

Nigeria

Correlation and Path Coefficient Analysis in Tomato (Solanum lycopersicon L. Karst) under Fruit Worm (Heliothis zea Buddie) Infestation in a Line × Tester

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## MATERIALS AND METHODS

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# ABSTRACT

## **OBJECTIVES**

## RESULTS

| TRAITS       | Trich<br>Count | Fl.<br>Cluster | Leaves<br>/Plant | Plant<br>Ht | Fruits<br>/Plant | Yield<br>/Plant | % Dam.<br>Fruits | Days to<br>Harv |
|--------------|----------------|----------------|------------------|-------------|------------------|-----------------|------------------|-----------------|
| Trich. Count | 1.00           | 0.90**         | 0.90**           | 0.69        | 0.04             | 0.16            | 0.92**           | 0.02            |
| Fl. Clusters | 0.90**         | 1.00           | 0.92**           | 0.91**      | 0.99**           | -0.91**         | -0.92**          | 0.92**          |
| Leaves/Plant | 0.90**         | 0.90**         | 1.00             | 0.91**      | 0.99**           | -0.91**         | -0.92**          | -0.91**         |
| Plant Ht.    | 0.71**         | 0.81**         | 0.99**           | 1.00        | 0.95**           | 0.56            | 0.91**           | 0.91**          |
| Fruits/Plant | 0.91**         | 0.91**         | 0.91**           | 0.90**      | 1.00             | 0.92**          | 0.97**           | 0.99**          |
| % Dam. Fruit | 0.22           | 0.22           | 0.29             | -0.10       | 0.91**           | 1.00            | -0.91**          | -0.90**         |
| Days to Harv | -0.90**        | -0.90**        | -0.90**          | 0.66        | 0.92**           | -0.32           | 1.00             | -0.92**         |
|              | 0.90**         | -0.81**        | -0.91**          | 0.57        | 0.92**           | -0.07           | -0.77**          | 1.00            |

| TRAITS        | Trich.<br>Count | Fl.<br>Cluster | Leaves<br>/Plant | Plant<br>Ht. | Fruits<br>/Plant | % Dam.<br>Fruits | Days to<br>Harv | Corr.<br>Coef. |
|---------------|-----------------|----------------|------------------|--------------|------------------|------------------|-----------------|----------------|
| Trich. Count  | 0.092           | 0.631          | 0.239            | -0.001       | 0.047            | 0.991            | 0.010           | 0.22           |
| Fl. Cluster   | 0.087           | 0.651          | 0.315            | -0.030       | 0.267            | -0.991           | 0.440           | 0.22           |
| Leaves/Plant  | 0.087           | 0.586          | 0.346            | -0.027       | -0.290           | -0.991           | -0.435          | 0.29           |
| Plant Ht.     | 0.069           | 0.527          | -0.030           | -0.030       | -0.279           | -0.980           | 0.435           | -0.10          |
| Fruits/ Plant | 0.088           | 0.592          | 0.315            | 0.027        | -0.293           | -0.945           | 0.473           | 0.91**         |
| % Dam. Fruit  | -0.087          | -0.586         | -0.311           | 0.020        | 0.270            | 0.977            | 0.440           | -0.32          |
| Days to Harv. | 0.087           | -0.527         | -0.315           | -0.017       | 0.270            | 0.829            | 0.478           | -0.07          |

#### DISCUSSIONS

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## CONCLUSIONS

Vield contributing traits such as flower clusters/plant, leaves/plant and plant height were found to have positive and significant genotypic and phenotypic correlations with number of fruits/plant. The resistance trait i.e. trichome count was genotypically correlated with flower clusters/plant, leaves/plant and % damaged fruits. Number of fruits/plant were phenotypically and positively correlated with flower clusters, leaves/plant, plant height and days to final harvest. Trichome count was significantly and negatively phenotypically correlated with number of fruits/plant. The % damaged fruit exhibited the highest positive direct effect on fruit yield. The direct effects of flower clusters/plant, days to final harvest, and leaves/plant were positive on fruit yield. These traits could be exploited concurrently when selecting for high yielding genotypes in tomato.

#### **ACKNOWLEDGEMENTS**

