed, 2012; ElDakkak et al., 2014; Dash et al., 2020; Kumar et al., 2020; Khan et al., 2023). The majority of features under study were found to have high GCV and PCV. For the majority of traits, there were slight variations between GCV\% and PCV\% at the three spaces, suggesting the significance of genetic factors in regulating the inheritance of these traits. According to our findings, the majority of the examined qualities exhibited high GCV/PCV percent in all three spaces. Such values ranged from (75.3, 62.4 and $55.5 \%$ ) for branches to ( $98.6,98.4$ and $97.0 \%$ ) for fruit weight at narrow ( 30 cm ), mediate ( 45 cm ) and wide ( 60 cm ) planting spaces, respectively. These findings showed that genetic variations accounted for roughly $55 \%$ of phenotypic variability. As a result, there may

Table 2. Different genetic parameters for eleven vegetative and quantitative traits at three planting spaces

|  | PH | BN | FL | Dim. | FN | FW | Y kg/p | DM | Y/fed | LN | LA |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 30 cm |  |  |  |  |  |  |  |  |  |  |  |
| Mean | 87.7 | 5.6 | 9.8 | 6.1 | 13.8 | 138.2 | 1.5 | 9.5 | 26.7 | 80.06 | 47.01 |
| SE | 2.4 | 0.3 | 0.2 | 0.3 | 0.3 | 1.9 | 0.0 | 0.3 | 0.4 | 1.68 | 1.66 |
| PCV | 9.79 | 10.96 | 28.22 | 41.4 | 35.94 | 23.11 | 113.95 | 12.48 | 35.45 | 23.24 | 20.20 |
| GCV | 9.62 | 8.26 | 27.68 | 40.4 | 35.10 | 22.79 | 95.34 | 12.07 | 32.27 | 22.34 | 19.84 |
| GCV/PCV | 98.24 | 75.3 | 98.1 | 97.6 | 97.7 | 98.6 | 83.7 | 96.7 | 91.0 | 96.13 | 98.23 |
| GAM\% | 16.63 | 10.9 | 47.8 | 69.4 | 60.4 | 39.6 | 140.1 | 20.6 | 51.4 | 37.80 | 34.29 |
| $\mathbf{h}^{2}{ }_{(b)}$ | 96.52 | 56.8 | 96.2 | 95.3 | 95.4 | 97.2 | 70.0 | 93.6 | 82.8 | 92.42 | 96.49 |
| 45 cm |  |  |  |  |  |  |  |  |  |  |  |
| Mean | 85.0 | 5.5 | 9.8 | 6.1 | 17.2 | 137.3 | 1.8 | 9.5 | 24.2 | 94.43 | 48.82 |
| SE | 2.3 | 0.5 | 0.3 | 0.3 | 0.5 | 2.5 | 0.0 | 0.4 | 0.3 | 1.60 | 1.45 |
| PCV | 11.34 | 13.86 | 26.46 | 41.47 | 38.48 | 22.50 | 103.12 | 11.41 | 32.52 | 26.27 | 17.87 |
| GCV | 10.18 | 8.66 | 25.84 | 40.42 | 36.26 | 22.13 | 80.98 | 10.81 | 23.73 | 23.74 | 17.50 |
| GCV/PCV | 89.8 | 62.4 | 97.7 | 97.5 | 94.2 | 98.4 | 78.5 | 94.8 | 73.0 | 90.37 | 97.92 |
| GAM\% | 16.1 | 9.5 | 44.4 | 69.4 | 60.1 | 38.3 | 112.0 | 18.1 | 30.5 | 37.76 | 30.16 |
| $\mathbf{h}^{2}{ }_{\text {b }}{ }^{\text {a }}$ | 80.6 | 39.0 | 95.4 | 95.0 | 88.8 | 96.8 | 61.7 | 89.8 | 53.2 | 81.67 | 95.87 |
| 60 cm |  |  |  |  |  |  |  |  |  |  |  |
| Mean | 83.8 | 6.4 | 9.8 | 6.1 | 21.0 | 135.2 | 2.3 | 9.4 | 19.7 | 114.81 | 43.47 |
| SE | 1.1 | 0.4 | 0.3 | 0.3 | 0.5 | 2.7 | 0.0 | 0.2 | 0.3 | 1.97 | 1.69 |
| PCV | 11.54 | 10.66 | 28.64 | 39.88 | 37.62 | 21.71 | 101.35 | 10.98 | 32.03 | 22.53 | 22.33 |
| GCV | 9.60 | 5.92 | 26.12 | 38.03 | 35.32 | 21.06 | 68.85 | 10.40 | 22.09 | 20.30 | 19.24 |
| GCV/PCV | 83.2 | 55.5 | 91.2 | 95.4 | 93.9 | 97.0 | 67.9 | 94.7 | 69.0 | 90.11 | 86.15 |
| GAM\% | 14.0 | 5.8 | 42.0 | 63.9 | 58.4 | 36.0 | 82.2 | 17.3 | 26.8 | 32.19 | 29.17 |
| $\mathbf{h}^{\mathbf{2}}{ }_{\text {b }}$ | 69.2 | 30.9 | 83.2 | 90.9 | 88.1 | 94.1 | 46.2 | 89.7 | 47.6 | 81.19 | 74.23 |

PH: plant height, BN: branch number/plant, FL: fruit length (cm), Dim: fruit diameter (cm), FN: fruit number/plant, FW: fruit weight (g), Y/p: yield (kg/ plant), DM: fruit dry matter, Y/fed.: yield (ton/fed.), LN: leaves number/plant, and LA: leaf area.
be room for improvement in certain traits, which may be more genotypically dominant. In general, the GCV/PCV exhibited the highest percentage at narrow planting spaces and decreased with increasing spaces in all studied traits (Table 2).

Plant breeders use the heritable proportion of variation as a foundation for their selection process based on phenotypic performances. Data demonstrated that broad sense heritability ( $h^{2}$ ) exhibited the highest percentage at narrow planting spaces and decreased with increasing spaces in all studied traits (Table 2). Broad sense heritability based on Stansfield (1983), were
moderate magnitude for number of branches ( $39 \%$ and $30.9 \%$ ) at 45 and 60 cm spaces, respectively as well as yield/plant (46.2\%) and yield/fed. (47.6\%) at 45 and 60 cm space and high for each of all other traits in all traits at the three planting spaces. High heritability suggested that these features would advance quickly through selection. These findings suggested that environmental influences had little bearing on the inheritance of these qualities and that mean-based selection would be effective in enhancing these features. These parameters have a high heritability, which suggests that choosing them would be more beneficial than choosing the other factors. At 30,

45 , and 60 centimeter spacing, the genotypes showed a significant and nearly equivalent genetic gain in yield/plant, ranging from 140.1, 112.0 , and $82.2 \%$, respectively. Genetic advance based on Hadiati et al. (2003) was low for number of branches (5.8\%) at wider space (60 cm ), and moderate magnitude at both 30 cm (10.9\%) and 45 cm (9.5\%) and high for all other studied traits (Table 2).

Johnson et al. (1955) found that in order to predict the impact of selection, heritability values in conjunction with estimates of genetic advancement were more helpful than heredity alone.

In general, high heritability was achieved with all traits except number of branches ( 30 and 45 cm ) and both yield/plant and yield/ fed. $(60 \mathrm{~cm})$ along with high genetic advance as percent of the mean (GAM\%). It follows that choosing these characters would be more successful due to their high genetic advance percentage and heritability. Based on the current study's findings, it can be said that there was a significant range of variability among various genotypes for practically all of the features that were investigated. There is some similarity between these findings and those of Yadav et al. (2016), Pujer et al. (2017), Samlindsujin et al. (2017), Mahmoud et al. (2018), Tirkey et al. (2018) and Hassan et al. (2021).

## Correlation

Phenotypic, genotypic and environmental correlation coefficients among traits over the two years under 30,45 and 60 cm intra row spacing's were estimated (Tables 3 and 4). The findings showed that the signals and magnitudes of the values were either equivalent or quite near to one another for every pair of traits. The genotypic coefficient values were higher than the phenotypic one for almost all the studied traits (in each of the three spaces) revealing that environment plays minor role in the modification of the expression of the genes, thereby establishing strong inherent relationship among the traits studied under all spaces, and these results agree with those obtained by Khan et al. (2023), Kumar et al. (2020), Onyia et al. (2020) and Tiwari and Upadhyay (2011).

The positive and highly significant correlation of yield/fed., with diameter, fruit weight and
yield/plant was obtained in the three spaces implying that these traits have stable relations and these traits are yield indicators, hence could be selected simultaneously for improvement in breeding programs (Tasisa et al., 2012) and corroborated with results obtained by Ghosh et al. (2010). Therefore, the selection for fruit weight or yield/plant, simultaneously or individually should improve yielding ability of the genotypes (Mpayo, 2010). These results were in conformity with findings of Alam and Paul (2019), Jeberson et al. (2016), Khan et al. (2023), Tadesse et al. (2014) and Yadav et al. (2010). However, the picture was different under various spaces, the correlation of yield/ plant was high positive with both fruit diameter (0.946) and average fruit weight (0.929), low and positive with leaf area (+0.320) under narrow spacing ( 30 cm ) and decreased to 0.687 , 0.701 and -0.359 , respectively under the wider spacing ( 60 cm ) where, correlation decreased with increasing spaces as mentioned above. On the opposite, the correlation of yield/plant was negative with dry matter ( -0.362 ) and number of branches $(-0.294)$ as well as weak or neglected correlation with leaves number ( 0.07 ) under narrow space ( 30 cm ) increased under wider one to positive $[0.012$ (dry matter), 0.336 (leaves number) and 0.322 (branches number)] correlated values. Furthermore, both genotypic and phenotypic correlations were larger in the majority of 45 and 60 cm spacing cases than in the environmental ones (Table 4), suggesting an innate link between multiple traits independent of environmental factors effect under wide spaces.

In fact, out of a total of 55 possible combinations between the nine traits studied, only 27,18 and 27 environmental correlation coefficients in 30, 45 and 60 cm spacing, respectively were greater than their corresponding genotypic correlation coefficient.

While there was a negative association with the other growth and yield parameters examined, the fruit diameter, fruit weight, yield/plant, and yield/fed. all shown extremely strong and significant positive correlation values with each other (Table 3). Fruit diameter and average fruit weight were substantially positively correlated. These results confirmed by Danquah and Ofori (2012) and Muniappan et al. (2010) both for fruit diameter. Branches number was significantly

Table 3. Phenotypic (above diagonal) and genotypic (below diagonal) correlations among yield and its attributes of eggplant

|  |  | PH | BN | FL | Dim. | FN | FW | Y/p | DM | Y/fed | LN | LA |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| PH | 30 cm |  | 0.386 | 0.127 | -0.186 | 0.061 | -0.087 | -0.32 | -0.424 | -0.324 | -0.041 | 0.35 |
|  | 5 cm | 1 | -0.08 | 0.432 | -0.034 | -0.198 | 0.071 | -0.273 | -0.411 | -0.274 | 0.166 | -0.039 |
|  | 60 cm |  | -0.141 | 0.281 | 0.048 | -0.564 | 0.222 | -0.375 | -0.425 | -0.376 | -0.151 | 0.017 |
| BN | 30 cm | 0.375 |  | -0.004 | -0.243 | 0.441 | -0.298 | -0.205 | 0.148 | -0.207 | 0.373 | 0.53 |
|  | 45 cm | -0.004 | 1 | -0.375 | 0.014 | 0.147 | -0.117 | -0.052 | 0.189 | -0.06 | -0.151 | 0.216 |
|  | 60 cm | -0.351 |  | -0.594* | 0.118 | 0.323 | -0.113 | 0.208 | 0.181 | 0.201 | 0.465 | 0.116 |
| FL | 30 cm | 0.147 | 0.014 |  | -0.273 | -0.319 | 0.071 | -0.128 | -0.144 | -0.125 | -0.496 | -0.27 |
|  | 45 cm | 0.446 | -0.589* | 1 | -0.27 | -0.441 | 0.116 | -0.234 | -0.126 | -0.235 | -0.218 | -0.181 |
|  | 60 cm | 0.292 | -0.913** |  | -0.389 | -0.262 | -0.055 | -0.31 | -0.249 | -0.31 | -0.217 | -0.328 |
| Dim. | 30 cm | -0.202 | -0.35 | -0.283 |  | -0.655* | 0.916** | 0.940** | -0.328 | 0.938** | 0.04 | 0.368 |
|  | 45 cm | -0.042 | 0.064 | -0.267 | 1 | -0.650* | 0.887** | 0.701* | -0.481 | 0.701* | -0.357 | 0.751* |
|  | 60 cm | 0.05 | 0.153 | -0.398 |  | -0.651* | 0.881** | 0.680* | -0.223 | 0.679* | -0.025 | 0.086 |
| FN | 30 cm | 0.054 | 0.56* | -0.32 | -0.663* |  | -0.868** | -0.703* | 0.573* | -0.702* | 0.151 | 0.172 |
|  | 45 cm | -0.195 | 0.221 | -0.443 | -0.657* | 1 | -0.882** | -0.438 | 0.392 | -0.439 | 0.582 | -0.528* |
|  | 60 cm | -0.576* | 0.504 | -0.265 | -0.663* |  | -0.877** | -0.359 | 0.395 | -0.359 | 0.141 | 0.223 |
| FW | 30 cm | -0.09 | -0.396 | 0.07 | 0.923** | -0.871** |  | 0.927** | -0.523 | 0.928** | -0.118 | 0.22 |
|  | 45 cm | 0.074 | -0.196 | 0.117 | 0.894** | $-0.885^{* *}$ | 1 | 0.728* | -0.524 | 0.728* | -0.434 | 0.640* |
|  | 60 cm | 0.228 | -0.165 | -0.056* | 0.895** | -0.878** |  | 0.700* | -0.219 | 0.701* | -0.093 | -0.113 |
| Y/p | 30 cm | -0.34 | -0.294 | -0.131 | 0.946** | -0.708* | 0.929** |  | -0.35 | 0.999** | 0.07 | 0.313 |
|  | 45 cm | -0.285 | -0.098 | -0.236 | 0.707* | -0.441 | 0.729* | 1 | -0.291 | 0.999** | -0.011 | 0.221 |
|  | 60 cm | -0.38 | 0.322 | -0.314 | 0.687* | -0.361 | 0.701* |  | 0.016 | 1.000** | 0.334 | -0.348 |
| DM | 30 cm | -0.468 | 0.148 | -0.14 | -0.348 | 0.595* | -0.539* | -0.362 |  | -0.35 | -0.006 | -0.054 |
|  | 45 cm | -0.494 | 0.414 | -0.154 | -0.522* | 0.431 | -0.567* | -0.313 | 1 | -0.288 | 0.077 | -0.297 |
|  | 60 cm | -0.453 | 0.354 | -0.268 | -0.232 | 0.414 | -0.229 | 0.012 |  | 0.016 | -0.131 | 0.333 |
| Y/fed | 30 cm | -0.341 | -0.293 | -0.127 | 0.947** | -0.707* | 0.929** | 1.000** | -0.357 |  | 0.067 | 0.314 |
|  | 45 cm | -0.285 | -0.092 | -0.236 | 0.707* | -0.441 | 0.728* | 1.000** | -0.315 | 1 | -0.011 | 0.222 |
|  | 60 cm | -0.379 | 0.327 | -0.314 | 0.688* | -0.361 | 0.701* | 1.000** | 0.012 |  | 0.334 | -0.35 |
| LN | 30 cm | -0.038 | 0.526* | -0.502* | 0.04 | 0.157 | -0.12 | 0.07 | -0.005 | 0.068 |  | 0.335 |
|  | 45 cm | 0.18 | -0.24 | -0.222 | -0.36 | 0.585* | -0.435 | -0.011 | 0.083 | -0.011 | 1 | -0.728* |
|  | 60 cm | -0.156 | 0.692* | -0.219 | -0.029 | 0.141 | -0.092 | 0.336 | -0.143 | 0.336 |  | -0.482 |
| LA | 30 cm | 0.380 | 0.732* | -0.281 | 0.374 | 0.171 | 0.225 | 0.320 | -0.043 | 0.320 | 0.347 |  |
|  | 45 cm | -0.021 | 0.295 | -0.186 | 0.764* | -0.541* | 0.653* | 0.225 | -0.343 | 0.225 | -0.746* | 1 |
|  | 60 cm | 0.019 | 0.105 | -0.331 | 0.08 | 0.227 | -0.114 | -0.359 | 0.37 | -0.358 | -0.499 |  |

[^1]Table 4. Environmental correlations among yield and its attributes of eggplant

|  | PH | BN | FL | Dim. | FN | FW | Y kg/p | FDM | Y/fed. | LN |
| :--- | ---: | ---: | ---: | ---: | ---: | ---: | ---: | ---: | ---: | ---: | LA

PH: plant height, BN: branch number/ plant, FL: fruit length (cm), Dim: fruit diameter (cm), FN: fruit number/plant, FW: fruit weight (g), Y/p: yield (kg/plant), DM: fruit dry matter, Y/fed: yield (ton/fed.), LN: leaves number/plant, and LA: leaf area.
positive related with leaf area ( 30 cm ) and leaf number (both 30 and 60 cm intra-row spaces). Also, leaf area was significantly positive genotypic correlated with fruit diameter and average fruit weight as well as Leaves number was significantly positive related with fruit number under 45 cm intra-row spaces.

The average fruit weight was positively correlated with plant height $(60 \mathrm{~cm})$, fruit yield per plant, and fruit yield per feed (all spaces). The fruit diameter had positive significant correlation with fruit weight and yield (all spaces) as well as number of branches per plant ( 45 and 60 cm ) at both the genotypic and phenotypic levels. A highly substantial and strong correlation coefficient was found between fruit weight and yield by Islam et al. (2010), Dash et al. (2020), Khan et al. (2023) and Onyia et al. (2020). Also, Fruit diameter and fruit production per plant were found to be positively associated by Akpan et al. (2016), suggesting that selecting for these features may indirectly increase the amount of fruit produced by eggplant plants.This outcome is consistent with the findings of Danquah and Ofori (2012) who found a positive correlation between fruit diameter and fruit weight.

Fruit number is positively correlated with branches and fruit dry matter under 30 cm spacing and with leaves number in case of 45 cm . In the same time fruit number is negatively correlated with fruit diameter and fruit weight in all studied spaces. These relations of number of fruits with both fruit diameter and fruit weight were specific to genotypes and thus such associations depend on the fruit diameter. These findings are consistent with those of Prabhu and Natarajan (2007), Manna and Paul (2012) and Parihar et al. (2014), who found that yield was positively and significantly correlated with the majority of the attributes.

## Path Coefficient Analysis

To identify the specific factor causing the association, the phenotypic and genotypic correlation between yield and yield components were divided into direct and indirect effects (Tables 5 and 6 as well as Figs. 1-4). Both direct negative impacts ( -0.408 ) on yield/plant and indirect beneficial effects (number of branches, fruit diameter, and number of fruits) were attributed to plant height. The number of branches had a direct, positive impact on plant yield
(0.1613). Via plant height, fruit length, and fruit weight, it also shown detrimental indirect impacts. Fruit length shown favorable indirect impacts via fruit weight, fruit diameter, and number of branches, and a negative direct effect ( -0.552 ) on yield/plant. Fruit diameter applied negative direct effect ( -1.607 ) on yield/plant and also showed positive indirect effects via Plant height, fruit length and fruit weight (Table 5). On the other hand, fruit number showed genotypic positive direct effect (0.579) on yield/ plant under 30 cm . It also showed negative indirect effects via plant height at 30 cm ; fruit length at 45 cm and fruit weight at all planting spaces as well as positive indirect effects via corresponding other traits of 30,45 and 60 cm spacing. Fruit weight showed positive direct effect (2.984, 3.11 and -4.897) on yield/ plant and also showed negative indirect effects via number of branches in all spaces, fruit diameter and fruit number in in 30 and 45 cm , plant height in 30 cm and fruit length in both 45 and 60 cm .

Our findings and those of previous researchers have some similarities and differences, which could be related to the diverse environmental factors and breeding materials. Because of this anomalous circumstance, it was suggested that a restricted simultaneous selection model could be used to properly utilize the direct benefit while eliminating the unwanted indirect consequences. After examining the significant correlation and the desired direct effect of fruit weight, number of branches, and number of fruits on yield/plant, it is possible to draw the conclusion that these parameters could be taken into account for the creation of inbred lines through pure line selection in subsequent generations, or for the development of elite hybrids through heterosis breeding.

There have been prior reports providing supporting evidence of a direct positive relationship between the quantity of fruits on a plant and its yield (Singh et al., 2006; Rani et al., 2008; Eslam et al., 2010; Alam and Paul, 2019; Khan et al., 2023). The outcome agreed with the findings of (Hayadar et al., 2007; Saleem et al., 2013). Noticed a positive direct relationship between plant height and yield/plant, in contrast to Ghosh et al. (2010) finding of a negative relationship between plant height and yield/plant in tomatoes. To increase yield, direct selection of these qualities would be beneficial.

Table 5. Path coefficient analysis of fruit yield/plant (Y/p) vs plant height, branches number, fruit length, fruit diameter, average fruit weight and fruits number/plant at 30,45 and 60 cm planting spaces

| Pathways of association | 30 cm | 45 cm | 60 cm |
| :---: | :---: | :---: | :---: |
| Y/P vs. Plant height (PH) |  |  |  |
| Direct effect (P17) | -0.408 | -0.311 | -2.815 |
| Indirect effect via BN (P27*r12) | 0.061 | -0.001 | -1.875 |
| Indirect effect via FL (P37*r13) | -0.081 | 0.049 | 0.972 |
| Indirect effect via Dim. (P47*r14) | 0.325 | 0.045 | 0.031 |
| Indirect effect via FN (P57*r15) | 0.032 | -0.297 | 4.424 |
| Indirect effect via FW (P67*r16) | -0.269 | 0.23 | -1.117 |
| Total (r) | -0.34 | -0.285 | -0.38 |
| Y/P vs. Number of branches (BN) |  |  |  |
| Direct effect (P27) | 0.161 | 0.304 | 5.341 |
| Indirect effect via PH (P17*r12) | -0.153 | 0.001 | 0.988 |
| Indirect effect via FL (P37*r23) | -0.008 | -0.061 | -3.038 |
| Indirect effect via Dim. (P47*r24) | 0.563 | -0.069 | 0.094 |
| Indirect effect via FN (P57*r25) | 0.324 | 0.337 | -3.871 |
| Indirect effect via FW (P67*r26) | -1.181 | -0.61 | 0.808 |
| Total (r) | -0.294 | -0.098 | 0.322 |
| Y/P vs. Fruit length (FL) |  |  |  |
| Direct effect (P37) | -0.552 | 0.108 | 3.328 |
| Indirect effect via PH (P17*r13) | -0.06 | -0.14 | -0.823 |
| Indirect effect via BN (P27*r23) | 0.002 | -0.18 | -4.876 |
| Indirect effect via Dim. (P47*r34) | 0.455 | 0.287 | -0.251 |
| Indirect effect via FN (P57*r35) | -0.185 | -0.675 | 2.035 |
| Indirect effect via FW (P67*r36) | 0.209 | 0.364 | 0.273 |
| Total (r) | -0.131 | -0.236 | -0.314 |
| Y/P vs. Fruit diameter (Dim) |  |  |  |
| Direct effect (P47) | -1.607 | -1.076 | 0.625 |
| Indirect effect via PH (P17*r14) | 0.083 | 0.013 | -0.141 |
| Indirect effect via BN (P27*r24) | -0.056 | 0.02 | 0.817 |
| Indirect effect via FL (P37*r34) | 0.156 | -0.029 | -1.324 |
| Indirect effect via FN (P57*r45) | -0.384 | -1.002 | 5.093 |
| Indirect effect via FW (P67*r46) | 2.754 | 2.781 | -4.383 |
| Total (r) | 0.946 | 0.707 | 0.687 |
| Y/P vs. Number of fruits (FN) |  |  |  |
| Direct effect (P57) | 0.579 | 1.525 | -7.679 |
| Indirect effect via PH (P17*r15) | -0.022 | 0.061 | 1.622 |
| Indirect effect via BN (P27*r25) | 0.09 | 0.067 | 2.692 |
| Indirect effect via FL (P37*r35) | 0.177 | -0.048 | -0.882 |
| Indirect effect via Dim. (P47*r45) | 1.066 | 0.707 | -0.414 |
| Indirect effect via FW (P67*r56) | -2.598 | -2.753 | 4.3 |
| Total (r) | -0.708 | -0.441 | -0.361 |
| Y/P vs. Average fruit weight (FW) |  |  |  |
| Direct effect (P67) | 2.984 | 3.11 | -4.896 |
| Indirect effect via PH (P17*r16) | 0.037 | -0.023 | -0.642 |
| Indirect effect via BN (P27*r26) | -0.064 | -0.06 | -0.881 |
| Indirect effect via FL (P37*r36) | -0.039 | 0.013 | -0.186 |
| Indirect effect via Dim. (P47*r46) | -1.484 | -0.962 | 0.562 |
| Indirect effect via FN (P57*r56) | -0.505 | -1.349 | 6.744 |
| Total (r) | 0.929 | 0.729 | 0.701 |

Table 6. Partitioning of the phenotypic and genotypic correlation coefficients into direct effects (main diagonal, bold) and indirect effects (above and below the main diagonal) in ten eggplant genotypes

| Item |  | PH | BN | FL | Dim | FN | FW | Correlation |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Phenotypic level |  |  |  |  |  |  |  |  |
| PH | 30 cm | -0.299 | 0.063 | -0.024 | 0.048 | 0.014 | -0.121 | -0.320 |
|  | 45 cm | -0.185 | 0.001 | -0.092 | 0.026 | -0.185 | 0.162 | -0.273 |
|  | 60 cm | -0.485 | -0.034 | -0.026 | -0.026 | -0.145 | 0.341 | -0.375 |
| BN | 30 cm | -0.116 | 0.163 | 0.001 | 0.062 | 0.100 | -0.416 | -0.205 |
|  | 45 cm | 0.015 | -0.008 | 0.080 | -0.010 | 0.138 | -0.266 | -0.052 |
|  | 60 cm | 0.068 | 0.240 | 0.055 | -0.065 | 0.083 | -0.174 | 0.208 |
| FL | 30 cm | -0.038 | -0.001 | -0.186 | 0.070 | -0.073 | 0.100 | -0.128 |
|  | 45 cm | -0.080 | 0.003 | -0.213 | 0.206 | -0.412 | 0.262 | -0.234 |
|  | 60 cm | -0.136 | -0.143 | -0.093 | 0.214 | -0.067 | -0.085 | -0.311 |
| Dim | 30 cm | 0.056 | -0.040 | 0.051 | -0.255 | -0.149 | 1.277 | 0.940 |
|  | 45 cm | 0.006 | 0.000 | 0.057 | -0.763 | -0.607 | 2.008 | 0.701 |
|  | 60 cm | -0.023 | 0.028 | 0.036 | -0.549 | -0.167 | 1.355 | 0.680 |
| FN | 30 cm | -0.018 | 0.072 | 0.059 | 0.167 | 0.228 | -1.211 | -0.703 |
|  | 45 cm | 0.037 | -0.001 | 0.094 | 0.496 | 0.934 | -1.998 | -0.438 |
|  | 60 cm | 0.273 | 0.078 | 0.024 | 0.358 | 0.257 | -1.349 | -0.359 |
| FW | 30 cm | 0.026 | -0.049 | -0.013 | -0.234 | -0.198 | 1.394 | 0.927 |
|  | 45 cm | -0.013 | 0.001 | -0.025 | -0.677 | -0.824 | 2.265 | 0.728 |
|  | 60 cm | -0.108 | -0.027 | 0.005 | -0.483 | -0.225 | 1.538 | 0.700 |
| Genotypic level |  |  |  |  |  |  |  |  |
| PH | 30 cm | -0.408 | 0.060 | -0.081 | 0.325 | 0.031 | -0.268 | -0.340 |
|  | 45 cm | -0.311 | -0.001 | 0.048 | 0.045 | -0.298 | 0.231 | -0.285 |
|  | 60 cm | -2.816 | -1.874 | 0.971 | 0.031 | 4.426 | -1.118 | -0.380 |
| BN | 30 cm | -0.153 | 0.161 | -0.008 | 0.563 | 0.325 | -1.182 | -0.294 |
|  | 45 cm | 0.001 | 0.304 | -0.063 | -0.069 | 0.337 | -0.609 | -0.098 |
|  | 60 cm | 0.988 | 5.341 | -3.039 | 0.095 | -3.870 | 0.807 | 0.322 |
| FL | 30 cm | -0.060 | 0.002 | -0.552 | 0.455 | -0.186 | 0.210 | -0.131 |
|  | 45 cm | -0.139 | -0.179 | 0.108 | 0.287 | -0.676 | 0.363 | -0.236 |
|  | 60 cm | -0.821 | -4.878 | 3.328 | -0.248 | 2.033 | 0.273 | -0.314 |
| Dim | 30 cm | 0.083 | -0.057 | 0.156 | -1.607 | -0.384 | 2.756 | 0.946 |
|  | 45 cm | 0.013 | 0.019 | -0.029 | -1.076 | -1.002 | 2.781 | 0.707 |
|  | 60 cm | -0.140 | 0.816 | -1.323 | 0.625 | 5.090 | -4.381 | 0.687 |
| FN | 30 cm | -0.022 | 0.090 | 0.177 | 1.066 | 0.579 | -2.599 | -0.708 |
|  | 45 cm | 0.061 | 0.067 | -0.048 | 0.707 | 1.525 | -2.753 | -0.441 |
|  | 60 cm | 1.622 | 2.691 | -0.881 | -0.414 | -7.681 | 4.302 | -0.361 |
| FW | 30 cm | 0.037 | -0.064 | -0.039 | -1.484 | -0.505 | 2.984 | 0.929 |
|  | 45 cm | -0.023 | -0.060 | 0.013 | -0.962 | -1.350 | 3.110 | 0.729 |
|  | 60 cm | -0.643 | -0.880 | -0.186 | 0.559 | 6.748 | -4.897 | 0.701 |



Fig. 1. Genotypic path coefficient diagram representing cause and effect relationships among some vegetative and fruit traits with yield/plant [PH (1): plant height; BN (2): branches number; FL (3): fruit length; Dim. (4): fruit diameter; FN (5): number of fruits per plant; FW (6): average fruit weight)] at 30 cm (Upper) and 45 cm planting spaces (Down)


Fig. 2. Average genotypic path coefficient diagram representing cause and effect relationships among some vegetative and fruit traits with yield/ plant [(PH (1): plant height; BN (2): branches number; FL (3): fruit length; Dim (4): fruit diameter; FN (5): number of fruits per plant; FW (6): average fruit weight] at 60 cm planting spaces


Fig. 3. Percentage contribution of fruit yield and its attributing traits based on path coefficient analysis for three intra-row spacing.


Fig. 4. Relative importance ( $\mathbf{R \%}$ ) according to path analysis of fruit yield/plant and its attributed traits in eggplant at 30, 45 and 60 cm planting spaces

Consequently, these traits may be taken into account while choosing genotypes to increase yield. Conversely, in all spaces (phenotypic and/or genotypic level), fruit diameter, fruit length, and plant height had a negative direct effect on yield. All characters' negative direct effects, however, were offset by their indirect effects through other characters, leading to a highly substantial positive association with yield in the end. The direct effect points out that, with other variables held constant, increasing any of these traits will decrease yield. However, the more suitable indirect effects play a more important part and mask the direct influence (Dewey and Lu, 1959). A positive phenotypic (1.277 and 2.008) and genotypic (2.756 and 2.781) effects of 30 and 45 cm , respectively (Table 6), as well as the phenotypic of $60 \mathrm{~cm}(1.355)$ were recorded indirectly by fruit diameter on yield per plant through fruit weight due to fact the diameter was positively correlated $\left(\mathrm{r}=0.916^{* *}\right.$ and $\left.0.923^{* *}\right),\left(0.887^{* *}\right.$ and $\left.0.894^{* *}\right)$ and ( $0.881^{* *}$ and $0.895^{* *}$ ) in 30,45 and 60 cm spaces, respectively with fruit weight (Table 2 and Figs. 1,2 and 3 ) which in turn had a large direct effects (1.394 and 2.984), (2.265 and 3.110) upon yield in 30 and 45 cm , respectively as well as 1.538 in phenotypic level of 60 cm .

Relative importance ( $\mathrm{R} \%$ ) according to path analysis of fruit yield/plant and its attributed traits in eggplant at 30,45 and 60 cm planting spaces are shown in Fig. 5. The results revealed that the greatest parts of yield/plant variation were explained by the direct effect for fruit weight at 30 cm (32.2\%) and 45 cm (30.9\%) planting spaces followed by fruit diameter (9.3\%) and both number of fruits/ plant (7.4\%) and fruit diameter (3.7\%) at 30 and 45 cm planting spaces, respectively. As for 60 cm spaces, the direct effect of number fruits/plant ( $16.6 \%$ ) exhibited the highest part of yield/ plant variation followed by number of branches (8\%) and average fruit weight ( $6.7 \%$ ). These traits showed significant contributions to plant yield demonstrate their importance as selection factors in eggplant breeding programs. The other traits, however, noted insignificant or nonexistent direct effects on fruit yield.

Regarding the relative importance of joint effects, it is obvious that their effective parts were obtained by fruit diameter on yield/ plant through its associations with each of number of fruits/plant ( 4.5 and $6.9 \%$ ) and average of fruit weight ( 32 and $19.1 \%$ ) as well as fruit number via its association with average fruit weight
(10.9 and $26.8 \%$ ) at 30 and 45 cm planting spaces, respectively. As for 60 cm planting space, the effective parts of the relative importance of joint effects were obtained by fruit number via its association with average fruit weight (18.6\%), number of branches through its associations with each of number of fruits/plant (11.6\%), fruit length ( $9.1 \%$ ) and average of fruit weight ( $2.1 \%$ ) as well as by plant height via both fruit number ( $7 \%$ ) and branches (3\%) and fruit length through its association with fruit number (3.8\%). The highest value of the indirect effects was recorded by fruit diameter via average of fruit weight ( $32 \%$ ) at 30 cm space and fruit number via fruit weight (26.8 and 18.6\%) at 45 and 60 cm spaces, respectively. Small values of relative importance ranging from 0.0 to $1.2 \% ~(45 \mathrm{~cm})$ and $1.8 \%$ ( 30 and 60 cm ) were obtained by the other direct and indirect effects.

The picture was different under various spaces where, total direct effects decreased with increasing plant spaces from 30 cm ( $44.6 \%$ ), 45 $\mathrm{cm}(42.6 \%)$ to $60 \mathrm{~cm}(36.8 \%)$ and reverse trend was observes for total indirect effects which increased with increasing plant spaces; i.e., 55.5 $(30 \mathrm{~cm}), 57(45 \mathrm{~cm})$ and $63.4 \%(60 \mathrm{~cm})$ as shown in Figs. 3 and 4.

It is clear from the present work that the correlation analysis gave a different picture of the role of number of fruit per plant, branches and fruit diameter on fruit yield than that the given by path coefficient analysis. Consequently, it is suggested to use indirect selection using other component qualities that also have similar traits and have good indirect impacts in order to increase yield. At the phenotypic level, it was shown that the majority of the direct effects were smaller than one, suggesting that phenotypic inflation resulting from multi-collinearity was negligible. Totally, the studied characters accounted for $100.02,99.57$ and $100.14 \%$ of fruit yield/ plant variability. The residual content ( 0.02 , 0.43 and $0.14 \%$ ) may be returned to unknown factors (random error) and/or some other traits that were not included in the present study.

## Conclusion

In conclusion: There is a wide range of phenotypic diversity for morphological, yield and quality traits among the genotypes. Some
genotypes produced higher yields and desirable traits compared to others. Almost all traits showed strong associations with other agronomic and yield traits, and higher values of expected genetic advances (GAM\%) were observed for those traits, indicating higher genetic gains in the breeding programme. In general, it is proposed to expand the genetic base of the existing Egyptian germplasm using these genotypes to develop a successful and sustainable breeding programme.

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# در اســـــة الاختــلافــــات الـور اثية و الارتباط في بعض التر اكيب الور اثية للباذنجان 

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يعد تقنير الاختلافات الور اثثة هام لتطوير تر اكيب ور اثثة جديدة ذات صفات محصولية وجودة مرنفــة، ويسـاعد مربي

 بـلاك لـونج (BVL)، أسـود (As)، جابـانيز و ايـت ايـج (JWE)، ابـل جـرين (AG)، روتونــــا بيانكـا سـتوماتا دي روز ا (RSB)، كورين ريد (KRL) و بلاك اوبلنج (BO)، تحت ثلاثثة مسـافات بين النباتات داخل الخط ( 30، 45 و 60 سم)، وأجريت هذه الدر اسة تحت ظروف منطقة الزقازيق - مصر أثنـاء الموسمي الصيف 2017 و 2018 ، وتم حسـاب النقدم الور اثي المتوقع والإرتباط ومعامل المرور للصفات تحت اللار اســة. أظهرت النتـائج وجود إختنالفات عاليـة المعنويـة بين
 و انخفضت مع زيادة المسافات في جميع الصفات المدروسة، وكانت نسبة درجة النوريث بالمعني الو اسع (h ${ }^{\text {الو }}$ ( ${ }^{2}$ ) لصفة عدد الأفر ع (39 و \% \% \%) تحت مسـافات زر اعة 45 و 60 سم علي التو الي، كان اللتقام الور اثي كنسبة مئوية (\%GA) مر تفعا لجميع الصفات المدروسـة باستثناء عدد الأفر ع، وكانت قيم GCV/PCV عاليـة في معظـ الصـفات المدروسـة لجميع المسافات، أظهرت التز اكيب الور اثثة فيم عاليـة لـقـدار التحسين الور اثي لصفات المحصـول وجودة الثمـار ، أظهر معامل الإرتباط الور اثي إرتباط قوي بين الصفات المحصولية و المورفولوجية. أظهرت نتائج تحليل معامل المرور أن صفة وزن الثمـار كـان لها التأتُثر المباثشر الأكبر على محصول النبات (2.984 و 3.11) يليها صفات عدد الثمـار / النبـات (0.579 و 1.525)، و عدد الأفر ع/نبات (0.161 و0.304) عند مسافات الزر اعة 30 و 45 سم ، علي النو الي. في حين أعطت صفة
 (0.625) عند مسـافة الزر راعة 60 سم، ممـا يدل على فعاليـة الانتخـاب المباشر لهذه الصفات في تحسين و إنتاج تر اكيب ور اثية عالية المحصول تحت كثافات زر اعة مختلفة.


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[^1]:    ** Significant at $1 \%$ level of significance; * Significant at 5\% level of significance.
    PH: plant height, BN: branch number/plant, FL: fruit length ( cm ), Dim: fruit diameter ( cm ), FN: fruit number/ plant, FW: fruit weight (g), Y/p: yield (kg/plant), DM: fruit dry matter, Y/fed: yield (ton/fed.), LN: leaves number/plant, and LA: leaf area.

