

## Estimates of Genetic Variability, Correlation and Path Analysis in Sunflower (*Helianthus annuus* L.)

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**T**WELVE sunflower genotypes were evaluated for two years; 2013 and 2014 at three locations; Agricultural Research Stations of Shandaweel (location 1), Al-Arish (location 2) and El-Ewiyat east (location 3). Genotypic and phenotypic variances, their coefficient of variation, heritability and genetic advance were assessed. As well as study the interrelationships among oil yield attributers using genotypic and phenotypic correlation and path coefficients analysis were done. The experiments were conducted in a randomized complete block design with three replications in the field. Results proved that significant differences were observed among sunflower genotypes, for all studied characters in the three locations over two seasons. The elite genotype No 7 (Line120) surpassed the two check cultivar (Sakha 53 and Giza 102) recording the maximum values of seed yield in the first location with produced the maximum values in the second location for head diameter and seed yield and recording the maximum values of head diameter, seed yield and oil yield in the third location.

Estimates of heritability in broad sense at the three locations were varied from; 26 - 97%, 29 - 96% and 57-97%, respectively. Results also indicated that, highly significant and positive correlation coefficients were obtained between oil yield and some studied traits at the genotypic and phenotypic levels in the three locations. According to path coefficients analysis (at genotypic and phenotypic levels), the traits, *i.e.* days to 50% flowering and plant height in location 1, days to 50% flowering, plant height, head diameter and 100-seed weight in location 2 and 100-seed weight and seed yield/plant in location 3 were important predictors of oil yield indicating their magnitude as selection criteria to obtain a valuable gain of selection for oil yield in sunflower.

**Keywords:** Sunflower, Genetic parameters, Genotypic and phenotypic correlation coefficients, Path coefficients analysis, Oil yield components.

Sunflower (*Helianthus annuus* L.) is considered one of the most important edible oilseed crops after soybean and palm in the world. Its seeds contain a high content of a good quality oil, *i.e.* 40-50 % and ~20 % of protein. Sunflower

breeders focus all their interest in the development of sunflower genotypes with the maximum seed and oil yields. The variability of initial materials playing the major role in the success of any breeding program (Fick & Miller, 1997 and Vranceanu, 2000). Oil yield as a polygenic trait is influenced by several characters called oil yield contributing traits. These components are related among themselves and with oil yield either positively or negatively. Correlation analysis does not provide the clear picture of complex associations among the plant traits. Path coefficient analysis is considered as a more precise method to dividing the direct and indirect influences of independent variables upon the dependent variable. In this respect, the association between oil yield and some other attributers using correlation and path analysis was studied by Fick *et al.* (1974), Skoric (1974), Green (1980), Marinkovic (1992), Punnia & Gill (1994), El-Hosary *et al.* (1999), Farhatullah & Khalil (2006), Habib *et al.* (2007), Hlandi *et al.* (2010), Hassan *et al.* (2013) and Kang & Ahmed (2014).

The objectives of this research work were study the variations among the evaluated genotypes through estimates of genotypic and phenotypic variances ( $V_g$  and  $V_{ph}$ ), genotypic and phenotypic coefficient of variation (GCV and PCV), broad sense heritability and genetic advance as well as understanding the interrelationships among oil yield attributers via study the genotypic and phenotypic correlation and path coefficients analysis.

### Materials and Methods

The present work was performed at the three locations; Agricultural Research Stations of Shandaweel (location 1), Al-Arish (location 2) and El-Ewyinat east (location 3) during two summer seasons; 2013 and 2014. Some soil properties of the experimental sites are shown in Table 1.

**TABLE 1. Some soil properties of the experimental sites (average the two growing seasons).**

Locations Soil properties	Shandaweel	Al-Arish	El-Ewyinat east
Sand (%)	55.91	87.20	64.70
Silt (%)	11.84	7.20	28.00
Clay (%)	32.25	5.60	7.30
Soil texture	Sandy clay	Sandy	Sandy loam
pH (1:1)	7.60	8.37	8.27
EC (ds m <sup>-1</sup> )	0.39	1.77	0.69

The materials used in this work comprised twelve sunflower genotypes. The origins and some descriptions of these genotypes are given in Table 2. The treatments were distributed in a split plot design with three replications.

**TABLE 2. Origins and some descriptions of the used genotypes.**

No.	Genotype	Origin	Agronomic characteristics			
			Days to 50% flowering	Plant height (cm)	Stem diameter (cm)	Head diameter (cm)
1	Line125	Egypt	54.00	160.67	1.80	19.17
2	Line167	Egypt	53.00	159.33	2.27	16.87
3	Line235	Egypt	52.00	162.00	1.97	20.83
4	Line350	Egypt	55.00	148.33	1.97	19.33
5	Line460	Egypt	56.00	171.00	2.30	19.83
6	Line465	Egypt	56.33	185.67	2.40	20.93
7	Line120	Egypt	55.33	185.67	2.07	18.47
8	Line880	Egypt	56.00	196.00	2.40	20.40
9	Line885	Egypt	55.67	179.00	2.10	21.60
10	Line990	Egypt	58.00	132.33	1.63	19.30
11	Sakha 53	Local variety	52.00	168.33	2.07	22.30
12	Giza 102	Local variety	44.00	133.67	1.63	21.77

*Cultural practices*

Each sub-plot consisted of 4 ridges, 3 m length and 0.6 m apart (plot area = 7.2 m<sup>2</sup>). The agricultural practices were maintained as recommended for sunflower in the three locations. Surface irrigation system was applied in Shandaweel while in the others, sprinkler irrigation system was used. Days to 50% flowering as number of days from sowing to flowering 50 % of plants for each genotype were recorded. At harvest, 10 guarded plants were chosen from the inner two ridges to collect data on the following characters:

- 1- Plant height in cm (PH).
- 2- Head diameter in cm (HD).
- 3- Stem diameter in cm (SD).
- 4- 100-seed weight in grams (100- SW).
- 5- Seed yield/plant in grams (SY/P).
- 6- Seed oil content (oil %): Oil was extracted by using petroleum ether (60-80°C) and Soxhelt apparatus according to AOAC (1980).
- 7- Seed yield (SY) was primarily calculated from plot area and then converted to the unit of (kg/fed).
- 8- Oil yield (OY): It was calculated by multiplying seed yield (kg/fed) by seed oil percentage (%).

### *Statistical analysis*

Combined analysis of variance over growing seasons and locations was done as outlined by Snedecor & Cochran (1989). Significant differences among treatment means were detected using least significant difference test (LSD) at 5% probability level. Assumption of homogeneity of variances was examined before running the combined analysis according to Levene (1960). The interrelationships among oil yield and its related traits were studied at the genotypic and phenotypic levels using the following methodologies:

- 1- Correlation coefficients between all pairs of studied traits were computed as suggested by Johnson *et al.* (1955).
- 2- Path coefficient analysis was subjected as suggested by Wright (1921) and rediscovered by Dewey & Lu (1959). The method permits to separate the genotypic and phenotypic correlation coefficient between the oil yield (as a resultant variable) and each of related traits (as explanatory variables) into direct effect (path coefficient) and indirect effects (that exerted through the other variables). The genotypic and phenotypic variances ( $V_g$  and  $V_{ph}$ ), and genotypic and phenotypic coefficient of variation (GCV and PCV), broad sense of heritability were estimated according to Johnson *et al.* (1955). Genetic advance in terms of percentage of means (with 10 % selection intensity) was estimated as described by Brim *et al.* (1959).

A BASIC program (Atia, 2007) was used to automate the computations of genotypic and phenotypic correlation coefficients and their corresponding path analyses.

### **Results and Discussion**

The results of Levene test (1960) proved the homogeneity of variances over growing seasons and locations for all studied characters that permits to apply combined analysis. Accordingly, the mean values of oil yield and its related characters for twelve sunflower genotypes are listed in Table 3.

#### *Mean performance*

It is evident from Table 3 that the differences among studied genotypes were clear and significant for all studied characters indicating wide genetic variation among tested genotypes.

Results displayed that genotype No 10 (Line 990) had the latest in flowering recording (60.5, 48.7 and 54.8 days) over both seasons in the three locations, respectively. On the other hand, the genotype No 12 (Giza 102) was the earliest in flowering recording averages of (46.5, 37.2 and 42.3 days) over both seasons in the three locations, respectively.

**TABLE 3. The mean performance of the studied genotypes obtained from combined analysis over two seasons for some yield traits through three locations.**

Location	Genotypes	DF	PH	H D	SD	100-SW	SY/P	Oil %	SY (kg/ fed)	Oil Y (kg/fed)
Shandaweel	1	56.7 d	185.7 e	20.2 e	2.0 e	5.7 fg	62.0 f	31.8 f	1176.0 a	451.5 bcd
	2	55.5 e	184.3 g	17.9 g	2.5 b	7.6 a	69.3 c	38.4 c	1137.7 abc	448.0 cd
	3	54.5 f	187.2 c	21.8 c	2.2 d	5.7 fg	68.0 cd	37.5 c	1077.7 de	438.7 d
	4	57.8 c	173.3 e	20.3 e	2.2 p	5.6 g	59.0 g	38.1 c	1112.0 cd	449.0 cd
	5	58.5 bc	195.3 d	21.1 d	2.5 b	7.1 b	63.0 f	31.9 f	1158.0 ab	476.2 a
	6	58.8 b	210.7 c	21.9 c	2.6 a	7.4 ab	64.0 ef	32.7 f	1175.2 a	466.8 ab
	7	57.8 c	210.0 f	19.5 f	2.3 c	6.4 d	66.3 de	34.0 e	1179.3 a	460.6 abc
	8	58.5 bc	221.0 cd	21.4 cd	2.6 a	5.9 e	67.0 cd	35.2 d	1118.3 bcd	451.5 bcd
	9	58.2 bc	204.0 b	22.6 b	2.3 c	6.0 e	74.7 b	42.5 b	1065.0 e	440.8 d
	10	60.5 a	157.3 e	20.3 e	1.8 f	5.9 ef	78.0 a	46.2 a	1121.0 bcd	449.2 cd
	11	54.5 f	193.3 a	23.3 a	2.3 c	6.7 c	72.7 b	41.8 b	1142.7 abc	415.8 e
	12	46.5 g	158.7 ab	22.8 ab	1.8 f	7.4 a	78.67 a	46.8 a	987.3 f	388.7 f
Al-Arish	1	27.3 bc	88.3 g	11.2 h	1.6 e	4.4 e	26.4 d	34.0 c	527.7 b	213.4 a
	2	46.0 d	90.0 g	12.1 g	2.1 b	5.4 b	25.8 d	33.0 e	534.0 b	210.7 a
	3	39.8 f	101.7 e	9.1 i	1.8 d	4.5 cde	23.9 e	30.5 f	446.3 de	183.0 e
	4	45.8 d	96.7 f	13.8 e	1.8 d	4.4 de	31.1 c	37.9 d	453.7 d	176.8 ef
	5	46.8 c	110.0 d	14.7 cd	2.1 b	5.1 b	34.3 b	41.2 b	524.3 b	201.3 bcd
	6	48.0 ab	113.3 cd	15.0 bc	2.2 a	6.3 a	34.0 b	41.2 b	531.0 b	193.8 d
	7	47.0 c	116.7 bc	15.7 a	1.9 c	5.4 b	31.7 c	39.2c	555.0 a	205.5 abc
	8	45.3 d	118.3 b	14.6 d	2.2 a	4.8 c	31.0 c	37.8 d	520.2 b	198.3 cd
	9	47.3 bc	125.0 a	14.8 bcd	1.9 c	4.7 cd	35.2 b	41.7 b	430.7 e	161.5 g
	10	48.7 a	110.0 d	15.1 b	1.4 f	4.0 f	31.2 c	38.3 cd	457.3 d	173.3 f
	11	43.3 e	90.0 g	10.9 h	1.9 c	4.5 cde	34.7 b	41.5 b	500.7 c	182.5 e
	12	37.2 g	101.7 e	12.8 f	1.4 f	6.2 a	37.0 a	43.8 a	554.0 a	208.4 ab
El- Ewiyinat east	1	53.0 cd	155.0 f	14.0 f	2.1 e	5.4 e	29.0 f	39.7 cd	434.2 bc	303.1 e
	2	52.0 f	156.7 f	20.3 e	2.6 b	6.3 bc	50.0 ef	35.2 e	839.5 b	306.4 de
	3	46.0 h	168.3 d	17.5 g	2.3 d	5.7 d	47.0 g	41.7 b	765.5 f	308.8 d
	4	52.5 e	165.0 e	21.3 d	2.3 d	5.7 d	53.4 d	38.8 d	742.7 g	280.9 h
	5	53.2 c	175.0 c	22.2 c	2.6 ab	6.4 bc	59.5 a	40.8 b	820.5 d	311.9 cd
	6	53.8 b	172.5 c	22.5 bc	2.7 a	7.1 a	56.7 b	43.3 a	825.2 cd	314.9 c
	7	52.7 de	174.2 c	23.4 a	2.4 c	6.5 b	56.0 bc	35.7 e	862.5 a	337.9 a
	8	52.3 ef	180.0 b	22.7 bc	2.7 a	6.2 c	50.7 e	36.2 d	833.2 bc	331.9 b
	9	53.8 b	192.5 a	23.0 ab	2.4 c	5.8 d	58.8 a	33.3 f	720.5 h	262.9 I
	10	54.8 a	179.2 b	22.9 ab	1.9 f	5.4 e	54.7 cd	38.8 d	750.3 g	291.4 f
	11	49.5 g	157.5 f	18.9 f	2.4 c	5.9 d	56.9 b	43.2 a	791.2 e	286.1 g
	12	42.3 i	167.5 de	21.0 d	1.9 f	7.2 a	60.2 a	40.7 bc	843.8 b	310.7 cd

Means followed by the same letters within each column do not differ significantly according to Duncan's Multiple Range test at the 5% level.

Genotype No 11 check cultivar (Sakha 53) had the tallest plants recording (193.3 cm) at the first location, while Genotype No 9 (Line 885) had the highest values (125.0 and 192.5 cm) in the second and third location, respectively. Considering head diameter, check cultivar genotype Sakha 53 gave the highest values at the first location (23.3 cm) and genotype Line120 had the best value at the second and third locations with value being (15.7 and 23.4 cm), respectively, without significant differences between them.

The elite genotypes Line 465 and Line 880 surpassed the two check cultivars (Sakha 53 and Giza 102) recording the maximum values of stem diameter (2.6, 2.2 and 2.7 cm). For 100-seed weight, genotype no. 12 recorded the highest values (7.4 and 7.4 g) in the first and third locations, while in the second location genotype no. 6 came in the first (6.3 g). With respect to seed yield/plant, the best values reached to 78.0 and 78.67 g by the genotypes Line 990 and Giza 102 in the first location, 37.0 g by Giza 102 in the second location and 59.5, 58.8 and 60.2 g by Line 460, Line 885 and Giza 102 in the third location.

Regarding seed oil content percentage, the best genotypes were; Line 999 and Giza 102 (46.5 and 46.8%), Giza 102 (43.8%) and Line 465 and Sakha 53 (43.3 and 43.2%) at three locations, respectively.

Concerning Seed yield/fed., it is obvious that genotypes no.1, 6 and 7, no.7 and 12 and no. 7 had the maximum values (1176.0, 1175.2 and 1179.3 kg/fed), (555.0 and 554.0 kg/fed) and (862.5 kg/fed), respectively. With regard to oil yield, data in Table 3 indicated that genotypes no. 5, 1 and 2 and no. 7 surpassed the other genotypes, where it gave the highest estimates by 476.2 kg/fed, 213.4 and 210.7 kg/fed and 337.9 kg/fed at the three locations, respectively. Similar results have been concluded by Sharief (1998), Vega *et al.* (2002), Allam *et al.* (2003) and Abdou *et al.* (2011) who found significant differences among studied genotypes of sunflower for seed and oil yields and most studied characters.

#### *Genetic parameters*

Such considerable range of variation provided a good opportunity for yield improvement. Genotypic and phenotypic coefficients of variation (GCV and PCV), broad sense heritability ( $h^2$ ), and genetic advance expressed as percent of mean for the six characteristics are shown in Table 4. The phenotypic coefficient of variation (PCV) was generally higher than the genotypic coefficient of variation (GCV) for the studied characters, but in many cases, the two values differed only slightly which reflect some what the effect of environment on the expression of traits. According, the selection would be effective to improve these traits among the studied genotypes. Similar results were reported by Humera *et al.* (2014). High values of genotypic and phenotypic coefficients of variation were shown for head diameter (7.15, 15.88 and 8.61) and (7.69, 19.08 and 9.56), seed yield/plant (9.49, 14.79 and 7.71) and (10.14, 20.92 and 8.55), 100-seed weight (11.06, 14.18 and 8.94) and (11.94, 15.3 and 10.79), stem diameter

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(11.89, 14.05 and 11.05) and (12.07, 14.00 and 11.63), plant height (10.59, 12.33 and 5.98) and (12.25, 16.63 and 7.04), oil yield (5.12, 9.46 and 6.49) and (6.17, 15.23 and 7.55) and seed yield/fed (5.46, 8.87 and 5.71) and (6.17, 15.23 and 7.55) in the three locations, respectively.

While low estimates were observed with the seed oil content percentage (2.53, 3.50 and 3.34) and (5.00, 4.65 and 4.43) and days to 50% flowering (6.45, 7.60 and 7.01) and (6.58, 7.76 and 7.12) in the three locations, respectively.

With regard to the broad sense heritabilities, the data also revealed that estimates ranged from moderately to high for all studied traits at the three locations, except for seed oil content % and seed yield (kg/fed.) in the first and second locations, respectively which had low values. In order to predicting the selection effect, heritability accompanied with genetic advance is rather used than heritability alone.

In the same frame, it has been emphasized that without considering genetic advance, the heritability values ( $h^2$ ) would not be practically important in selection based on phenotypic appearance (Johnson *et al.*, 1955).

The data in the Table 4 present that, high values of heritability accompanied with high values of genetic advance (as % of mean) were obtained with stem diameter and weight of 100 seeds in three locations, indicating the importance of the additive gene effects. High heritability in the third location for stem diameter (90%) and head diameter (81%) coupled with high genetic advance (18.48 %) and (13.64%), respectively. High heritability estimates accompanied with high genetic advance is rather useful than high heritability alone for predicting the selection effect (Farhatullah *et al.*, 2006 and Humera *et al.*, 2014).

As shown in Table 4, high estimates of heritability accompanied with moderate genetic advance were observed with seed yield/plant in the first location, with days to 50% flowering in the second location, also in the third location, seed yield/plant had high estimates of heritability coupled with moderate genetic advance. These results indicate that additive gene effects are more important and the improvement can be applied through selection for this trait (Labana *et al.*, 1980). On the other hand, low values of heritability accompanied with low genetic advance in the three locations were obtained with seed oil percentage indicative of non-additive gene effects. Therefore, a limited scope for improvement of this trait is expected under the studied genotypes. The current conclusions are supported by Johnson *et al.* (1955) who emphasized that selection can be making safely with high values of heritability and genetic advance.

**TABLE 4. Estimates of genetic parameters for some yield traits in sunflower genotypes through three locations over two seasons.**

Traits	Locations	Genetic parameters				
		Mean	GCV	PCV	h <sup>2</sup>	GA (% mean)
DF	Shandaweel	56.49	6.45	6.58	0.96	11.14
	El-arash	45.22	7.60	7.76	<b>0.96</b>	<b>13.11</b>
	El- Ewyinat east	51.33	7.01	7.12	0.97	12.14
PH	Shandaweel	190.07	10.59	12.25	0.75	16.10
	El-arash	105.14	12.33	16.63	0.55	16.09
	El- Ewyinat east	170.28	5.98	7.04	0.72	8.93
H D	Shandaweel	21.09	7.15	7.69	0.86	11.69
	El-arash	13.31	15.88	19.08	0.69	23.25
	El- Ewyinat east	21.23	8.61	9.56	0.81	13.64
S D	Shandaweel	2.25	11.89	12.07	0.97	20.60
	El-arash	1.85	14.05	14.00	0.91	23.53
	El- Ewyinat east	2.35	11.05	11.63	<b>0.90</b>	<b>18.48</b>
100-SW	Shandaweel	6.46	11.06	11.49	0.93	18.73
	El-arash	4.96	14.18	15.03	0.89	23.55
	El- Ewyinat east	6.12	8.94	10.79	0.69	13.04
SY/P	Shandaweel	68.56	9.47	10.14	<b>0.87</b>	<b>15.58</b>
	El-arash	31.37	14.79	20.92	0.50	18.40
	El- Ewyinat east	54.40	7.71	8.55	<b>0.81</b>	<b>12.24</b>
Oil %	Shandaweel	39.81	2.53	5.00	<b>0.26</b>	<b>2.25</b>
	El-arash	38.36	3.50	4.65	<b>0.57</b>	<b>4.65</b>
	El- Ewyinat east	37.88	3.34	4.43	<b>0.57</b>	<b>4.42</b>
SY(kg/fed)	Shandaweel	1120.8	5.46	6.47	0.71	8.11
	El-arash	502.90	8.87	16.35	0.29	8.46
	El- Ewyinat east	802.42	5.71	6.05	0.89	9.50
Oil Y (kg/fed)	Shandaweel	444.73	5.12	6.17	0.69	7.49
	El-arash	192.37	9.46	15.23	0.39	10.35
	El- Ewyinat east	303.91	6.49	7.55	0.74	9.82

*Correlation matrix*

Genotypic (above diagonal) and phenotypic (below diagonal) correlation coefficients among oil yield and its related characters, estimated in location 1, location 2 and location 3 are given in Table 5. Generally, there was clear convergence between most genotypic and phenotypic correlation coefficients considering the value or sign in three locations indicating that the observed associations among most studied characters may be mostly attributed to genetic effects. Results showed that the most effective relationships to sunflower breeder, in location1, were those between oil yield and each of 50% flowering (0.91\*\* and 0.76\*\*), plant height (0.689\*\* and 0.324\*\*), stem diameter (0.609\*\* and 0.527\*\*), head diameter (-0.498\*\* and -0.29\*) and seed yield/ plant (-0.473\*\* and -0.443\*\*) at *Egypt. J. Agron.* **37**, No.2 (2015)



the genotypic and phenotypic levels, respectively. The high positive genotypic correlation between each of the aforementioned characters and oil yield reflected the inherent associations; therefore, the breeder can obtain high yielding genotypes through selection for one or more of these characters, especially if they proved to be more contributors to yield variation as lately shown.

On the other side, the yield contributors exhibited various trends of correlations among themselves. There was negative and highly significant genotypic or phenotypic association between 50% flowering and each of head diameter (-0.3\*\* and -0.26\*), 100-seed weight (-0.34\*\* and -0.33\*\*) and seed weight /plant (-0.28\* and -0.27\*) and plant height with seed weight/ plant (-0.27\* and -0.27\*) while on the reverse, 50% flowering had positive and highly significant genotypic and phenotypic associations with each of plant height (0.50\*\* and 0.42\*\*) and stem diameter (0.45\*\* and 0.44\*\*). However, plant height was found to be highly significant and positively correlated with stem diameter (0.90\*\* and 0.74\*\*) and the same results showed with stem diameter with 100-seed weight (0.33\*\* and 0.31\*\*) at the genotypic and phenotypic, respectively.

**TABLE 5. Genotypic (above diagonal) and phenotypic (below diagonal) correlation coefficients among oil yield (kg/fed) and its related characters evaluated individually under three locations over the two seasons.**

Locations	Traits	DF	PH	H D	S D	100-SW	SY/P	Oil Y
Shandaweel	DF	1	<b>0.50**</b>	<b>-0.3**</b>	<b>0.45**</b>	<b>-0.34**</b>	<b>-0.28*</b>	0.91**
	PH	<b>0.42**</b>	1	<b>0.19</b>	<b>0.90**</b>	0.04	<b>-0.27*</b>	0.69**
	H D	<b>-0.26*</b>	0.05	1	-0.04	0.01	<b>-0.34**</b>	-0.498**
	S D	<b>0.44**</b>	<b>0.74**</b>	-0.03	1	<b>0.33**</b>	<b>0.25*</b>	0.609**
	100-SW	<b>-0.33**</b>	0.06	-0.01	<b>0.31**</b>	1	<b>0.25*</b>	-0.069
	SY/P	<b>-0.27*</b>	-0.10	<b>0.32**</b>	<b>-0.33**</b>	0.21	1	-0.473**
	Oil Y	<b>0.76**</b>	0.324**	<b>-0.29*</b>	<b>0.527**</b>	-0.112	-0.443**	
	Al-Araish	DF	1	<b>0.37**</b>	<b>0.64**</b>	<b>0.37**</b>	-0.22	0.09
PH	<b>0.29*</b>	1	<b>0.78**</b>	<b>0.46**</b>	0.22	<b>0.55**</b>	<b>-0.098</b>	
H D	<b>0.54**</b>	<b>0.83**</b>	1	<b>0.36**</b>	<b>0.29**</b>	<b>0.62**</b>	<b>0.032</b>	
S D	<b>0.34**</b>	0.18	0.18	1	<b>0.34**</b>	0.21	0.326**	
100-SW	0.22	<b>0.25*</b>	<b>0.29**</b>	<b>0.27*</b>	1	<b>0.42**</b>	<b>0.519**</b>	
SY/P	<b>0.86**</b>	<b>0.74**</b>	<b>0.73**</b>	-0.01	<b>0.37**</b>	1	0.085	
Oil Y	<b>0.022</b>	<b>0.438**</b>	<b>0.426**</b>	0.038	<b>0.413**</b>	0.561**		
El-ewynat east	DF	1	<b>0.31*</b>	<b>0.56**</b>	<b>0.40**</b>	<b>-0.37**</b>	-0.01	-0.129
	PH	<b>0.28*</b>	1	<b>0.74**</b>	0.16	-0.04	<b>0.41**</b>	-0.166
	H D	<b>0.49**</b>	<b>0.70**</b>	1	<b>0.23*</b>	<b>0.28*</b>	<b>0.58**</b>	0.082
	S D	<b>0.37**</b>	0.19	<b>0.27*</b>	1	<b>0.34**</b>	-0.01	0.301**
	100-SW	<b>-0.26*</b>	0.14	<b>0.35**</b>	<b>0.35**</b>	1	<b>0.53**</b>	0.493**
	SY/P	-0.02	<b>0.37**</b>	<b>0.50**</b>	0.01	<b>0.42**</b>	1	-0.198
	Oil Y	-0.108	<b>-0.001</b>	<b>0.223</b>	0.310**	<b>0.51**</b>	-0.122	

\* and \*\*: Significant and highly significant at 0.05 and 0.01 probability levels, respectively.

Results also cleared that the most effective relationships in location 2, were those observed between oil yield and each of 100-seed weight (0.519\*\* and 0.413\*\*) at the genotypic and phenotypic levels, respectively. Furthermore, high positive and highly significant genotypic correlation were obtained between oil yield and each of stem diameter (0.326\*\*) and 100-seed weight (0.519\*\*). Also, high positive and highly significant phenotypic correlation between oil yield and each of plant height (0.438\*\*), head diameter (0.426\*\*), 100-seed weight (0.413\*\*) and seed yield/plant (0.561\*\*).

The yield contributors exhibited various trends of correlations among themselves. There was positive and highly significant genotypic or phenotypic association between days to 50% flowering had positive and highly significant genotypic and phenotypic associations with each of plant height (0.37\*\* and 0.29\*), head diameter (0.64\*\*, 0.54\*\*) and stem diameter (0.37\*\* and 0.34\*\*). However, plant height was found to be highly significant and positively correlated with head diameter (0.78\*\* and 0.83\*\*) and seed weight /plant (0.55\*\* and 0.74\*\*). Head diameter was found to be highly significant and positively correlated with 100-seed weight (0.29\*\* and 0.29\*\*) and seed yield/plant (0.62\*\* and 0.73\*\*). Also highly significant positive correlation were found between stem diameter and 100-seed weight (0.34\*\* and 0.27\*\*) and between 100-seed weight and seed weight /plant (0.42\*\* and 0.37\*\*) at the genotypic and phenotypic levels, respectively.

The results in Table 5 also proved that the relationships in location 3. Highly significant positive associations were obtained between oil yield and each of stem diameter (0.301\*\* and 0.310\*\*) and 100-seed weight (0.493\*\* and 0.510\*\*) at the genotypic and phenotypic levels, respectively. The high positive genotypic correlation between each of the aforementioned characters and oil yield reflected the inherent associations; therefore, the breeder can obtain high yielding genotypes through selection for one or more of these characters, especially if they proved to be more contributors to yield variation as lately shown.

On the other side, the yield attributers exhibited various trends of correlations among themselves. The remainder correlation coefficients among studied characters were mostly negligible and insignificant. Days to 50% flowering had positive and highly significant genotypic and phenotypic associations with each of head diameter (0.56\*\* and 0.49\*\*) and stem diameter (0.40\*\* and 0.37\*\*). Also, plant height was found to be highly significant and positively correlated with stem diameter (0.74\*\* and 0.70\*\*) and seed weight /plant (0.41\*\* and 0.37\*\*) at the genotypic and phenotypic levels, respectively. Highly significant positive correlations were also found between head diameter and each of with stem diameter (0.23\* and 0.27\*), 100-seed weight (0.28\* and 0.35\*\*) and seed yield/plant (0.58\*\* and 0.50\*\*) at the genotypic and phenotypic levels, respectively. The same trend were obtained between stem diameter and 100-seed weight (0.34\*\* and 0.35\*\*), and between 100-seed weight and seed weight / plant (0.53\*\* and 0.42\*\*) at the genotypic and phenotypic, respectively.

Generally, the highly significant positive genotypic relationship between any characters indicates that the improvement predicted under selection for one of them, would automatically extended to the other. These findings are in conflict with those obtained by Hladni *et al.* (2010) who found highly significant and negatively associations between stem diameter, total leaf area, head diameter and 100 seed weight on one side and seed oil content on the other. This discrepancy in results may be attributed to the used breeding materials and the environmental conditions. In fact, decisions of selection depending only on correlation coefficients may not always be effective because it measures the mutual association between a pair of traits neglecting the complicated interrelationships among all traits (Kang, 1994). Therefore, the correlation procedure may not provide a deep imagine about the importance of each component in the structure of oil yield. The path analysis can efficiently play this vital role.

#### *Path analysis*

Information that has been obtained from the correlations can be augmented by dividing correlations into direct and indirect influences of a particular set of causal interrelationships.

In such cases, the correlation coefficients may be confounded with indirect effects due to common association inherent in trait interrelationships. So, path coefficient analysis has proven useful in giving more information that describes the casual relationships such as yield and its attributers. In the present study, the resultant variable was oil yield while the remaining characters represented the casual variables. The matrix of direct and joint effects six predictor characters on oil yield is shown in Table 6.

Positive direct effects were recorded for all oil yield attributes except head diameter and seed yield/plant (-0.37 and -0.15) in genotypic level, plant height, head diameter and seed weight /plant (-0.18, -0.02 and -0.20) in phenotypic path coefficients at location 1. Results in Table 6 showed that positive direct effects for all oil yield attributes except plant height, head diameter (-0.31 and -0.20) in genotypic level and head diameter and stem diameter (-0.019 and -0.09) in phenotypic path coefficients at location 2.

On the other hand, data of location 3 cleared that positive direct effects were obtained for all oil yield attributes except days to 50% flowering , plant height and seed yield/plant (-0.10, -0.14 and -0.71) and (-0.31, -0.24 and -0.51) in genotypic and phenotypic levels, respectively.

The maximum direct effects were observed with days to 50% flowering (0.65 and 0.70) at location 1 and with 100-seed weight at location 2 and 3 (0.62 and 0.32) and (0.69 and 0.40) considering the genotypic and phenotypic levels.

It is noticed that, the path analysis gave different picture from what the correlation coefficient did. For example, the simple correlation coefficients (at genotypic and phenotypic levels) between oil yield and each of plant height and stem diameter which were positive and highly significant at location 1, also between oil yield and 100-seed weight at location 2 and also between oil yield and each of stem diameter and 100-seed weight at location 3 (Table 5). However, separation of the indirect effects from correlation coefficients through the path analysis gave different picture, where these traits had trivial effect on oil yield (Table 6).

**TABLE 6. The direct and indirect effects of six predictor characters on oil yield (kg/fed) at genotypic and phenotypic levels evaluated individually under three locations over the two seasons.**

Location	Level	Traits	DF	PH	H D	S D	100-SW	SY/P
Shandaweel	Genotypic	DF	<b>0.65</b>	0.16	0.08	0.02	-0.02	0.03
		PH	<b>0.33</b>	<b>0.44</b>	-0.05	0.01	0.01	0.06
		H D	-0.19	0.08	<b>-0.37</b>	-0.01	0.01	-0.04
		S D	0.29	0.40	0.02	<b>0.05</b>	0.02	0.04
		100-SW	-0.22	0.02	-0.01	0.02	<b>0.09</b>	-0.03
		SY/P	-0.18	-0.12	-0.15	-0.02	0.02	<b>-0.15</b>
	Phenotypic	DF	<b>0.70</b>	-0.08	0.01	0.11	-0.30	0.05
		PH	<b>0.29</b>	<b>-0.18</b>	-0.01	0.19	0.01	<u>0.02</u>
		H D	-0.18	-0.01	<b>-0.02</b>	0.01	0.01	-0.07
		S D	0.31	-0.13	0.01	<b>0.26</b>	0.23	0.07
		100-SW	-0.23	-0.01	0.01	0.08	<b>0.09</b>	-0.04
		SY/P	-0.19	0.02	-0.08	-0.08	0.12	<b>-0.20</b>
Al-Araish	Genotypic	DF	<b>0.30</b>	-0.12	-0.12	0.08	-0.14	0.01
		PH	0.11	<b>-0.31</b>	-0.15	0.10	0.13	0.02
		H D	0.19	-0.24	<b>-0.20</b>	0.08	0.18	0.03
		S D	0.11	-0.14	-0.07	<b>0.21</b>	0.21	0.01
		100-SW	-0.07	-0.07	-0.06	0.07	<b>0.62</b>	0.02
		SY/P	0.03	-0.17	-0.12	0.05	<b>0.26</b>	<b>0.04</b>
	Phenotypic	DF	<b>0.14</b>	0.04	-0.10	-0.23	-0.07	0.04
		PH	0.04	<b>0.15</b>	-0.16	-0.02	0.08	0.34
		H D	0.08	0.12	<b>-0.19</b>	-0.02	0.10	0.34
		S D	0.05	0.03	-0.03	<b>-0.09</b>	0.09	-0.01
		100-SW	-0.03	0.04	-0.06	-0.02	<b>0.32</b>	0.17
		SY/P	0.01	0.11	-0.14	.01	<b>0.12</b>	<b>0.46</b>
El-ewynat east	Genotypic	DF	<b>-0.10</b>	-0.04	0.22	0.01	-0.23	0.01
		PH	-0.03	<b>-0.14</b>	0.29	0.01	-0.02	-0.27
		H D	-0.05	-0.10	<b>0.43</b>	0.01	0.18	-0.39
		S D	-0.04	-0.02	0.10	<b>0.04</b>	0.22	0.01
		100-SW	0.04	0.01	0.12	0.01	<b>0.69</b>	-0.35
		SY/P	0.01	-0.06	0.25	0.01	<b>0.37</b>	<b>-0.71</b>
	Phenotypic	DF	<b>-0.31</b>	-0.07	0.30	0.07	-0.11	0.01
		PH	-0.09	<b>-0.24</b>	-0.42	0.03	0.06	-0.17
		H D	-0.13	-0.17	<b>0.60</b>	0.05	0.14	-0.25
		S D	-0.12	-0.04	0.16	<b>0.18</b>	0.14	-0.01
		100-SW	0.08	-0.03	0.21	0.06	<b>0.40</b>	-0.21
		SY/P	0.01	-0.09	0.29	0.01	<b>0.17</b>	<b>-0.51</b>

Shadwy and bold cell indicate to the direct effects.

Concerning the indirect effects in location 1, it is noted that the highest positive effect on oil yield were recorded from steam diameter through each of plant height and days to 50% flowering ( 0.40 and 0.29) at genotypic level, while via each of days to 50% flowering and 100-seed weight (0.31 and 0.23) at genotypic and phenotypic levels, respectively. Furthermore, high values of genotypic and phenotypic positive indirect effects were noticed with days to 50% flowering via plant height and plant height via steam diameter. However, 100-seed weight had maximum negative indirect effects on oil yield via days to 50% flowering (-0.22 and -0.23) at genotypic and phenotypic levels, respectively. In location 2, the results revealed that, steam diameter via 100-seed weight had the highest genotypic positive indirect effect on oil yield (0.21). On the other hand, the highest phenotypic indirect effects were recorded from each of plant height and head diameter through seed yield/plant (0.34). Furthermore, high estimates of genotypic and phenotypic positive indirect effects were observed by head diameter through each of days to 50% flowering and 100-seed weight and by 100-seed weight via seed weight/plant. However, the maximum negative indirect effects on oil yield were obtained from head diameter via plant height (-0.24) and days to 50% flowering through steam diameter (-0.23) at genotypic and phenotypic levels, respectively. Regarding location 3, path analysis proved that plant height followed by seed yield/plant via head diameter had the highest genotypic positive indirect effects on oil yield (0.29 and 0.25). On the other side, each of days to 50% flowering followed by seed yield/plant through head diameter gave the maximum phenotypic positive indirect effects on oil yield (0.30 and 0.29). Furthermore, high estimates of genotypic and phenotypic positive indirect effects were observed by days to 50% flowering through head diameter, steam diameter via 100-seed weight and head diameter via 100-seed weight and each of 100-seed weight and steam diameter through head diameter. However, the highest negative indirect effects on oil yield were recorded from head diameter via seed yield/plant (-0.39) and from plant height via head diameter (-0.42) at genotypic and phenotypic levels, respectively. The remainder indirect effects were very small and little importance. An overall view on the results of path analysis, it is proved that the traits, *i.e.* days to 50% flowering, head diameter, steam diameter, 100 seed weight and seed yield/plant are considered as direct and indirect selection criteria for oil yield in sunflower. Similar results were reported by Punnia & Gill (1994), El-Hosary *et al.* (1999), Hladni *et al.* (2010) and Hassan *et al.* (2013) who confirmed the importance of path analysis when deciding upon selection criteria using yield components.

The relative importance (RI %) according to genotypic and phenotypic path analysis are presented in Table 7. It is evident that the most oil yield variation (genotypic and phenotypic) was explained by the direct effects for 50% flowering (28.10 and 33.14) followed by plant height (13.21 and 2.22). For location 2 it is evident that the most oil yield variation (genotypic and phenotypic) was explained by the direct effects for weight 100 seeds (19.43 and

6.39) followed by plant height (4.81 and 1.39), 50% flowering (4.40 and 1.28) and head diameter (1.94 and 2.25). The data of relative importance (RI %) also revealed that for location 3 the most oil yield variation (genotypic and phenotypic) was explained by the direct effects for seed yield/plant (17.43 and 8.87) followed by weight 100 seeds (16.01 and 5.65).

**TABLE 7. The relative importance (RI %) of six predictor characters on oil yield (kg/fed) at genotypic and phenotypic levels evaluated individually under three locations over the two seasons**

Locations		Shandaweel		Al-Araish		El-ewynat east	
		G	Ph	G	Ph	G	Ph
DF		<b>28.10</b>	<b>33.14</b>	<b>4.40</b>	1.28	0.32	<b>3.30</b>
PH		<b>13.21</b>	2.22	<b>4.81</b>	1.39	0.67	1.97
H D		<b>9.33</b>	0.04	1.94	<b>2.25</b>	<b>6.26</b>	<b>12.50</b>
H D		0.14	<b>4.44</b>	2.23	0.48	0.05	1.08
100-SW		0.60	0.58	<b>19.43</b>	<b>6.39</b>	<b>16.01</b>	<b>5.65</b>
SY/P		1.56	<b>2.85</b>	0.09	<b>13.55</b>	<b>17.43</b>	<b>8.87</b>
X1	X2	<b>13.84</b>	<b>7.19</b>	3.42	0.77	0.27	1.41
X1	X3	<b>6.94</b>	0.60	3.71	1.83	<b>11.45</b>	<b>6.33</b>
X1	X4	1.26	<b>10.72</b>	2.30	0.53	0.10	1.41
X1	X5	1.99	2.90	4.14	1.67	1.54	2.25
X1	X6	<b>2.68</b>	<b>5.15</b>	0.12	0.71	0.05	0.21
X2	X3	<b>2.95</b>	0.03	4.75	2.92	<b>2.79</b>	<b>6.90</b>
X2	X4	1.72	<b>4.63</b>	2.98	0.30	0.05	0.54
X2	X5	0.17	0.13	4.20	1.48	0.21	0.96
X2	X6	1.77	0.51	0.71	6.41	<b>2.61</b>	<b>3.08</b>
X3	X4	0.07	0.03	1.50	0.37	0.26	1.94
X3	X5	0.03	0.01	3.53	2.23	<b>5.17</b>	<b>5.90</b>
X3	X6	<b>2.20</b>	0.21	0.51	8.09	<b>11.30</b>	<b>10.32</b>
X4	X5	0.14	0.99	4.45	0.94	0.58	1.72
X4	X6	0.22	<b>2.35</b>	0.19	0.02	0.03	0.09
X5	X6	0.34	0.54	1.10	6.85	<b>16.49</b>	<b>5.93</b>
Direct + Indirect		<b>89.23</b>	<b>79.23</b>	70.51	<b>60.02</b>	83.63	82.33
Residuals		10.77	20.77	29.49	39.80	16.37	17.67
<b>Total</b>		<b>100</b>	<b>100</b>	<b>100</b>	100	<b>100</b>	<b>100</b>

Also, the great genotypic and phenotypic components of joint effects were expressed by 50% flowering on oil yield via its association with plant height (13.84 and 7.19), stem diameter (1.26 and 10.72) and seed yield/plant (2.68 and 5.15) and by plant height via stem diameter (1.72 and 4.63) in location 1. For location 2 the great genotypic and phenotypic components of joint effects were expressed by 50% flowering on oil yield via its association with head diameter (3.71 and 1.83) and weight 100 seeds (4.14 and 1.67), plant height via head diameter (4.75 and 2.92) and weight 100 seeds (4.20 and 1.48), head diameter via weight 100 seeds (3.53 and 2.23) and weight 100 seeds via seed yield/plant (1.10 and 6.85). Moreover, in location 3 great genotypic and phenotypic components of joint effects were expressed by 50% flowering on oil yield via its association with head diameter (11.45 and 6.33), and weight 100 seeds (1.54 and 2.25), plant height via head diameter (2.79 and 6.90) and seed yield/plant (2.61 and 3.08), head diameter via weight 100 seeds (5.17 and 5.90) and seed yield/plant (11.30 and 10.32), and weight 100 seeds via seed yield/plant (16.49 and 5.93).

Negligible values of relative importance were observed for the other direct and indirect influences. Totally, the studied six characters explained (89.23% and 79.23 %), (70.51% and 60.02 %) and (83.63% and 82.33 %) in the three locations, respectively of oil yield variation at the genotypic and phenotypic levels, respectively. In accordance, the residual part may be attributed to unknown variation (random error), committing of errors during measuring the studied characters and/or some other traits that were not incorporated in the present study.

In conclusion, among the studied characters days to 50% flowering, weight 100 seeds and seed yield/plant were the most reliable oil yield components as selection criteria in sunflower breeding programs.

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## تقدير الاختلافات الوراثية و الارتباط و معامل المرور في دوار الشمس

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تم تقييم 12 تركيب وراثي من دوار الشمس خلال الموسم الصيفي لعامي 2013 و2014 م بمحطات البحوث الزراعية بشندويل والعريش وشرق العينات باستخدام تصميم القطاعات كاملة العشوائية في ثلاثة مكررات.

استهدف العمل البحثي دراسة التباين عبر تقدير معاملات الاختلاف المظهرية والوراثية ، معامل التوريث و درجة التحسين الوراثي مع تحديد أكثر الصفات اسهاما في محصول الزيت عبر دراسة الارتباط وتحليل معامل المرور .

وتتلخص أهم النتائج المتحصل عليها فيما يلي:

وجود فروق معنوية بين التراكيب الوراثية تحت الدراسة لجميع الصفات في المواقع الثلاثة. حيث أعطى التركيب الوراثي رقم 7 (**Line120**) أعلى قيمة لمحصول البذور في الموقع الأول بينما أعطى أعلى معدل لكل من قطر الساق و محصول البذور في الموقع الثاني وسجل تفوق ملحوظ في الموقع الثالث لكل من قطر الساق و محصول البذور و محصول الزيت مما يشير إلى تفوقه و كونه من التراكيب الوراثية المبشرة التي يوصى بالتوسع في إستخدامها في برامج تربية عباد الشمس.

تراوحت تقديرات درجات التوريث من 26 - 97% و 29 - 96% و 57 - 97% في المواقع الثلاثة على التوالي . كما أظهرت النتائج وجود علاقة ارتباط موجبة عالية المعنوية بين محصول الزيت و مكوناته على المستويين الوراثي والمظهري في المواقع الثلاثة. كما أشارت نتائج تحليل معامل المرور إلى أن صفتي 50% تزهير وارتفاع النبات في الموقع الأول ، و صفات عدد الأيام حتى 50% تزهير و ارتفاع النبات و قطر القرص و وزن 100 بذرة في الموقع الثاني و صفتي وزن 100 بذرة و محصول بذور النبات في الموقع الثالث كانت الأكثر اسهاما في محصول الزيت سواء عن طريق التأثير المباشر او غير المباشر (وراثيا و مظهريا) , مما يشير إلى أهمية وضع هذه الصفات في الاعتبار كمعايير انتخابية لتحسين محصول الزيت في عباد الشمس وذلك بالنسبة لكل موقع .