

## Combining ability for seed yield and its components in sesame (*Sesamum indicum* L.)

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### SUMMARY

Hybrids were synthesized in a 10 x 10 diallel fashion excluding reciprocals and analysis of combining ability was undertaken for yield and its component traits in sesame. Non-additive genetic variance was of greater importance for plant height, length of main branch, number of capsules on main branch, number of capsules per plant, capsule length, number of seeds per capsule, seed yield per plant and harvest index. Based on general combining ability, parents C 1013, AT 123 and GT 2 were found to be good general combiners for improving yield and yield contributing traits. The cross combinations viz., GT 2 x GT 10, Mrug 1 x PT 64 and C 1013 x ABT 23 had highly significant sca effect for seed yield and most of the yield attributing traits.

**Key words :** Diallel, Combining ability, Sesame

Sesame an ancient oilseed crop of India is known as “Queen of oilseeds” by virtue of its oil quality. Rajasthan is the major sesame growing state (314.9 thousand ha. area) in India with a production of 129.9 thousand tones but the productivity (403 kg/ha) is very low (Anonymous, 2007-08).

Combining ability analysis is considered a very useful technique in classifying parental lines according to their potential to yield good hybrids and to aid in selecting parents which when crossed could give rise to better segregants in later generations. In the present study, an attempt has been made to examine the combining ability of some sesame accessions and their hybrids over environments.

### MATERIALS AND METHODS

Genetic material for the present investigation comprised of 10 sesame genotypes viz., GT 1, GT 2, C 1013, Mrug 1, GT 10, PT 64, Tapi, ABT 23, ABT 26 and AT 123. The 10 parents and 45 hybrids were raised in randomized block design with 3 replications under 4 environments created by four different dates of sowing during *Kharif*-2005. The field experiment was conducted at the Main Castor and Mustard Research Station, S. D. Agricultural University, Sardarkrushinagar with 45 x 15

cm spacing. All the agronomic practices were followed uniformly. The observations were recorded on 5 randomly selected plants of each genotype in each replication and each environment for different characters. The data were analyzed for combining ability for single environment using Griffing (1956), Model 1 and Method II. Griffing method of combining ability was later elaborated by Daljit Singh (1973, 1979) which was utilized for combining ability for pooled over environments.

### RESULTS AND DISCUSSION

The pooled analysis of variance for combining ability revealed presence of significant mean squares due to gca and sca for all the characters under study (Table 1), thereby suggesting that both additive and non additive gene actions were important for the expression of these traits. The mean squares due to interaction of environments with gca and sca were also significant for all the characters under study. However, the ratio of  $s^2_{gca}/s^2_{sca}$  variance showed that non additive type gene effects were more important in the expression of all the characters under study except for days to 50 per cent flowering, days to maturity and number of effective branches per plant.

Three parental lines viz., C 1013, AT 123 and GT 2 were found to be good general combiners for seed yield along with a number of its component traits (Table 2). Among these, C 1013 was also found to be best general combiner for days to maturity, number of effective branches per plant, number of capsules per plant, capsule length and harvest index. The parent, AT 123 manifested desirable and significant gca effects for plant height, length of main branch, number of effective branches per plant, number of capsules per plant, capsule length and number

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