DIALLEL CROSS ANALYSIS FOR YIELD AND ITS COMPONENTS IN SIX FLAX GENOTYPES

Maysa S. Abd Al-Sadek

ABSTRACT

This study was conducted with the objective of estimating combining ability and gene action for both of seed and straw yields as with as their components in six flax genotypes. This was achieved via evaluating six parents ($P_1$ = Giza 8, $P_2$ = Sakha 2, $P_3$ = S.413/1/2/3, $P_4$ = S.420/12, $P_5$ = Cizar and $P_6$ = Bombay) with their 15 $F_2$ progenies. In 2014/2015 season, the six parents and their 15 $F_2$ crosses were planted in a randomized complete block design with three replications at Giza Res. Station Farm.

The collected data indicated that the additive effects were more important than non-additive effects for each of straw yield, plant height, technical stem length, seed yield, no. of capsules per plant, 1000-seed weight and no. of seeds per capsule indicating that effective selection should be possible within these $F_2$ and subsequent generations for these characters. While, both additive and non-additive effects were the same almost effect in inheritance of no. of basal branches per plant.

$P_3$ and $P_4$ showed high general combining ability for each of straw yield per plant, plant height and technical stem length. Also, $P_1$ proved to be good combiner for seed yield and its two important components, no. of capsules per plant and 1000-seed weight in addition $P_5$ exhibited high general combiner for all seed yield characters except 1000-seed weight, suggesting the importance of these parents for increasing the previous traits in flax breeding programs. The simple correlation between GCA values and parental means for all characters studied were significantly positive. These results indicated that the parents showing higher mean performance proved to be the highest general combiners for these traits. One cross ($P_3$x$P_6$) exhibited significant and positive SCA effect for each of straw yield, plant height and technical stem length as well as this cross included high x low general combiners parents for GCA effects for straw yield. For seed yield, one cross ($P_1$x$P_3$) showed significant positive SCA values for both seed yield and no. of capsules per plant as well as this cross involved high x high general combiner for these two traits in addition one cross ($P_1$x$P_2$) had high x high general combiner parents for 1000-seed weight. Therefore, these crosses ($P_2$x$P_3$, $P_1$x$P_5$ and $P_1$x$P_3$) may prove useful for simultaneous improvement of the above-mentioned traits.

Straw yield per plant was significantly positively correlated with both of plant height and technical stem length. Also, plant height exhibited high positive correlation with technical stem length. On the other hand, seed yield per plant exhibited highly significant and positive correlation with each of no. of capsules, 1000-seed weight and no. of basal branches. Therefore, the possibility of using the two traits (plant height and technical stem length) as selection indices for improving straw yield per plant as well as, using 1000-seed weight and capsules number per plant as selection indices for improving seed yield per plant.

Keywords: Combining ability, Gene action, Diallel cross, Correlation, Flax.
INTRODUCTION

Flax (Linum usitatissimum L.) has always had industrial uses; recently its uses have widened to include a range of new possibilities such as cigarette papers, car door panels and compressed boards. But more and more, flax is carving a niche as a health food. Alpha-linolenic acid (an omega-3 fatty acid found in seed flax) is essential in the human diet. It can reduce heart disease and lower cholesterol. Flax is cultivated in Egypt for two purposes, seeds and fibers.

The diallel cross technique proposed by Griffing (1956) has been widely used for evaluation of general combining ability which is due to additive gene action and specific combining ability which is due to non-additive gene effects. The difficulty in producing enough F₁ hybrid seeds in some self pollinated crops have limited the use of diallel analysis, and in such cases, F₂ diallels may be more appropriate (Shehata and Comstock, 1971 and Patil and Chopde,1981). However F₂ diallels estimate dominance with only half the efficiency of F₁ diallels (Allard 1960).

The additive genetic variance had more important role in the inheritance of straw yield, plant height, technical stem length, seed index as reported by Foster et al (1998), Abo-Kaied et al (2007), Abo-Kaied and El-Refaie,Amany (2008) and El-Kady, Eman and Abo-Kaied (2010). On the other hand, non-additive variance had an important role in the inheritance of no. of basal branches per plant, seed yield per plant and no. of capsules per plant as reported by Shehata and Comstock (1971), Roa and Singh (1987), Mishra and Rai (1996), El-Kady and Abo-Kaied (2010) and Abdel-Moneam (2014).

This study aimed to estimate the combining ability and the type of gene action for yield and yield components traits for six flax genotype and their 15 crosses in F₂ generations with an ultimate goal of selecting suitable parents and the superior crosses which can be used in breeding program.

MATERIALS AND METHODS

The materials used for the present study consisted of (6 parents and 15 F₂).s. These six parents represent a wide genetic variability for yield and yield component traits of flax. Genotype characteristics of these parents used according to their pedigree, classification and origin are presented in Table 1.

<table>
<thead>
<tr>
<th>Genotypes</th>
<th>Pedigree</th>
<th>Type</th>
<th>Origin</th>
</tr>
</thead>
<tbody>
<tr>
<td>P₁=Giza 8</td>
<td>Giza 5 x Santa Catalina 6</td>
<td>oil</td>
<td>Local variety</td>
</tr>
<tr>
<td>P₂=Sakha</td>
<td>I. 2348 x Hera</td>
<td>dual</td>
<td>Local variety</td>
</tr>
<tr>
<td>P₃=S.413/1/2/3</td>
<td>S.5281/1 x S.40/9</td>
<td>dual</td>
<td>Local strain</td>
</tr>
<tr>
<td>P₄=S.420/12</td>
<td>S.162/12 x S.2467/1 (Hira)</td>
<td>dual</td>
<td>Local strain</td>
</tr>
<tr>
<td>P₅=Cizar</td>
<td>Introduction from Holland</td>
<td>fiber</td>
<td>Holland</td>
</tr>
<tr>
<td>P₆=Bombay</td>
<td>B12 x C.I. 42</td>
<td>fiber</td>
<td>USA</td>
</tr>
</tbody>
</table>
In 2012/2013 season, the six parents were crossed in a diallel mating design excluding reciprocals to obtain 15 $F_1$ crosses at Giza Res. Sta. of Agric. Res. Center. The $F_1$ seed were sown in 2013/2014 for obtaining $F_1$ hybrid plants and consequently to obtain $F_2$ seeds by selfing. In 2014/2015 season, the six parents and 15 $F_2$ were planted in a randomized complete block design with three replications at Giza Res. Station Farm. Seeds of each parent and $F_2$ was sown in two rows. The rows were 2 m long and 20 cm apart. Single seeds were hand drilled in 5 cm spacing within rows. Ten random guarded plants from each row (twenty plants for each parent and each $F_2$) used for measuring data. The following traits were recorded: (1) straw weight per plant (g), (2) plant height (cm), (3) technical stem length (cm), (4) no. of basal branches, (5) seed yield per plant (g), (6) 1000-seed weight (g), (7) no. of capsules per plant, and (8) no. of seeds per capsule.

**Statistical Analysis**

Plot means were used for statistical analysis. Combining abilities, general (GCA) and specific (SCA) were calculated according to Griffing's method 2, model 1 (fixed effects). Phenotypic correlation coefficients were calculated according to the formula suggested by Al-Jibouri et al., (1958).

**RESULTS AND DISCUSSION**

**Straw yield and its components :**

**Analysis of variances:**

Analysis of variance exhibited that mean squares due to entries (parents and $F_2$s) are highly significant for straw yield/plant and its components (Table 2). This indicates that those parental genotypes as well as the $F_2$s crosses showed a reasonable degree of variability for these traits. Also, mean squares due to parents and crosses were highly significant for all characters with exception for no. of basal branches/plant was significant only.

General (GCA) and specific (SCA) combining ability variances for these traits were highly significant, indicating the presence of both additive and non-additive type of genetic variance. The ratio of general to specific (GCA/SCA) combining ability variances for each of straw yield/plant, plant height and technical stem length showed that the additive effects were more important than non-additive effects. While, both additive and non-additive effects were the same almost effect in inheritance of no. of basal branches/plant. Although SCA mean squares were significant for plant height and technical stem length, the magnitude of GCA mean squares were greater than SCA mean squares for these two important components of straw yield (plant height and technical stem length). Therefore, the magnitude of additive genetic effects must be of considerable value for both characters. Consequently, effective selection should be possible within these $F_2$ and subsequent populations for these two traits which indicate to high yielding ability for straw yield. Similar results were reported by Gaafar et al (1992) Foster et al (1998), Zahana, Afaf (2006), Abo-Kaied et al (2007), Abo-Kaied and El-Refaie, Amany (2008), El-Kady, Eman and Abo-Kaied (2010) and Abdel-Moneam (2014).
The estimates of GCA effects are presented in Table 3. $P_3$ (S.413/1/2/3) and $P_4$ (S.420/12) showed high general combining ability for each of straw yield/plant, plant height and technical stem length, suggesting the importance of these two parents ($P_3$ and $P_4$) for increasing straw yield, plant height and technical length in flax breeding programs. Also, $P_1$ (Giza 8) and $P_6$ (Bombay) showed significant and positive GCA effects for no. of basal branches per plant. The simple correlation between GCA values (Table 3) and parental means (Table 5) for straw yield and its components were significantly positive. Similar findings were reported by Abo-Kaied and El-Refaie, Amany (2008) in flax. These results indicated that the parents showing higher mean performance proved to be the highest general combiners for these traits. Therefore, high mean performance of the parents could be transferred to hybrids in such cases. For mean performances, $P_3$ gave high mean value for each of straw yield/plant, plant height and technical stem length (Table 5).

SCA effects:
Specific combining ability (SCA) effects for 15 $F_2$’s crosses of straw yield per plant and its components are present in Table (4). Out of the 15 $F_2$ crosses, six crosses ($P_1 $$P_3$, $P_1 $$P_4$, $P_2 $$P_3$, $P_2 $$P_6$, $P_3 $$P_5$ and $P_4 $$P_3$) for straw yield/plant, five crosses ($P_2 $$P_4$, $P_2 $$P_5$, $P_2 $$P_6$, $P_3 $$P_5$ and $P_3 $$P_6$) for plant height, four crosses ($P_2 $$P_4$, $P_1 $$P_6$, $P_2 $$P_3$ and $P_3 $$P_3$) for technical stem length and one cross ($P_1 $$P_6$) for no. of basal branches/plant were exhibited significant and positive SCA effect. In general one cross ($P_3 $$P_5$) exhibited significant and positive SCA effect for each of straw yield, plant height and technical stem length as well as this cross ($P_3 $$P_5$) included high x low general combiners parents for GCA effects for straw yield, however, one cross ($P_1 $$P_6$) included two high general combiner parents for GCA effects for no. of basal branches. For mean performances, two crosses ($P_1 $$P_3$ and $P_3 $$P_4$) showed high mean performances for all straw characters traits with except no. of basal branches/plant (Table 5).

From the breeding point of view as suggested by Bhatada and Bahale (1983) for crosses exhibiting significant and positive SCA effects which resulted from high x high general combiners, the breeding procedure which utilize both additive and non-additive genetic variance would be more useful for improvement of straw yield. The available additive genetic variance should be exploited by adopting mass selection in early generations and some form of inter-se mating may be followed among elite selections in later generations, which may help in fixing non-additive effects.
2-3
Table 4. Estimation of specific combining ability ($s_{ij}$) effects for straw seed yields and their components in 15 flax crosses.

<table>
<thead>
<tr>
<th>Crosses</th>
<th>Straw yield (g)</th>
<th>Plant height (cm)</th>
<th>Technical length (cm)</th>
<th>No. of basal branches</th>
<th>Seed yield/plant (g)</th>
<th>No. of capsules/plant</th>
<th>1000-seed weight</th>
<th>No. of seeds/capsule</th>
</tr>
</thead>
<tbody>
<tr>
<td>P×P₁</td>
<td>0.188ns</td>
<td>2.199ns</td>
<td>-0.847ns</td>
<td>0.135ns</td>
<td>0.140ns</td>
<td>1.100ns</td>
<td>0.261**</td>
<td>-0.008ns</td>
</tr>
<tr>
<td>P×P₂</td>
<td>0.292**</td>
<td>1.955ns</td>
<td>-6.051**</td>
<td>0.199ns</td>
<td>0.201*</td>
<td>3.315*</td>
<td>0.179ns</td>
<td>-0.312ns</td>
</tr>
<tr>
<td>P×P₃</td>
<td>0.369**</td>
<td>2.194ns</td>
<td>2.237*</td>
<td>0.161ns</td>
<td>0.241**</td>
<td>2.069ns</td>
<td>0.192ns</td>
<td>0.128ns</td>
</tr>
<tr>
<td>P×P₄</td>
<td>0.025ns</td>
<td>1.253ns</td>
<td>0.932ns</td>
<td>0.180ns</td>
<td>0.213**</td>
<td>2.966**</td>
<td>0.094ns</td>
<td>-0.047ns</td>
</tr>
<tr>
<td>P×P₅</td>
<td>0.096ns</td>
<td>1.801ns</td>
<td>3.004**</td>
<td>0.193*</td>
<td>0.232**</td>
<td>0.247ns</td>
<td>0.746**</td>
<td>-0.048ns</td>
</tr>
<tr>
<td>P×P₆</td>
<td>0.438**</td>
<td>-7.293**</td>
<td>4.362**</td>
<td>0.118ns</td>
<td>0.134ns</td>
<td>6.878**</td>
<td>-0.032ns</td>
<td>-1.051**</td>
</tr>
<tr>
<td>P×P₇</td>
<td>-0.022ns</td>
<td>3.342**</td>
<td>1.397ns</td>
<td>0.120ns</td>
<td>0.124ns</td>
<td>-0.343ns</td>
<td>0.359**</td>
<td>0.231ns</td>
</tr>
<tr>
<td>P×P₈</td>
<td>-0.112ns</td>
<td>2.401*</td>
<td>-1.441ns</td>
<td>0.139ns</td>
<td>0.188*</td>
<td>0.564ns</td>
<td>0.415**</td>
<td>0.057ns</td>
</tr>
<tr>
<td>P×P₉</td>
<td>0.226*</td>
<td>2.685*</td>
<td>1.057ns</td>
<td>0.152ns</td>
<td>0.200*</td>
<td>2.208ns</td>
<td>0.083ns</td>
<td>0.056ns</td>
</tr>
<tr>
<td>P×P₁₀</td>
<td>-0.145ns</td>
<td>1.537ns</td>
<td>-5.911**</td>
<td>0.144ns</td>
<td>0.089ns</td>
<td>6.930**</td>
<td>-0.507**</td>
<td>-0.847**</td>
</tr>
<tr>
<td>P×P₁₁</td>
<td>0.325**</td>
<td>3.940**</td>
<td>3.854**</td>
<td>0.163ns</td>
<td>0.247**</td>
<td>-0.072ns</td>
<td>0.392**</td>
<td>0.287ns</td>
</tr>
<tr>
<td>P×P₁₂</td>
<td>-0.224*</td>
<td>2.394*</td>
<td>-8.674**</td>
<td>0.177ns</td>
<td>0.196*</td>
<td>0.196ns</td>
<td>0.284*</td>
<td>0.286ns</td>
</tr>
<tr>
<td>P×P₁₃</td>
<td>0.268**</td>
<td>0.992ns</td>
<td>0.000ns</td>
<td>0.165ns</td>
<td>0.162**</td>
<td>1.252ns</td>
<td>0.296**</td>
<td>0.000ns</td>
</tr>
<tr>
<td>P×P₁₄</td>
<td>-0.018ns</td>
<td>1.224ns</td>
<td>1.022ns</td>
<td>0.178ns</td>
<td>0.157**</td>
<td>1.847ns</td>
<td>0.187ns</td>
<td>0.000ns</td>
</tr>
<tr>
<td>P×P₁₅</td>
<td>0.174ns</td>
<td>1.428ns</td>
<td>1.907ns</td>
<td>0.099ns</td>
<td>-0.31**</td>
<td>-1.229ns</td>
<td>-0.417**</td>
<td>-0.093ns</td>
</tr>
<tr>
<td>LSD</td>
<td>0.280</td>
<td>1.343</td>
<td>0.284</td>
<td></td>
<td>0.229</td>
<td>4.304</td>
<td>0.289</td>
<td>0.514</td>
</tr>
<tr>
<td>5%</td>
<td>0.364</td>
<td>0.215</td>
<td>0.135</td>
<td>0.609*</td>
<td>0.620*</td>
<td>0.407</td>
<td>0.370</td>
<td>0.654**</td>
</tr>
</tbody>
</table>

# = Number refer to parent codes, Table 3.
*,** Significant at 0.05 and 0.01 levels of probability, respectively.
r : Simple correlation coefficients between SCA values and means of crosses.

Seed yield and its components:
Analysis of variances:
Analysis of variance showed that mean squares due to entries (6 parents and 15 $F₂$'s) were highly significant for seed yield/plant and its components (Table 2). Also, parents and crosses mean squares were significant for all characters. These results indicated that the parental genotypes as well as $F₂$'s crosses possess a reasonable degree of variability for these characters. Also, general (GCA) and specific (SCA) combining ability variances for these traits were highly significant, indicating the presence of both additive and non-additive type of genetic variance. Also, analysis of variances due to combining ability showed highly significant mean squares for both general and specific combining ability, revealing the important role of both additive and non-additive effects in the expression of seed yield and its components. The ratio of GCA/SCA for seed yield and its components, revealed that the inheritance of these traits were mainly controlled by additive genes effects. Abo-Kaied et al (2007), Abo-Kaied and El-Refaie, Amany (2008), El-Kady, Eman and Abo-Kaied (2010) and Abdel-Moneam (2014) reported similar results in combining ability for these traits in flax.
Table 5. Mean performances of 21 flax genotypes (6 parents and 15 F$_2$'s crosses) for straw, seed yield and their components.

<table>
<thead>
<tr>
<th>Genotypes</th>
<th>Straw yield and its components</th>
<th>Seed yield and its components</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Straw yield / plant (g)</td>
<td>Plant height (cm)</td>
</tr>
<tr>
<td>Parents</td>
<td></td>
<td></td>
</tr>
<tr>
<td>P1 #</td>
<td>1.83</td>
<td>94.90</td>
</tr>
<tr>
<td>P2</td>
<td>1.38</td>
<td>83.84</td>
</tr>
<tr>
<td>P3</td>
<td>3.43</td>
<td>124.00</td>
</tr>
<tr>
<td>P4</td>
<td>2.89</td>
<td>111.10</td>
</tr>
<tr>
<td>P5</td>
<td>1.61</td>
<td>86.95</td>
</tr>
<tr>
<td>P6</td>
<td>2.18</td>
<td>99.79</td>
</tr>
<tr>
<td>Crosses</td>
<td></td>
<td></td>
</tr>
<tr>
<td>P1 x P2</td>
<td>2.21</td>
<td>94.75</td>
</tr>
<tr>
<td>P1 x P3</td>
<td>3.34</td>
<td>114.39</td>
</tr>
<tr>
<td>P1 x P4</td>
<td>3.09</td>
<td>109.87</td>
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<tr>
<td>P1 x P5</td>
<td>2.16</td>
<td>96.99</td>
</tr>
<tr>
<td>P1 x P6</td>
<td>2.41</td>
<td>103.84</td>
</tr>
<tr>
<td>P2 x P3</td>
<td>3.20</td>
<td>98.09</td>
</tr>
<tr>
<td>P2 x P4</td>
<td>2.41</td>
<td>103.97</td>
</tr>
<tr>
<td>P2 x P5</td>
<td>1.73</td>
<td>91.09</td>
</tr>
<tr>
<td>P2 x P6</td>
<td>2.25</td>
<td>97.67</td>
</tr>
<tr>
<td>P3 x P4</td>
<td>3.30</td>
<td>122.04</td>
</tr>
<tr>
<td>P3 x P5</td>
<td>3.19</td>
<td>112.51</td>
</tr>
<tr>
<td>P3 x P6</td>
<td>2.82</td>
<td>117.26</td>
</tr>
<tr>
<td>P4 x P5</td>
<td>2.80</td>
<td>104.80</td>
</tr>
<tr>
<td>P4 x P6</td>
<td>2.70</td>
<td>111.33</td>
</tr>
<tr>
<td>P5 x P6</td>
<td>2.42</td>
<td>99.35</td>
</tr>
<tr>
<td>G. Means</td>
<td>1.91</td>
<td>75.14</td>
</tr>
<tr>
<td>LSD 0.05</td>
<td>0.38</td>
<td>4.45</td>
</tr>
</tbody>
</table>

# = Number refer to parent codes, Table 3.

GCA effects:

Estimates of GCA effects for each parent are presented in Table 3. The data indicated that P$_1$ (Giza 8) and P$_5$ (Cizar) showed significant and positive GCA effects for seed yield and no. of capsules/plant also P$_1$, P$_2$ (Sakha 2) and P$_3$ (S.413/1/2/3) for 1000-seed weight as well as P$_5$ and P$_6$ (bombay) for no. of seeds/capsule. In general, P$_1$ proved to be good combiner for seed yield and its two important components (no. of capsules/plant and 1000-seed weight). Also, P$_5$ exhibited high general combiner for all seed yield characters except 1000-seed weight. Therefore, using such parents in varietal improvement programs may result in isolating desirable segregates for these traits owing to the breeders desire. The correlation coefficient between GCA values (Table 3) and parental means for seed yield and its characters (Table 5) were significantly positive. These results indicated that the superiority of a parent in cross combinations could be directly predicted its per se performance.
SCA effects:
Specific combining ability effects calculated for each cross are presented in Table (4). The data showed that positive significant SCA values have been registered in ten crosses for seed yield/plant, four crosses for no. of capsules per plant and seven crosses for 1000-seed weight. In general, one cross \( (P_1 \times P_3) \) showed significant positive SCA values for both seed yield and no. of capsules/plant as well as this cross involved high x high general combiner for these two traits in addition one cross \( (P_1 \times P_2) \) had high x high general combiner parents for 1000-seed weight. Therefore, these crosses \( (P_1 \times P_5) \) for seed yield and no. of capsules as well as \( P_1 \times P_2 \) for 1000-seed weight) are likely to throw good segregates for these traits if the allelic genetic systems are present in good combination and epistatic effects present in the crosses act in the same direction to maximize the desirable characteristics. Therefore, these crosses \( (P_1 \times P_5 \) and \( P_1 \times P_2) \) may prove useful for simultaneous improvement of the above-mentioned traits. As well as, the cross \( (P_1 \times P_5) \) gave high mean performance for both seed yield/plant and no. of capsules/plant (Table 5). The correlation between cross means (Table 5) and their SCA values (Table 4) was significant and positive for seed yield and no. of seeds/capsule indicating that high performing crosses were high specific combinations. Therefore, the choice of promising cross combinations would be based on SCA effects or mean performance of a cross.

Correlation studies
Phenotypic correlation coefficients among eight traits in flax are shown in Table 6. straw yield/plant was significantly positively correlation with both of plant height and technical stem length. Also, plant height exhibited high positive correlated with technical stem length. On the other hand seed yield/plant exhibited highly significant and positive correlation with each of no. of capsules and 1000-seed weight. The no. of basal branches showed highly r values with each of seed yield and no. of capsules/plant. Therefore, the possibility of using the two traits (plant height and technical stem length) as selection indices for improving straw yield/plant as well as, using 1000-seed weight, capsules number per plant and no. of basal branches as selection indices for improving seed yield/plant. While, no. of seeds/capsule exhibited highly significant and negative correlation with each of seed yield/plant, no. of capsules/plant and 1000-seed weight. These results are in harmony with that reported by Momtaz et al. (1977), Mourad et al. (1987) and Abo-Kaied and El-Refaie, Amany (2008).
Table 6. Simple correlation coefficients among eight traits in 21 flax genotypes.

<table>
<thead>
<tr>
<th>Characters</th>
<th>Straw yield/plant (g)</th>
<th>Plant Height (cm)</th>
<th>Technical length (cm)</th>
<th>No. of basal branches</th>
<th>Seed yield/plant (g)</th>
<th>No. of capsules/plant</th>
<th>1000-seed weight</th>
</tr>
</thead>
<tbody>
<tr>
<td>1-Straw yield/plant (g)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>2-Plant height (cm)</td>
<td>0.89**</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>3-Technical stem length (cm)</td>
<td>0.83**</td>
<td>0.80**</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>4-No. of seeds/capsule</td>
<td>0.40</td>
<td>0.37</td>
<td>-0.03</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>5-Seed yield/plant (g)</td>
<td>0.12</td>
<td>-0.02</td>
<td>-0.25</td>
<td>0.58**</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>6-No. of capsules/plant</td>
<td>-0.01</td>
<td>-0.15</td>
<td>-0.42</td>
<td>0.61**</td>
<td>0.85**</td>
<td>0.28</td>
<td></td>
</tr>
<tr>
<td>7-1000 seed-weight (g)</td>
<td>0.22</td>
<td>0.07</td>
<td>0.16</td>
<td>0.03</td>
<td>0.61**</td>
<td>0.26</td>
<td></td>
</tr>
<tr>
<td>8-No. of seeds/capsule</td>
<td>-0.04</td>
<td>0.16</td>
<td>0.12</td>
<td>-0.14</td>
<td>-0.56**</td>
<td>-0.64**</td>
<td>-0.66**</td>
</tr>
</tbody>
</table>

*,** Significant at 0.05 and 0.01 levels of probability, respectively.

REFERENCES


تحليل الهجن الدائرية للمحصول ومكوناته في ستة تراكيب وراثية من الكتان

ماسية سعيد عبدالصادق

معهد بحوث المحاصيل الحقلية - مركز البحوث الزراعية - الجيزة

تُرصد هذه الدراسة تأثير قدرة على الانتفلاج الفعل الفعلي في التكيف النباتي لمحصول القش والبذور ومكوناته في التракيبات الوراثية. استخدمت النباتات من نوع لينوم (Linum usitatissimum L.) في اثنان من 15 نموذج في الجيل الثاني، تأثر بالنسب الوراثية للفعل الوراثي. باستخدام نموذج الهجنة، تم تقسيم نباتات الدراسة إلى جداول زراعية مدفوعة في محطة البحوث الزراعية في سيناء خلال موسم 2019/2020.

استنتجت النتائج أن التأثيرات الوراثية المؤثرة أكبر على مستواة القش النباتي والطول الكلي والطول الفعال، وعدد الكبسولات للنبات. تأثرت حيزاًً أصغر عدد الورقة الأساسية والمتوسطة بالنظام. تم استخدام نتائج الدراسة للتنبؤ بمستوى القش النباتي والطول الكلي والطول الفعال، وعدد الكبسولات للنبات. يمكن استمرار الاختبار في البرامج التربوية للتحسين في مستويات القش النباتي والطول الفعال، وعدد الكبسولات للنبات.
Table 2. Mean squares for 21 genotypes (parents and crosses), general (GCA) and specific (SCA) combining ability for straw and seed yields and their components in flax

<table>
<thead>
<tr>
<th>Character</th>
<th>S.O.V.</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Reps. 2#</td>
</tr>
<tr>
<td>Straw yield/plant (g)</td>
<td>0.34**</td>
</tr>
<tr>
<td>Plant height (cm)</td>
<td>18.62**</td>
</tr>
<tr>
<td>Technical stem length (cm)</td>
<td>7.64</td>
</tr>
<tr>
<td>No.of basal branches</td>
<td>0.17**</td>
</tr>
<tr>
<td>seed yield/plant (g)</td>
<td>0.13**</td>
</tr>
<tr>
<td>No.of capsules/plant</td>
<td>18.27</td>
</tr>
<tr>
<td>1000seed-weight (g)</td>
<td>0.16*</td>
</tr>
<tr>
<td>No.of seeds /capsule</td>
<td>0.22</td>
</tr>
</tbody>
</table>

Note: * The degrees of freedom

Table 3. Estimation of general combining ability effects ($\bar{g}_i$) for straw and seed yields and their components in 6 flax genotypes.

<table>
<thead>
<tr>
<th>Parents</th>
<th>Straw yield / plant (g)</th>
<th>Plant Height (cm)</th>
<th>Technical length (cm)</th>
<th>No. of basal branches</th>
<th>Seed yield / plant (g)</th>
<th>No. of capsules /plant</th>
<th>1000-seed weight</th>
<th>No. of seeds /capsule</th>
</tr>
</thead>
<tbody>
<tr>
<td>P1=Giza 8</td>
<td>-0.110**</td>
<td>-2.971**</td>
<td>-3.819**</td>
<td>0.113**</td>
<td>0.346**</td>
<td>6.194**</td>
<td>0.671**</td>
<td>-0.748**</td>
</tr>
<tr>
<td>P2=Sakha 2</td>
<td>-0.399**</td>
<td>-9.118**</td>
<td>-3.581**</td>
<td>-0.175**</td>
<td>-0.086**</td>
<td>-2.734**</td>
<td>0.591**</td>
<td>-0.236**</td>
</tr>
<tr>
<td></td>
<td>P3-S.413/1/2/3</td>
<td>P4-S.420/12</td>
<td>P5-Cizar</td>
<td>P6-Bombay</td>
<td>LSD (gi - gi)</td>
<td>r</td>
<td></td>
<td></td>
</tr>
<tr>
<td>----------------</td>
<td>----------------</td>
<td>-------------</td>
<td>----------</td>
<td>-----------</td>
<td>--------------</td>
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<td></td>
<td></td>
</tr>
<tr>
<td><strong>0.05</strong></td>
<td>0.621**</td>
<td>0.292**</td>
<td>-0.261**</td>
<td>-0.113**</td>
<td>0.112</td>
<td>0.149</td>
<td></td>
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<tr>
<td></td>
<td>10.762**</td>
<td>6.001**</td>
<td>-5.938**</td>
<td>0.364ns</td>
<td>1.295</td>
<td>1.719</td>
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<tr>
<td></td>
<td>8.967**</td>
<td>5.559**</td>
<td>-5.852**</td>
<td>-1.475**</td>
<td>1.198</td>
<td>1.599</td>
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<tr>
<td></td>
<td>-0.003ns</td>
<td>0.006ns</td>
<td>-0.017ns</td>
<td>0.076**</td>
<td>0.107</td>
<td>0.143</td>
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<tr>
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<td>-0.016 ns</td>
<td>-0.082**</td>
<td>0.158**</td>
<td>0.320**</td>
<td>1.627</td>
<td>2.176</td>
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<tr>
<td></td>
<td>-2.108**</td>
<td>-1.596**</td>
<td>2.186**</td>
<td>-1.941**</td>
<td>0.109</td>
<td>0.146</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>0.500**</td>
<td>-0.001ns</td>
<td>-0.374**</td>
<td>-1.388**</td>
<td>0.194</td>
<td>0.260</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>0.055ns</td>
<td>0.010ns</td>
<td>0.347**</td>
<td>0.573**</td>
<td>0.194</td>
<td>0.260</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

*,** Significant at 0.05 and 0.01 levels of probability, respectively.

r : Simple correlation coefficients between GCA values and parental means.