Variability and genetic divergence in paprika (*Capsicum annuum* L.)

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**Abstract**

Ninety four paprika (*Capsicum annuum*) accessions were evaluated for 17 characters for variability, heritability, genetic advance and genetic divergence at Lam, Guntur (Andhra Pradesh). Higher phenotypic and genotypic coefficient of variation (PCV and GCV) and heritability coupled with high genetic advance was observed for number of fruits plant\(^{-1}\), fresh fruit yield plant\(^{-1}\), dry fruit yield plant\(^{-1}\), 100 seed weight, number of seeds fruit\(^{-1}\) and capsanthin, capsaicin and oleoresin contents indicating the higher magnitude of variability for these traits and consequently more scope for their improvement through selection. Plant height, plant spread and fruit diameter exhibited moderate PCV and GCV estimates suggesting the possible role of environment in expression of these characters. Cluster analysis through Mahalanobis D\(^2\) analysis classified the 94 genotypes into 10 clusters indicating considerable genetic diversity in the material studied.

**Keywords:** *Capsicum annuum*, genetic advance, genotypic coefficient of variation, heritability, paprika, phenotypic coefficient of variation.

There is need to develop a long term strategy for development of paprika (*Capsicum annuum* L.) in view of its increasing demand in the domestic and export sectors. Hence a genetic divergence study which is a basic requirement for effective selection, was undertaken along with an attempt to estimate genetic variability, heritability and genetic advance in the available germplasm of paprika at Andhra Pradesh Horticultural University, Horticultural Research Station, Lam, Guntur (Andhra Pradesh).

The material chosen for the investigation was a set of 94 paprika genotypes comprising of established varieties, advanced true breeding lines and local collections, collected and developed at Horticultural Research Station, Lam, Guntur. The experiment was laid out in a completely randomized block design with 94 paprika accessions as treatments during *kharif* 2005. Each treatment was randomly replicated thrice. Each treatment or a genotype in each replication was represented by a plot of four rows each of 5 m length. The crop

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\(^1\)Part of PhD thesis submitted by the senior author to ANGRAU, Hyderabad, India.
<table>
<thead>
<tr>
<th>Character</th>
<th>Range</th>
<th>Genotypic variance</th>
<th>Phenotypic variance</th>
<th>Genotypic coefficient of variation (%)</th>
<th>Phenotypic coefficient of variation (%)</th>
<th>Broad sense heritability (%)</th>
<th>Genetic advance (%)</th>
<th>GA as per cent mean (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Plant height (cm)</td>
<td>95.20 - 128.8</td>
<td>199.60</td>
<td>210.20</td>
<td>14.84</td>
<td>15.23</td>
<td>95.0</td>
<td>28.37</td>
<td>29.79</td>
</tr>
<tr>
<td>Plant spread (cm)</td>
<td>124.9 - 172.8</td>
<td>677.20</td>
<td>694.60</td>
<td>20.83</td>
<td>21.09</td>
<td>97.5</td>
<td>52.94</td>
<td>42.37</td>
</tr>
<tr>
<td>Days to 50% flowering</td>
<td>75 - 82</td>
<td>15.54</td>
<td>17.86</td>
<td>5.24</td>
<td>5.62</td>
<td>87.02</td>
<td>7.57</td>
<td>10.01</td>
</tr>
<tr>
<td>Days to maturity</td>
<td>110 - 132</td>
<td>81.63</td>
<td>85.11</td>
<td>8.20</td>
<td>8.37</td>
<td>95.91</td>
<td>18.22</td>
<td>16.34</td>
</tr>
<tr>
<td>Fruits plant¹</td>
<td>126 - 255</td>
<td>1674.31</td>
<td>1712.15</td>
<td>32.46</td>
<td>32.82</td>
<td>97.80</td>
<td>83.36</td>
<td>66.13</td>
</tr>
<tr>
<td>Fruit length (cm)</td>
<td>8.61 - 13.17</td>
<td>3.96</td>
<td>4.40</td>
<td>22.69</td>
<td>24.37</td>
<td>86.68</td>
<td>3.74</td>
<td>43.50</td>
</tr>
<tr>
<td>Fruit girth (cm)</td>
<td>2.32 - 4.6</td>
<td>1.77</td>
<td>2.02</td>
<td>28.63</td>
<td>30.61</td>
<td>87.50</td>
<td>2.56</td>
<td>51.16</td>
</tr>
<tr>
<td>Fruit shape index</td>
<td>3.70 - 8.8</td>
<td>0.55</td>
<td>0.65</td>
<td>36.97</td>
<td>40.07</td>
<td>85.10</td>
<td>1.42</td>
<td>70.28</td>
</tr>
<tr>
<td>Fresh fruit wt plant¹ (g)</td>
<td>517.65 - 1195.5</td>
<td>33819</td>
<td>33892</td>
<td>35.53</td>
<td>35.56</td>
<td>99.80</td>
<td>378.42</td>
<td>73.10</td>
</tr>
<tr>
<td>Dry fruit wt plant¹ (g)</td>
<td>160.37 - 91.04</td>
<td>2412</td>
<td>2465</td>
<td>30.62</td>
<td>30.96</td>
<td>97.82</td>
<td>100.05</td>
<td>62.39</td>
</tr>
<tr>
<td>Recovery % of fresh to dry</td>
<td>31.84 - 46.3</td>
<td>25.06</td>
<td>27.66</td>
<td>15.73</td>
<td>16.52</td>
<td>90.60</td>
<td>9.82</td>
<td>30.83</td>
</tr>
<tr>
<td>Wt. of dry stalkless chilles (g)</td>
<td>124.55 - 1271.4</td>
<td>1244.4</td>
<td>1271.40</td>
<td>28.32</td>
<td>28.63</td>
<td>97.90</td>
<td>71.89</td>
<td>57.72</td>
</tr>
<tr>
<td>Seeds fruit¹</td>
<td>71.52 - 188.33</td>
<td>894.13</td>
<td>915.13</td>
<td>41.81</td>
<td>42.30</td>
<td>97.71</td>
<td>60.89</td>
<td>85.13</td>
</tr>
<tr>
<td>100 seed wt. (g)</td>
<td>0.45 - 0.15</td>
<td>0.07</td>
<td>0.08</td>
<td>60.11</td>
<td>61.80</td>
<td>94.63</td>
<td>0.54</td>
<td>120.46</td>
</tr>
<tr>
<td>Oleoresin (%)</td>
<td>8.14 - 17.34</td>
<td>8.93</td>
<td>9.00</td>
<td>36.71</td>
<td>36.86</td>
<td>99.20</td>
<td>6.13</td>
<td>75.32</td>
</tr>
<tr>
<td>Capsanthin (EOA)</td>
<td>33625 - 88733</td>
<td>388379003 - 392060600</td>
<td>58.61</td>
<td>58.89</td>
<td>99.10</td>
<td>40406</td>
<td>120.20</td>
<td></td>
</tr>
<tr>
<td>Capsaicin (%)</td>
<td>0.18 - 0.44</td>
<td>0.005</td>
<td>0.005</td>
<td>39.83</td>
<td>40.31</td>
<td>97.62</td>
<td>0.15</td>
<td>81.05</td>
</tr>
</tbody>
</table>
received timely management practices as per the package of practices and recommendations of the Andhra Pradesh Horticultural University. Five plants were randomly selected per accession and observations were recorded on plant height, plant spread, days to 50% flowering, days to maturity, number of fruits plant\(^{-1}\), fruit length, fruit girth, fruit shape index, fresh fruit yield plant\(^{-1}\), dry fruit plant\(^{-1}\), weight of dry stalk-less chillies plant\(^{-1}\), number of seeds fruit\(^{-1}\), 100 seed weight and oleoresin, capsanthin and capsaicin contents. Analysis of variance in respect of various characters was done (Panse & Sukhatme 1967). Genetic variability for different characters was estimated as suggested by Singh & Choudary (1985). Heritability (broad sense) and genetic advance as percentage of mean were calculated as per Hanson et al. (1956) and Johnson et al. (1955), respectively. Assessment of genetic diversity in classifying the genotypes into different clusters was done by Mahalanobis D\(^2\) analysis (1928).

The analysis of variance in the 94 paprika genotypes indicated highly significant differences among the genotypes for all the 17 quantitative and qualitative characters studied (Table 1). The genotype LCA 432 recorded the highest number of fruits plant\(^{-1}\). The highest dry fruit yield plant\(^{-1}\) was recorded in LCA 427. The highest capsanthin content was recorded in LCA 424 and highest oleoresin content in KTPL 19.

The phenotypic coefficient of variation (PCV) in general, was higher than genotypic coefficient of variation (GCV) for all the traits, but the differences were very narrow indicating low environmental influence on the expression and are suggestive of the heritable nature. Similar results were reported by Chatterjee (2006) in chilli. Higher PCV and GCV were observed for number of fruits plant\(^{-1}\), fresh fruit yield, dry fruit yield, 100 seed weight, number of seeds fruit\(^{-1}\) and capsanthin, capsaicin and oleoresin contents indicating the higher magnitude of variability for these traits and consequently more scope for their improvement through selection.

Plant height, spread and fruit diameter exhibited moderate PCV and GCV estimates suggesting the possible role of environment in influencing these characters. Similar results were reported by other workers also (Gogoi & Gautam 2002; Rathod et al. 2002; Wasule et al. 2004; Mishra et al. 2005). Days to 50% flowering and days to maturity recorded low PCV and GCV suggesting limited variability indicating need to generate more variability for wider spectrum of selection (Chatterjee 2006).

The effectiveness of the selection depends upon genetic advance of the character selected along with heritability. The study revealed high heritability coupled with high genetic advance for several biometric characters including yield plant\(^{-1}\), fruits plant\(^{-1}\), fruit weight, number of seeds fruit\(^{-1}\), 100 seed weight and oleoresin, capsanthin and capsaicin contents indicating the effect of additive genes. These characters could be considered reliable for effective selection in paprika. High heritability and low genetic advance was observed for days to 50% flowering and days to maturity indicating that these characters are controlled by non-additive genes and improvement through selection for these characters will not be very much effective (Chatterjee 2006; Choudhary & Samadia 2004).

Based on D\(^2\) values, the 94 genotypes were grouped into 10 clusters (Table 2) which indicated considerable genetic diversity in the material studied. The largest was cluster I which comprised of 24 genotypes. The cluster distances ranged from 15789.6 (between cluster II and cluster X) to 856.7 (between cluster 1 and cluster II) (Table 3) (Sreelathakumary & Rajamony 2004; Varalakshmi & Babu 1991).

On the basis of present study, yield plant\(^{-1}\), fruits plant\(^{-1}\), number of seeds fruit\(^{-1}\), 100 seed weight and oleoresin, capsanthin and capsaicin contents were the most important quantitative and qualitative characters to be taken into consideration for effective selection in paprika. It is desirable to attempt crosses.
between desirable genotypes belonging to different clusters with desirable genotypes for getting highly heterotic crosses.

**References**

Chatterjee B 2006 Character association and genetic divergence in chilli (*Capsicum annuum* L.) MSc Thesis, Acharya N G Ranga Agricultural University, Hyderabad.


