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Screening for sources of tolerance to drought in sesame induced mutants: Assessment of indirect selection criteria for seed yield

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ABSTRACT

In African sub-Saharan countries, sesame is mostly grown under rain fed conditions where it is subjected to drought stress. The objective of this study was to identify the drought tolerance sources in induced-mutants of sesame. Sixteen induced mutants and their three respective parental sources were evaluated in separate experiments under drought stress and normal conditions in the field over two years (2010 and 2011). Seven drought tolerance indices including stress susceptibility index (SSI), stress tolerance index (STI), mean productivity (MP), geometric mean productivity (GMP), tolerance (TOL), yield index (YI) and yield stability index (YSI) were calculated based on yield under drought (Y_s) and yield in optimal conditions (Y_p). Factor analysis (FA) evidenced that first and second factors accounted for 98.7 and 98.5 % of the variation in the first and second year, respectively. Biplot and FA evidenced that genotypes LC 164, LC 162, BC 167, EF 147 and MT 169 had the highest grain yield under both DS and NS environments in 2010, whereas in 2011 the best performers in both environments were HC 108, 32-15, HB 168 and 38-1-7. FA and the mean rank method discriminated genotypes LC 164, LC 162 and BC 167 as the most drought-tolerant in 2010 whereas in 2011 the combined methods identified 32-15 as the highest drought-tolerant genotype. Plant height, the number of capsules per plant, and the length of the capsules should be considered in selection for obtaining high-yielding sesame cultivars in drought-stressed environments.

Key words: *Sesamum indicum L., drought stress, yield, criteria.*

INTRODUCTION

Sesame (*Sesamum indicum* L., Pedaliaceae) is also called orphan crop. Nowadays, however world demand for its seeds is interestingly increasing owing to its good quality oil (50 %), protein (25 %) and for content of antioxidants[3,11,33]. Beside these nutritional benefits, sesame cropping has many agricultural advantages: it grows well in tropical to temperate climates, it can grow on stored soil moisture without the need for rainfall or irrigation, and be grown in pure stands with low input, or else in mixed stands with diverse crops[4].

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Despite the many advantages of sesame seeds, less attention is dedicated to the crop by research centers so that genetic and breeding improvement efforts in sesame have been limited making the results of such efforts slow to emerge. Ashri[4] stated that the main reason for this limited success is that sesame is a crop mainly produced in developing countries and usually by smallholders. The total world production was about 4 756 751 tones from a planted area of 9 million hectares in 2013 and the average yields ranged from 385 kg ha^{-1} in 2000 to 506 kg ha^{-1} in 2013[12]. Despite the high yield potential of sesame, actual yields are quite low due to a combination of biotic and abiotic stresses.

The main sesame grower countries are India, Myanmar and Sudan, most growing areas are classed as arid or semi-arid[12] and in these regions, sesame is subjected to terminal and intermittent droughts. In these drought prone environments, breeders' primary interest is in grain yield which may be considered to be affected by three components including yield potential, appropriate phenology and drought tolerance[24]. To achieve high and durable yield in such environments drought-tolerant genotypes are needed. Unfortunately, the development of improved sesame cultivars for drought tolerance is hampered by the lack of efficient selection criteria. Two classical methods are usually followed to select for drought tolerance in crops: (i) utilization of grain yield as direct selection criteria, and (ii) indirect selection based on secondary traits[26] which are plant characteristics that are associated with yield, and they can provide additional information for breeders to use when they make selections[20].

Although some authors reported positive correlation between yield in optimal conditions and yield under drought[1,8,17,22], direct selection based on yield potential or mean yield under non-stress conditions may be misleading for the selection of drought-tolerant genotypes because drought tolerance is a complex quantitative trait, involving interactions of many metabolic pathways related to stress tolerance genes[1].

The identification of a standard evaluation assay has been the most pressing problem for the selection of drought-tolerant genotypes[18]. Different indices have been employed for selecting drought-tolerant genotypes. Rosielle and Hamblin[25] defined stress tolerance (TOL) as the difference between yields under optimal (Y_p) and stress conditions (Y_s) and the mean productivity (MP) as the average yield between Y_s and Y_p. Fischer and Maurer[15] proposed a stress susceptibility index (SSI) and Fernandez[14] introduced a stress tolerance index (STI) as a selection criterion to identify genotypes with high yield and stress tolerance potentials. The latter author stated that MP has an upward bias due to a relatively larger difference between Y_p and Y_s and proposed a novel index, the geometric mean productivity (GMP) which is less sensitive to large extreme values.

All these indices are based on grain yield though, it is useful to screen for secondary traits as well because grain yield under drought is a complex quantitative trait whose repeatability is low relative to yield in non-stress environments, reducing selection efficiency[16]. Also, high-yielding cultivars in well-watered conditions are not necessarily the top performers in drought-stressed conditions. Hence much effort has been focused on the genetic analysis of secondary traits. In a drought breeding program, secondary traits are valuable for many reasons: If observed before or at flowering, they can be used for selecting desirable crossing parents; if observed before maturity, they can be used for preliminary selection.

Bänziger and Lafitte[5] reported that secondary traits can help to improve the precision with which drought-tolerant genotypes are identified, compared to measuring only grain yield under drought. Therefore, the understanding of the relationship between yield and secondary traits is crucial for developing an adequate breeding program and this relationship is traditionally explained by means of correlation, regression and path coefficient analyses[29,30,38]. Path coefficient analysis[36] is helpful in partitioning the direct and indirect contribution of yield components to seed yield[29] and gives more realistic relationship between characters than the phenotypic correlation.

The objective of this study was to evaluate and select high yielding sesame genotypes and identify secondary traits to be used as selection criteria for seed yield under both drought and optimal conditions. Phenotypic correlations, path coefficients and factor analysis will be used for this assessment.

MATERIALS AND METHODS

Germplasm and location

The experiments were carried out with 19 sesame lines at the experimental station of the Centre National de Recherche Agronomique (CNRA), Bambey, Senegal (latitude 14° 42' North and longitude 16° 28' West) during the dry season of 2010 (04 November 2010 to January 25, 2011) and the wet cropping season of 2011 (July, 01-November, 30). These lines included 16 gamma-ray-induced mutants and their three respective parental sources: 32-15; 38-1-7 and Birkan. Mutants were induced in 2008 using 300 and 400 Gy gamma rays doses irradiating seeds of the three mother sources cited above. These mutants were confirmed in 2009 and 2010[7,10]. Cultivars 32-15 and 38-1-7 belonged to the sesame germplasm collection of the Institut Sénégalais de Recherches Agricoles (ISRA) and are widely grown by senegalese sesame growers for their rusticity and their marketable seeds' value (white seeds). Cultivar Birkan is a non-photosensitive mutant-cultivar introduced from the Faculty of Agriculture, Akdeniz University (Turkey).

Experimental design and set-up

In both 2010 and 2011, the layout was a factorial design consisted of adjacent non-stressed (NS) and drought-stressed (DS) blocks separated by a buffer of 10 m to prevent lateral movement of water from the NS to the DS plots. Within each block, plants were assigned to experimental plots using a randomized complete block design with three replications. Each plot consisted of four rows, spaced 0.6 m apart with 0.20 m between plants in the row. Row length was 4 m in both years. All trials were established in the field and plots were kept free from weeds, diseases and insects pests by a combination of preventive chemicals treatments (Decis, 1ml l^{-1}) and hand labour.

Prior to seeding, the soil was ploughed at a 15 cm depth. A composite N-P-K fertilizer (15-15-15) was applied at a rate of 80 kg ha^{-1} before sowing. In the dry season of 2010-2011 (04 November 2010 to January 25, 2011), plots were irrigated with 40 mm water one day prior to seeding using an oscillating ramp system. After emergence, both NS and DS blocks were irrigated with 20 mm of water twice a week until flowering time. Thereafter, DS plots did not receive water until harvest, whereas NS plants were kept well watered by receiving 20 mm of water twice a week until physiological maturity. Non-stressed plots received a total amount of 460 mm water, while DS plots had received a total of 180 mm before flowering.

Environmental data, including daily rainfall (mm), minimum and maximum temperatures (°C) and relative air humidity (%) were obtained from an automated weather station (Hobo H21-002) placed on the experimental site. The minimum air temperature and humidity were 13.4 °C and 8 %, respectively and the maximum were 39.1°C and 100 %, respectively.

In the 2011 wet cropping season, NS plots were sown in July as normal planting. To simulate a terminal drought, DS plots were planted with a delay of two months (September, 5). In all DS plots, 50 % flowering occurred between October, 9 and 15. After this period, no rainfall was recorded until crop maturity. Total rainfall was 584.2 mm in NS plots while DS plots received 159.9 mm before flowering. The minimum air temperature and humidity were 15.5 °C and 13 %, respectively and the maximum were 40.4°C and 100 %, respectively.

In both seasons, data were recorded for plant height (PH), height to the first capsule on the main stem (SLFC), number of branches per plant (NB), number of capsules per plant (NCP), number of seed per capsule (NSC), capsule length (LC), 1000-seed weight (SW) and seed yield. For seed yield measurement, 2.4 m^2 where harvested from the two central rows. Prior to harvesting, the first two plants at the borders of the row were discarded.

In addition, the drought intensity index (DII) defined by Fischer and Maurer[15] was determined for each season.

$DII = \frac{1 - X_{ds}}{X_{ns}}$, where X_{ds} and X_{ns} are the mean yields of all genotypes under drought-stressed and non-stressed conditions, respectively.

The stress susceptibility index (SSI) was calculated according to Fischer and Maurer[15] and the stress tolerance index (STI) was determined for each genotype following Fernandez[14]:

$SSI = \frac{\left(\frac{1-Y_s}{Y_p}\right)}{DII}$, where Y_s and Y_p are the mean yields of a given genotype in DS and NS environments, respectively.

$STI = \frac{Ys \times Yp}{X_p^2}$, where X_p is the mean yield of all genotypes under non-stressed conditions.

Geometric mean productivity (GMP) was calculated for seed yield according to Fernandez[14]:

$GMP = \sqrt{(Ys \times Yp)}$

Data were analyzed using MINITAB statistical program by one-way ANOVA and t test. Differences between mean values of treatments were evaluated using least significant difference (LSD) at a 0.05 significance level. Path coefficient and phenotypic correlation analyses were carried out to determine the relationship between the traits studied and their direct and indirect contribution to seed yield[9].

RESULTS

Variation in yield and yield components under non-stressed (NS) and drought-stressed (DS) environments

For both years, the results of ANOVA showed significant differences between genotypes in respect to yield and yield components under normal and drought stress conditions (Tables 1 & 2) except for plant height and number of branches in 2010 and for number of capsules per plant in 2011 in drought stress conditions.

Comparison of means grain yield per genotypes indicated that in well watered conditions (NS) HB 168 in 2010 and 38-1-7, HB 168, and 32-15 in 2011 had the highest grain yield while Birkan, MC 114 in 2010; MT 169 in 2011 had the lowest yield. In drought stress conditions (DS), LC 164 and LC 162 in 2010; HC 107 in 2011 had the highest yield while MC 114 had the lowest in both years (Tables 1 & 2). The range in yield under normal and drought stress conditions showed that there is a genotypic variability between genotypes for productivity.

In 2010, genotypes LC 164, LC 162, BC 167, EF 147 and MT 169 had the highest grain yield under both DS and NS environments while ICN 130, MC 112, 32-15, ICN 115, HC 107, Birkan, and MC 114 had the lowest grain yields under both DS and NS conditions (Table 1, Fig. 1a). HC 108, SHI 165 and EF 153 had the highest grain yields only in DS environment in contrast to VGR 156, 38-1-7, HSC 105 and HB 168 which were the best performers only in NS environment.

In 2011, HC 108, 32-15, HB 168 and 38-1-7 had the highest grain yield under both DS and NS conditions. In contrast MT 169, MC 112, EF 147, SHI 165, and MC 114 had the lowest grain yield in both DS and NS conditions (Table 2, Fig. 1b). HC 107, LC 162, LC 164, HSC 105, and ICN 115 had the highest grain yields in DS environments while Birkan, VGR 156, BC 167, ICN 130 and EF 153 revealed higher grain yields only in NS conditions (Table 2, Fig. 1b).

Stress tolerance indices and selection for drought resistance

To evaluate 19 sesame genotypes for drought tolerance, seven selection indices (SSI, STI, MP GMP, TOL, YI and YSI) were used. STI, MP, GMP and YI had significant positive correlation with both yield

under drought-stressed (Y_s) and yield under non stressed environments in both two years but there is no correlation between YI and Y_p in 2011 (Table 3 & 4). YI had the highest correlation ($r = 1^{**}$ in both years) with Y_s, whereas MP had the highest correlation with Y_p ($r = 0.87^{**}$ and $r = 0.94^{**}$, for 2010 and 2011, respectively).

The correlation between Y_p and Y_s was positively significant ($r = 0.49^{*}$) in 2010 but not significant at all in 2011. SSI and TOL were negatively correlated with Ys even though this correlation is not significant in the year 2011 (Table 4).

The estimates of drought tolerance attributes based on a single criterion are contradictory. In the 2010 trial and according to STI, MP and GMP genotypes LC 164, LC 162 and HB 168 were the most drought-tolerant genotypes whereas MC 114, Birkan and MC 112 were the most sensitive ones during the season 2010 (Table 5). Based on TOL scores HC 108, LC 164 and SHI 165 were the most desirable drought tolerant genotypes and HB 168, HSC 105 and 38-1-7 the most sensitive genotypes. According to SSI and YSI the desirable drought-tolerant genotypes were HC 108, BC 167 and SHI 165 (Table 5).

The same contradiction was highlighted in the 2011 cropping season when suitable drought-tolerant genotypes were selected based on a single drought tolerance index (Table 6).

The mean rank and standard deviation of ranks of all drought tolerance criteria were calculated and based on these two criteria the most desirable drought tolerant genotypes were identified. In consideration to all indices genotypes LC 164, LC 162 and BC 167 exhibited the best mean rank and low standard deviation of ranks (Table 5) under drought-stressed environment in 2010, hence they were considered as the suitable drought tolerant genotypes.

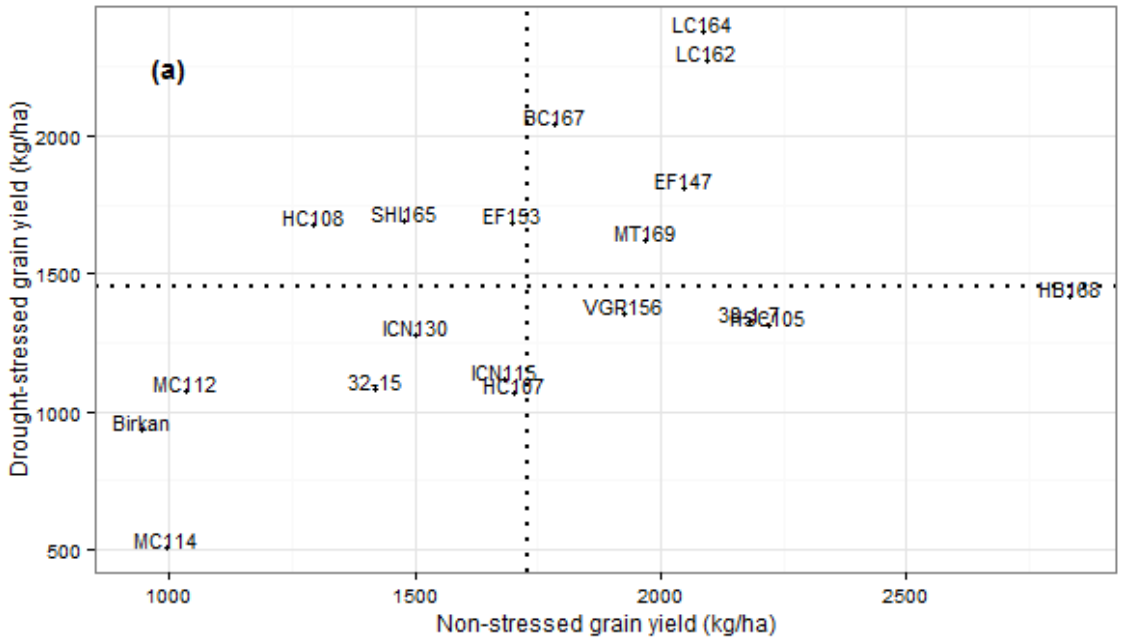
In 2011, LC 162, 32-15, HB 168 and HC 108 had the best mean rank and low standard deviation (Table 6) and were identified as the most drought tolerant genotypes.

Genotype LC 162 could therefore be identified as the best drought tolerant material, while MC 114 was the most sensitive for both years.

To use all indices simultaneously, factor analysis was also carried out. The two first factors explained 98.7 % and 98.5 % of the total variance in 2010 and 2011, respectively (Table 7). The relationship between the genotypes and all the drought tolerance indices is plotted in the same graph (Fig 2a & 2b). The first factor (FA1) was highly and positively correlated with Y_p, STI, MP and GMP in both years (Table). YI and Y_s were positively correlated by the first factor in 2010. Therefore, FA1 in both years was named as drought tolerance.

The second factor (FA2) was represented by SSI, TOL and YSI in 2010 and by Y_s, YI, STI, MP and GMP in 2011. SSI, TOL, YS, STI, MP, GMP and YI had negative coefficient with FA2. Thus the higher scores for FA1 and FA2 in 2010 were in accordance with higher drought tolerance while in 2011 it's higher score for FA1 and lower scores for FA2 which may be considered as higher drought tolerance. The sum of two first factors (FA1+FA2) are presented in Tables 5 and 6, respectively.

Coefficients of direct and indirect effects of path analysis in drought-stressed conditions are shown in Table 8. The number of capsules per plant had the highest positive and direct effect ($p = 0.519$) on seed yield in drought conditions. This trait was followed by plant height ($p = 0.332$), thousand seeds weight ($p = 0.276$), length of the capsule ($p = 0.233$) and number of seeds per capsule ($p = 0.176$). The stem length to the first capsule had negative direct effect (-0.283) on seed yield under drought-stressed conditions. The height of the Plant had the highest indirect effect (0.348) on seed yield via the number of capsules per plant. Similarly, the number of capsules per plant had a positive indirect effect (0.223) via plant height on seed yield followed by stem length to the first capsule (0.221) via plant height. Thousand seeds weight had negative indirect effect (-0.216) on seed yield via number of capsules per plant.



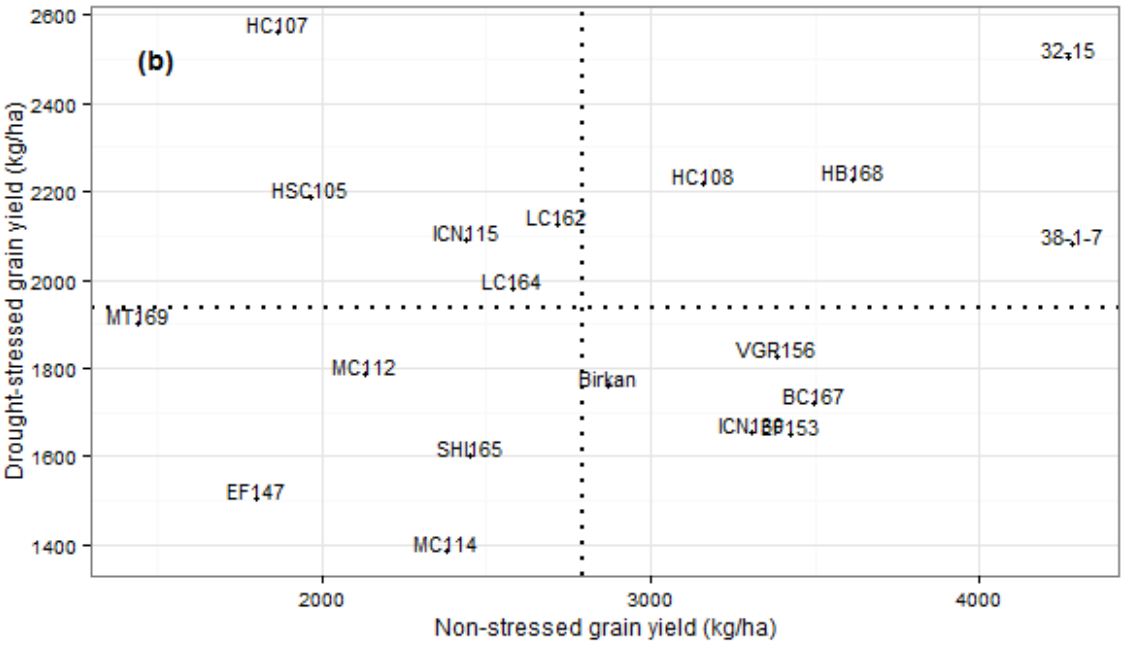


Fig. 1: Biplot for seed yield of 19 sesame genotypes in non-stressed and drought-stressed conditions in 2010 (a) and 2011 (b) Dotted lines represent overall yield mean in non-stressed (vertical) and drought-stressed (horizontal) conditions

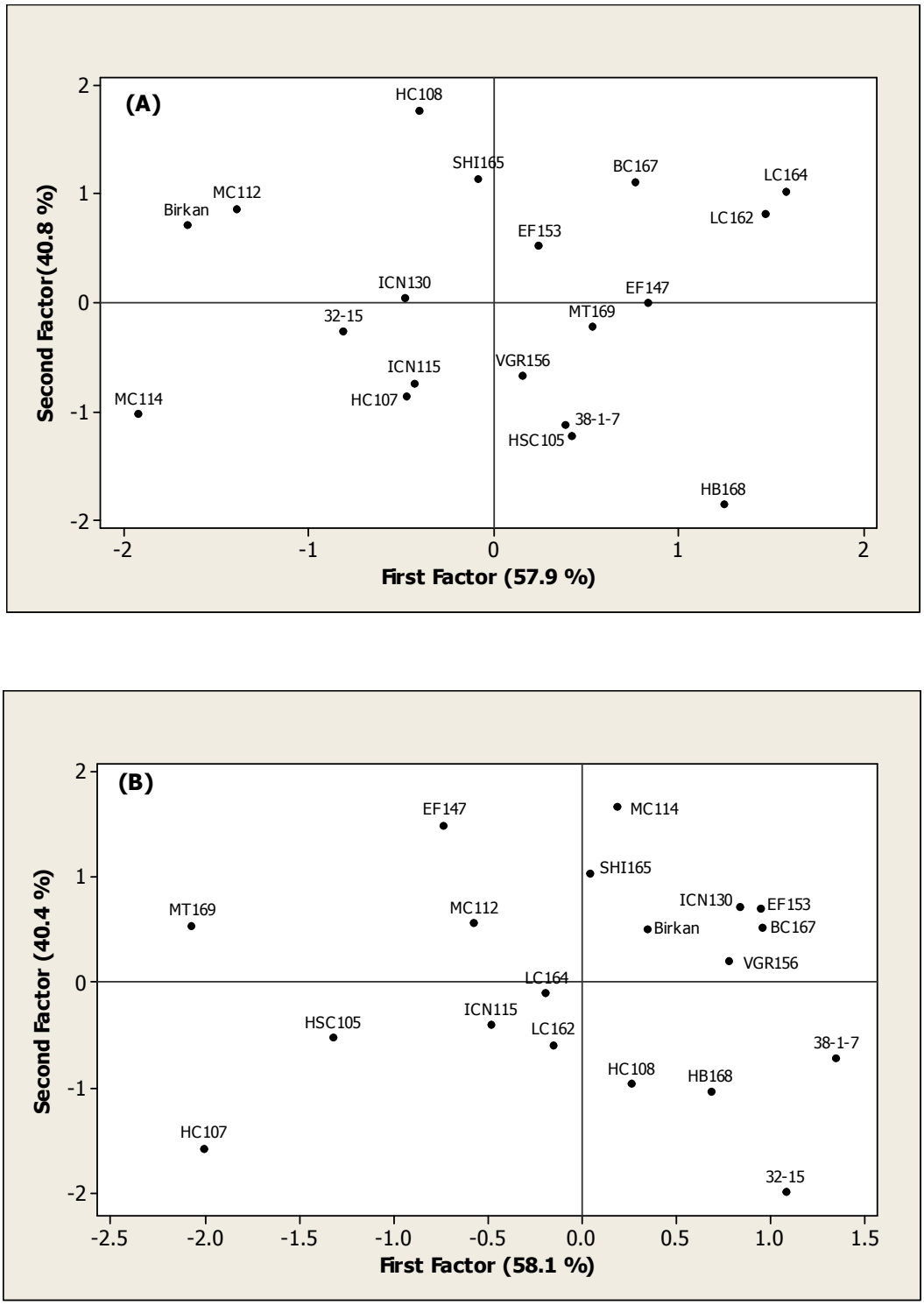


Fig. 2: Biplot based on first and second factor for 19 sesame genotypes during 2010 (A) and 2011 (B)

Table 1. Yield component and yield of sesame genotypes grown under well watered (WW) and water stress (WS) conditions during 2010

Genotype	PH(cm)		SLFC(cm)		NCP		NSC		NB		Lcap (cm)		1000-SW (g)		Yield(kg ha^{-1})	
	WW	WS	WW	WS	WW	WS	WW	WS	WW	WS	WW	WS	WW	WS	WW	WS
32-15	93.3	72.5	27.7	20.6	102.6	58.0	55.0	59.0	4.2	4.3	2.7	3.0	3.7	3.3	1418.9	1083.3
38-1-7	98.2	82.0	27.8	25.0	138.7	100.2	71.0	48.0	4.7	5.0	3.6	2.8	2.9	3.2	2182.1	1325.1
BC167	92.3	87.9	20.9	23.8	124.4	71.0	76.3	71.0	3.5	4.9	3.5	3.1	3.7	3.6	1782.7	2041.6
Birkan	83.5	74.5	17.7	17.6	80.0	52.6	61.7	56.0	3.4	3.0	3.2	2.7	3.7	4.2	943.5	935.7
EF147	93.3	90.8	19.9	18.4	90.7	99.1	71.3	77.3	3.6	4.7	3.5	3.9	3.8	3.7	2047.4	1809.3
EF153	102.9	90.9	24.9	25.6	102.4	103.9	76.7	68.3	3.9	6.4	3.0	3.0	3.0	3.2	1699.2	1686.1
HB168	109.6	82.2	26.0	23.0	152.1	85.8	60.3	52.0	4.6	6.1	3.2	2.9	3.6	3.5	2834.1	1419.6
HC107	89.1	72.3	25.0	20.8	89.9	52.2	64.7	75.0	3.9	3.9	2.9	3.4	3.5	3.4	1703.5	1070.7
HC108	101.2	94.1	24.2	23.8	120.7	116.8	69.7	37.0	4.7	6.8	2.8	2.7	3.2	3.3	1292.4	1678.2
HSC105	110.1	85.4	21.0	20.4	164.9	112.9	53.0	45.3	3.9	6.0	3.4	2.8	3.3	3.4	2218.8	1315.2
ICN115	88.9	76.2	18.0	16.1	108.9	76.0	64.3	57.3	3.7	4.2	3.5	3.2	3.6	3.6	1681.8	1116.0
ICN130	83.8	71.4	21.0	18.5	97.8	60.5	61.7	58.0	2.5	2.7	2.8	2.6	3.6	3.7	1500.5	1278.0
LC162	116.1	102.0	23.9	29.2	138.3	118.6	80.7	78.3	4.3	6.8	3.1	3.5	3.3	3.2	2093.4	2273.2
LC164	94.4	92.8	26.3	25.8	112.1	92.1	71.7	77.7	3.7	5.1	3.2	3.3	3.7	3.7	2086.1	2379.0
MC112	93.9	83.1	24.9	20.3	106.9	78.0	69.0	71.0	4.3	5.3	3.0	2.5	3.4	3.5	1033.7	1076.4
MC114	121.6	102.1	50.0	52.9	73.4	81.0	111.3	75.0	4.1	4.7	2.3	2.2	2.8	2.9	996.4	507.2
MT169	97.1	87.0	20.5	19.4	117.6	84.0	50.3	46.0	3.2	2.8	2.7	3.2	3.4	3.5	1970.9	1620.7
SHI165	95.5	80.2	25.7	24.6	134.4	104.9	57.0	45.0	3.6	3.7	2.5	2.4	3.6	3.5	1476.4	1690.0
VGR156	118.2	90.5	34.9	28.4	128.1	82.5	59.0	68.0	4.6	7.5	3.2	3.1	3.3	3.2	1926.6	1355.8
Mean	99.1	85.2	25.3	23.9	114.9	85.8	67.6	61.3	3.9	5.0	3.1	3.0	3.4	3.4	1731.0	1455.8
LSD(5%)	17.3	22.1	5.3	7.2	44.1	39.4	18.5	21.3	1.1	3.5	0.8	0.7	0.3	0.4	662.3	775.7

SLFC: stem length to the first capsule, PH: plant height, NCP: number of capsules per plant, NSC: number of seed per capsule, LC: length of the capsule, SW: 1000-seed weight.

Table 2. Yield components and yield of sesame genotypes grown under well watered (WW) and water stress (WS) conditions during 2011

Genotype	PH(cm)		SLFC(cm)		NCP		NSC		NB		1000-SW (g)		Yield(kg ha^{-1})	
	WW	WS	WW	WS	WW	WS	WW	WS	WW	WS	WW	WS	WW	WS
32-15	176	137.8	85.6	69.3	132.0	80.4	60.5	70.7	5.2	2.5	3.6	3.4	4270.0	2503.5
38-1-7	185.2	142.5	76.9	87.9	434.4	67.5	74.2	70.5	5.8	2.6	2.9	3.3	4281.9	2080.0
BC167	214.5	141.6	117.2	66.4	129.7	52.6	72.6	74.9	4.6	2.1	3.8	3.8	3494.2	1720.2
Birkan	124.5	116.8	31.2	37.6	76.3	79.4	74.4	78.6	2.9	3.5	3.7	3.9	2867.6	1759.2
EF147	148.2	112.8	61.5	50.3	89.6	50.7	73.6	72.4	3.2	2.6	3.8	3.7	1799	1505.4
EF153	218.7	151.2	122.8	94	151.0	96.6	68.8	74.5	4.1	4.2	2.9	3.4	3421.9	1649.2
HB168	185.1	134.6	87.7	62.4	122.2	73.7	61.7	66.9	3.5	2.9	3.7	3.6	3613.4	2227.0
HC107	142.4	131.5	72.9	70.7	95.6	68.7	63.0	67.1	4.8	1.9	3.7	3.4	1866.5	2561.4
HC108	186.6	147.6	105.7	72.8	111.5	84.1	70.3	72.3	5.0	2.5	3.3	3.2	3159.1	2216.4
HSC105	135.5	115.3	51.5	44.9	101.6	97.8	64.9	64.8	3.6	3.3	3.5	3.5	1964.5	2188.6
ICN115	124.6	117.9	42.8	39.8	96.6	90.6	74.9	68.2	3.9	2.9	3.7	3.3	2440.7	2090.8
ICN130	191.7	122.6	81.8	57.6	123.6	76.8	61.7	66.1	0.5	0.0	3.8	3.5	3305.5	1653.3
LC162	194.8	136.8	93.8	71.3	123.6	70.3	69.9	80.4	3.3	2.4	3.1	3.3	2715.2	2127.3
LC164	215.8	153.2	115.9	83.5	90.2	77.4	72.0	78.1	2.1	2.9	3.8	3.5	2581.6	1980.7
MC112	154.3	123.4	74.3	63.5	94.3	63.1	80.1	82.1	3.8	2.8	3.5	3.6	2134.5	1788.9
MC114	188.9	130.0	116.5	83.0	116.4	48.7	83.9	88.3	4.9	3.8	3.3	3.2	2381.7	1386.7
MT169	187.8	143.1	71.4	85.7	92.6	72.8	71.2	66.3	0.0	2.1	3.3	3.2	1441.1	1900.2
SHI165	172.2	131.1	87.1	62.6	90.8	81.1	57.5	71.9	1.9	1.5	3.5	3.3	2449.8	1599.9
VGR156	201.2	144.6	118.7	86.9	106.3	71.1	65.0	68.7	4.1	2.7	2.7	3.2	3383.7	1827.0
Mean	176.2	133.4	85	67.9	125.2	73.9	69.5	72.8	3.5	2.6	3.46	3.4	2793.7	1935.1
LSD (5%)	26.7	22.33	15.38	17.46	50.52	30.17	9.9	11.2	1.7	1.2	0.35	0.3	748.9	623.5

SLFC: stem length to the first capsule, PH: plant height, NCP: number of capsules per plant, NSC: number of seed per capsule, LC: length of the capsule, SW: 1000-seed weigh

Table 3. Simple correlation coefficients between Y_p, Y_s and drought tolerance/susceptibility indices of 19 sesame genotype in 2010

	Y_p	Y_s	SSI	STI	MP	GMP	TOL	YI
Ys	0.494*							
SSI	0.317	-0.630**						
STI	0.810**	0.894**	-0.235					
MP	0.868**	0.861**	-0.174	0.985**				
GMP	0.833**	0.891**	-0.231	0.988**	0.997**			
TOL	0.521*	-0.484*	0.937*	-0.063	0.029	-0.036		
YI	0.492*	1.000**	-0.632**	0.894**	0.860**	0.890**	-0.486*	
YSI	-0.316	0.631**	-1.000**	0.237	0.176	0.232	-0.937**	0.633**

Y_p: yield in optimal conditions, Y_s: yield under drought, SSI: stress susceptibility index, STI: stress tolerance index, MP: mean productivity, GMP: geometric mean productivity, TOL: tolerance, YI: yield index , YSI: yield stability index

Table 4. Simple correlation coefficients between Y_p, Y_s and drought tolerance/susceptibility indices of 19 sesame genotype in 2011

	Y_p	Y_S	SSI	STI	MP	GMP	TOL	YI
YS	0.186							
SSI	0.791**	-0.400						
STI	0.900**	0.576*	0.485*					
MP	0.940**	0.511*	0.552*	0.988**				
GMP	0.907**	0.578*	0.500*	0.994**	0.994**			
TOL	0.921**	-0.211	0.945**	0.667**	0.732**	0.673**		
YI	0.190	1.000**	-0.398	0.579**	0.514*	0.581**	-0.208	
YSI	-0.789**	0.402	-1.000**	-0.481*	-0.550*	-0.497*	-0.944**	0.400

Y_p: yield in optimal conditions, Y_s: yield under drought, SSI: stress susceptibility index, STI: stress tolerance index, MP: mean productivity, GMP: geometric mean productivity, TOL: tolerance, YI: yield index , YSI: yield stability index

Table 5. Ranks, ranks mean and standard deviation of ranks mean (SDR) of drought tolerance /susceptibility indices in 2010

Genotype	Y_p	Y_s	SSI	STI	MP	GMP	TOL	YI	YSI	Rank mean	SDR	FA1+FA2
32-15	15	15	12	16	16	16	11	15	12	14.22[16]	1.99	-1.08[16]
38-1-7	3	11	16	8	8	8	17	11	16	10.89[10]	4.70	-0.74[13]
BC167	9	3	2	5	5	5	3	3	2	4.11[3]	2.20	1.88[3]
Birkan	19	18	8	18	18	18	7	18	8	14.67[17]	5.27	-0.95[15]
EF147	6	4	9	4	4	4	10	4	9	6.00[4]	2.60	0.83[6]
EF153	11	6	7	9	9	9	8	6	7	8.00[7]	1.66	0.77[7]
HB168	1	9	19	3	3	3	19	9	19	9.44[9]	7.67	-0.61[12]
HC107	10	17	15	15	15	15	16	17	15	15.00[18]	2.06	-1.33[18]
HC108	16	7	1	12	12	12	1	7	1	7.67[6]	5.70	1.37[4]
HSC105	2	12	17	7	7	7	18	12	17	11.00[11]	5.61	-0.80[14]
ICN115	12	14	14	14	13	14	14	14	14	13.67[15]	0.71	-1.17[17]
ICN130	13	13	10	13	14	13	9	13	10	12.00[13]	1.80	-0.44[9]
LC162	4	2	5	2	2	2	5	2	5	3.22[2]	1.48	2.28[2]
LC164	5	1	4	1	1	1	2	1	4	2.22[1]	1.64	2.60[1]
MC112	17	16	6	17	17	17	6	16	6	13.11[14]	5.35	-0.52[11]
MC114	18	19	18	19	19	19	13	19	18	18.00[19]	1.94	-2.93[19]
MT169	7	8	11	6	6	6	12	8	11	8.33[8]	2.40	0.31[8]
SHI165	14	5	3	11	11	11	4	5	3	7.44[5]	4.25	1.06[5]
VGR156	8	10	13	10	10	10	15	10	13	11.00[11]	2.18	-0.51[10]

Table 6. Ranks, ranks mean and standard deviation of ranks mean (SDR) of drought tolerance /susceptibility indices in 2011

Genotype	Y_p	Y_s	SSI	STI	MP	GMP	TOL	YI	YSI	Rank mean	SDR	FA1+FA2
32-15	2	2	13	1	1	1	16	2	13	5.67[1]	6.32	3.07[1]
38-1-7	1	8	18	2	2	2	19	8	18	8.67[8]	7.70	2.07[2]
BC167	4	14	17	6	5	6	18	14	17	11.22[12]	5.85	0.44[6]
Birkan	9	13	12	12	10	12	12	13	12	11.67[14]	1.32	-0.16[12]
EF147	18	18	6	19	19	19	4	18	6	14.11[18]	6.62	-2.22[18]
EF153	5	16	19	8	7	8	17	16	19	12.78[16]	5.65	0.25[8]
HB168	3	3	11	3	3	3	13	3	11	5.89[2]	4.37	1.72[3]
HC107	17	1	1	13	13	13	1	1	1	6.78[4]	6.96	-0.43[13]
HC108	8	4	9	4	4	4	10	4	9	6.22[3]	2.68	1.24[4]
HSC105	16	5	3	14	14	14	3	5	3	8.56[7]	5.73	-0.79[14]
ICN115	13	7	4	11	12	11	6	7	4	8.33[6]	3.46	-0.07[10]
ICN130	7	15	16	9	8	9	15	15	16	12.22[15]	3.83	0.12[9]
LC162	10	6	7	7	9	7	7	6	7	7.33[5]	1.32	0.44[6]
LC164	11	9	8	10	11	10	8	9	8	9.33[9]	1.22	-0.08[11]
MC112	15	12	5	16	16	16	5	12	5	11.33[13]	5.00	-1.13[16]
MC114	14	19	14	17	17	17	11	19	14	15.78[19]	2.68	-1.47[17]
MT169	19	10	2	18	18	18	2	10	2	11.00[11]	7.55	-2.61[19]
SHI165	12	17	10	15	15	15	9	17	10	13.33[17]	3.12	-0.98[15]
VGR156	6	11	15	5	6	5	14	11	15	9.78[10]	4.32	0.58[5]

Table 7. Results of factor analysis for drought tolerance/susceptibility indices and yields of 19 sesame genotypes in two years

Index	2010			2011		
	Factor loading		Communalities	Factor loading		Communalities
	FA1	FA2		FA1	FA2	
Yp	0.875	-0.459	1	0.915	-0.368	1
Ys	0.851	0.521	1	-0.193	-0.981	1
SSI	-0.142	-0.987	1	0.97	0.216	0.983
STI	0.987	0.100	0.985	0.672	-0.718	1
MP	0.998	0.029	1	0.733	-0.664	0.999
GMP	0.995	0.088	0.999	0.685	-0.723	1
TOL	0.045	-0.973	1	0.987	0.023	1
YI	0.85	0.523	1	-0.19	-0.982	1
YSI	0.144	0.987	1	-0.97	-0.219	1
Variance	5.2139	3.6687		5.2254	3.6363	
Variance %	0.579	0.408		0.581	0.404	
Cumulative	0.579	0.987		0.581	0.985	

Table 8. Path coefficients of measured traits in mutant germplasm of sesame

Stem length to the first capsule vs yield	r = -0.195	Number of seeds per capsule vs yield	r = 0.157
Direct effect	- 0.283	Direct effect	0.176
Indirect effect via PH	0.221	Indirect effect via SLFC	-0.089
Indirect effect via NCP	0.100	Indirect effect via PH	0.106
Indirect effect via NSC	0.056	Indirect effect via NCP	-0.111
Indirect effect via LC	-0.092	Indirect effect via LC	0.099
Indirect effect via SW	-0.189	Indirect effect via SW	-0.021
Plant height vs yield	r = 0.409	Capsule length vs yield	r = 0.533
Direct effect	0.332	Direct effect	0.233
Indirect effect via SLFC	-0.188	Indirect effect via SLFC	0.112
Indirect effect via NCP	0.348	Indirect effect via PH	0.029
Indirect effect via NSC	0.056	Indirect effect via NCP	0.032
Indirect effect via LC	0.021	Indirect effect via NSC	0.075
Indirect effect via SW	-0.142	Indirect effect via SW	0.055
Number of cap. per plant vs yield	r = 0.531	Thousand seeds weight vs yield	r = 0.132
Direct effect	0.519	Direct effect	0.276
Indirect effect via SLFC	-0.055	Indirect effect via SLFC	0.193
Indirect effect via PH	0.223	Indirect effect via PH	-0.171
Indirect effect via NSC	-0.038	Indirect effect via NCP	-0.216
Indirect effect via LC	0.015	Indirect effect via NSC	-0.013
Indirect effect via SW	-0.115	Indirect effect via LC	0.046

SLFC: stem length to the first capsule, PH: plant height, NCP: number of capsules per plant, NSC: number of seed per capsule, LC: length of the capsule, SW: 1000-seed weight.

DISCUSSION

In sesame breeding, the goal is to attain high seed yield. The later character is therefore the most reliable measure for selecting for drought tolerance. Venuprazad *et al.*[35] stated that direct selection under dry season stress also gave similar response as under naturally occurring wet season stress. But our results are in contradiction with this later author according to a high genotype x season interaction (data not shown). In our research conditions, yield was significantly lower in drought-stressed conditions relative to non-stressed conditions. However, contrary to what was expected, some genotypes performed better under moderate drought-stressed conditions than in non-stressed conditions. Similar findings were reported by Urrea *et al.*[31] with the dry bean (*Phaseolus vulgaris* L.) cultivar SER 22 which performed well under drought-stressed conditions, but below average under non-stressed conditions. These genotypes are classified in the class C according to Fernandez[14].

In 2010, genotypes LC 164, LC 162, BC 167, EF 147 and MT 169 had the highest grain yield under both DS and NS environments, while in 2011 HC 108, 32-15, HB 168 and 38-1-7 had the highest grain yield irrespective to the environments. These genotypes could be therefore classified in the group A based on Fernandez[14] model. In contrast ICN 130, MC 112, 32-15, ICN 115, HC 107, Birkan, MC 114 and MT 169, MC 112, EF 147, SHI 165, MC 114 perform poorly in both DS and NS conditions in the first and second year, respectively, and were classified as group D.

HC 108, SHI 165, EF 153 and HC 107, LC 162, LC 164, HSC 105, ICN 115 perform favorably only when grown under DS conditions in 2010 and 2011, respectively, and were classified in the group C contrary to VGR 156, 38-1-7, HSC 105, HB 168 and Birkan, VGR 156, BC 167, ICN 130 and EF 153 which perform favorably only in NS environment and thus were classified in the group B according to Fernandez[14].

To determine the most suitable drought tolerant criteria, the correlation between Y_p, Y_s and other drought tolerance indices was computed. In other words, correlation studies between yield and drought tolerances indices can be a good criteria for screening the best genotypes and indices used[13]. Therefore, a discriminatory index must have a significant correlation with grain yield under both stressed and non-stressed environments[21].

STI, MP and GMP had significant and positive correlation with both yield under drought-stressed (Ys) and non-stressed environments in both two years. These results confirmed those of Abdolshahi *et al.*[1] stating that STI, MP, GMP and YI appeared to be the most efficient selection indices for identifying high yielding genotypes for both normal and drought-stressed environments.

GMP and STI had high correlation with MP and therefore STI, GMP and MP could produce similar results. Fernandez[14] stated that STI is estimated based on GMP and the correlation between STI and GMP is equal to 1. Akçura *et al.*[2] reported that YI, YSI, STI, GMP were significantly and positively correlated with stress yield and these indices showed that cultivars may be ranked only on the basis of their yield under stress and so does not discriminate genotypes of group A.

Based on STI, GMP and MP, LC 164 was the best drought tolerant genotype in the first year. Thus, it could be concluded that selection based on these indices results in genotypes with high yield potential as stated by Abdolshahi *et al.*[1]. STI is effective in selecting higher-yielding lines in both stressed and non-stressed environments and could thus discriminates group A with others (B,C,D). The higher the value of STI of a given genotype, the higher its stress tolerance and yield potential[14]. It is clear that STI is not efficient in selecting low yield lines even though their reduction percentage of seed yield across environments is lower. It's the case of genotypes hsc105 with yield reduction percentage (PR= 8 %, data not shown), mc112 (PR = 10 %) and mutant-cultivar Birkan (PR = 29 %).

The correlation between Y_p and Y_s was positively significant ($r = 0.49^*$) in 2010 but not in 2011. In other words, sesame genotypes with high yield potential may not necessarily perform favorably in drought-stressed environments. This result is supported by Belko *et al.*[6] who reported poor relationship between grain yield under NS and DS environments in both short and medium duration cowpea genotypes and opposed those claimed by certain authors who stated that genotypes with high yield potential are like to have high yield in drought-stressed conditions.

The correlation between YI and Y_s is equal to 1. Therefore, YI is a suitable criterion for drought tolerance. TOL had high positive correlation with Y_p and negative ones with Y_s. Fernandez[14] stated that selection based on TOL favours genotypes with low yield potential in non-stressed conditions and high yield under stress conditions. Based on these results STI, GMP and MP favour genotypes with high yield potential while TOL favours genotypes with low yield potential. Thus, different indices would not result in the same ranking.

Factor analysis and the mean ranking approach were used for selecting the suitable drought tolerant material across environments and years. These methods have the advantage to use all drought tolerance indices simultaneously. In the first year, LC 164 and LC 162 were identified as the best drought tolerant genotypes according to the two ranking methods.

In 2011, cultivar 32-15 was ranked first according to FA and the mean ranks method and thus identified as the most drought tolerant genotype. This ranking method was also used to identify drought-tolerant cultivars of bread wheat[13], spring canola cultivars[19] and Corn[23].

Genotypes ranking according to their drought tolerance/susceptibility were thus affected across the seasons. The two experiments were conducted in two different seasons contrasting for weather conditions. In 2010, the experiment was conducted in a hot and dry season under irrigated conditions while the second year corresponds to the normal rainy season in the semi-arid tropic. In other words, this seasonality may interact as a genotype by environment effect. The cultivar 32-15 is a well locally adapted variety grown largely by Senegalese sesame growers in rainfed conditions. All other genotypes were induced by mutagenesis from 32-15 and 38-1-7 as parents.

Path coefficient analysis in the present study showed that number of capsules per plant, plant height, thousand seed weight and length of the capsules were the most important components with direct and positive influence on seed yield in drought stress conditions. This was in accordance with the findings of Uzun and Cagirgan[34] and Yingzhong and Yishou[37]. Plant height is the character most contributing to seed yield in sesame because the species has an indeterminate growth habit[32]. Although this character prevents mechanized harvesting and the expansion of its cultivation, plant height may favoured high branching and capsule production. Thus, plant height, number of capsules per plant, and length of the capsules should be considered in selection for obtaining high- yielding sesame cultivars in drought-stressed environments. This was supported by the fact that plant height has a positive indirect effect (Table 8) on seed yield via number of capsules per plant. In other studies[27,37] higher number of capsules per plant and plant height showed a positive indirect effect on seed yield.

The correlation and the direct effect of stem length to the first capsule on seed yield were negative. However, the stem length to the first capsule had a positive indirect effect on seed yield via plant height with which it was strongly correlated. Thus, in indirect selection for high-yielding sesame cultivars, plant height and stem length to the first capsule traits should be considered together as a selection criterion.

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