

Mass Selection and Individual Plant Selection as Two Breeding Methods for Improving Lettuce (*Lactuca sativa* L.)

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ABSTRACT

Three field experiments were conducted in order to estimate the magnitudes of variability of some important characters of the "Balady" cultivar of lettuce and to study the efficiency of two cycles of mass selection and individual plant selection for two generations, as two breeding methods, on the studied economical characters. Heritability percentages in broad sense and phenotypic correlation coefficients among all possible pairs, of the studied characters were also estimated. This study was carried out at the Experimental Station Farm (at Abies), Faculty of Agriculture, Alexandria University, during the three successive winter seasons of 2010/2011, 2011/2012 and 2012/2013.

The obtained results, generally, illustrated that the estimated coefficients of variability and ranges for all characters; reflected high and enough variability to suggest high possibilities of conducting successful and individual plant selection to produce improved strains with better performances than the original population. Generally, the results indicated clearly that the studied characters were improved through the two practiced breeding methods; mass selection and individual-plant selection; but, with different magnitudes among the characters and the used breeding methods. The obtained results showed high heritability estimates, in broad sense, for the various studied characters. Also, the estimated values for the correlation coefficients, between the different pairs of characters, were found to be either significant and highly significant with positive or negative signs, which may help in selecting desirable traits in lettuce breeding programs.

Accordingly, improving production of lettuce can be achieved through purification of already established cultivars and introducing improved strains through *via* both mass and individual-plant selection programs.

Key words: lettuce, yield and its components, selection methods, variability, heritability, correlation

INTRODUCTION

Lettuce (*Lactuca sativa*, $2n = 18$) is a highly self-pollinated crop that originated in the Mediterranean area. It is belonging to the family Compositae and considered as the most important vegetable in the group of leafy vegetables in many regions of the world. Lettuce is an excellent source of vitamins and nutrients which are highly required for human health, and because of high cellulose content, it facilitates digestion. Moreover, lettuce contains lactucin and lactucopicorn which improve calm sleep (Sharma, 2002).

The literature illustrated that improvement of lettuce for field production, in Egypt, received little or no attention. Where, some local cultivars, such as "Balady" cultivar, exhibited a relatively low productivity level and a clear deterioration in its quality characters, and showed, also, unusual degrees of variability in morphological characters among the individual plants of the grown population. Therefore, estimating and investigating the magnitudes of variability among lettuce plants for economic traits are very necessary to plan effective breeding programs. Since, the wide ranges of variability available provide good scope to conduct genetical studies and to design the identification of superior genotypes. In this respect,

several investigators estimated the magnitude of variability among lettuce genotypes for economic and important traits; such as Tash (2010) and Kumar *et al.* (2010). They concluded that wide range of variability estimates among the studied characters of different genotypes of lettuce were observed.

The heritability of characters determines much the phenotype of plant is a guideline genotype and thus help the breeder to base a selection program on the phenotypic performance of the plant. High heritability, in broad sense, indicated that a large proportion of phenotypic variance was attributed to the genotypic variance and were less influenced by environment. Falconer (1960) illustrated that characters which have high heritability are dependable because their gene expression is superimposed by the environmental influences. Thus, the degree of success in selection depends upon the magnitude of the heritability value; therefore, the effect of selection is realized more quickly in those characters which have high heritability. Many studies were reported in this respect (Gupta *et al.*, 2008; Souza *et al.*, 2009; Tashi *et al.*, 2010 on lettuce; and Antonova *et al.*, 2009; Soni *et al.*, 2013; Meena, 2014 on cabbage).

Interrelationships among various agronomic traits are vital to plan an effective breeding program. So, using phenotypic correlation is an important tool for the breeder to help in selecting and determine difficult measured characters through the selection of another easier in measuring. Therefore, some researchers estimated the relationships among the different pairs of the studied characters; such as Souza *et al.* (2008) and Kumar *et al.* (2010). These authors concluded generally that there were significant correlations between pairs of some economic characters of lettuce; but, with different trends (positive and/or negative); which were useful in lettuce selection. Also, positive and significant association of cabbage yield was observed with all the characters except days to maturity and stem length (Meena *et al.*, 2014). These results indicated that selection based on these characters either in combination or alone; will result in identifying the genotypes having high yield potential.

The main objective of the present investigation was to estimate and compare the efficiency of mass selection and individual plant selection, as two breeding methods, on the improvement of some important traits of lettuce cultivar "Balady". Heritability percentages, in broad sense, were also estimated for growth and productivity characters. The phenotypic correlation coefficients among the various pairs of the studied traits were also estimated to assist lettuce breeders in their selection programs.

MATERIALS AND METHODS

This study was carried out during the successive winter seasons of 2010/2011 till 2012/2013 at the Experimental Station Farm of the Faculty of Agriculture, Alexandria University; at Abies, Alexandria, A.R.E.

The genetical material used in this study was the "Balady" cultivar of lettuce crop which constitutes the original population for conducting the two methods of selection: individual-plant selection and mass selection. This cultivar was chosen since it is commonly grown and well adapted to the Egyptian environmental conditions; but, expresses a lot of variation and deterioration, which were observed and reported by many growers and consumers.

Growing the original population and data recorded

Seeds of the "Balady" cultivar were sown in nursery on October 15th, 2010, and the seedlings were transplanted on November 25th. The experimental area consisted of 80 rows, 4.00 m long and 0.60 m width. The spacing within rows was 0.20 m between plants. All the recommended agronomical and plant protection practices of lettuce commercial production were made, for raising a

healthy crop, whenever they appeared necessary.

Initial visual selection was made according to the criteria; maturity, non-wrapper leaves, core length (cm), core diameter (cm), head (cm), head diameter (cm), head weight (g), dry matter (%), bitterness (scale) and firmness (scale). Bitterness was given a score from 1 to 5, where score 1= non-bitter, 2= bitter and 3= better (Damarany, 1989b). Firmness of the head was ranked from 1 to 5, using the hand compression method (Kader *et al.*, 1973); where, 1= soft compressed or spongy; 2 = fairly firm, neither soft nor firm; 3 = firm, and commercially acceptable; 4 = hard and solid; and 5 = extra-hard. Each of the previously studied characters were measured on a plant basis, and they were used to calculate statistical parameters; range, mean, standard deviation and coefficients of variation.

Selection methods

In the growing season of 2010/2011 seedlings of "Balady" cultivar (original population C_0) were transplanted into the experimental area. Both selection methods were performed as follows:
Individual plant selection method: From 40 plants, selfed seeds of the 40 best plants with maturity and desirable agronomic traits were to be sown in the next season. On November 2011/2012 growing season, seedlings of each selected plant were cultivated in the five separated families. Then the practices of selection were between and within those families, on the basis of the best in earliness of maturity character, a head weight and its components; were conducted. At the end of the growing season selfed seeds of each selected plant from each selected family separately collected to get the final selected lines which came out to be five lines.

Mass selection method: seeds of the 10 best plants, from 1600 plants (C_0), were selected according to the performances of the previously mentioned desirable traits to represent the first cycle of mass selection (C_1). These seeds were bulked and sown in the nursery on October 15th, 2011. Seedlings were transplanted on November 25th. Then the best plants to represent the second cycle of selection (C_2).

Evaluation of the various genetic populations

On October 15th, 2012; the seeds of the original population (C_0), the five individual selected lines (S_{2-1} to S_{2-5}) and the second mass selected line (C_2), which were selected from the cultivar in this investigation, were sown and their seeds were transplanted on November 25th. The experimental design was a randomized complete blocks (RCBD) with three replicates. Each experimental unit consisted of three rows, 4.00 meters long and 0.60 meter width and the spacing was 0.20 meter, on one side of the r

commercial cultural practices were performed whenever they were necessary. The measurements of the ten studied characters were recorded on five randomly selected plants from each genetic population in each replication.

Statistical analyses

The statistical analyses for all collected data of the above-mentioned characters were conducted by the standard method of the randomized complete blocks design, as illustrated by Al- Rawi and Khalf-Alla (1980); using Co-Stat software (2004), a computer program for statistics of the differences between means.

Data of the various given characters were recorded to be used for genetic evaluation and estimation of some of the most important biometrical parameters; such as phenotypic correlation coefficients between different pairs of studied characters; which were estimated as described by Mather and Jinks (1971). Heritability percentages, in broad sense (h_{bs}^2), were also calculated as suggested by Allard (1960) as follows:

$$H_{bs}^2\% = \delta_g^2 / \delta_p^2 \times 100$$

$$\delta_p^2 = \delta_g^2 + \delta_e^2$$

$$\delta_g^2 = MS_g - (MS_e)/r$$

Where:

δ_p^2 = phenotypic variance

δ_g^2 = genotypic variance

δ_e^2 = environmental variance (error mean square)

RESULTS AND DISCUSSION

Variability estimates in the original population:

The estimated values of the parameters, mean, range, standard deviation and coefficient of variation for the studied important characters of the original population are arranged in Table 1. The results reflected clearly that the original population (C_0) was characterized by pronounced variability for

most studied characters as shown from the estimated coefficients of variation values. This result to be related to that, the cultivar Balady has been grown for a long period without any selection improvement. The characters that showed high coefficients of variability, high than 35% were bitterness (45.66%) and core length (38.66%). Whereas, the characters core diameter, firmness and non-wrapper leaves number and head weight showed relatively moderate levels of variability with estimated values of 32.48%, 31.22%, and 29.59% for their C.V.%, respectively. On the other hand, the four remaining characters: maturity, head diameter, leaves dry matter and head length; showed the lowest coefficients of variation values, that were estimated as 8.62%, 10.65% and 12.32%, each in turn. Generally, it can be stated that all studied characters could be improved through suitable selection method with varying degree depending upon the amount of variation presented in the population. The most of the studied characters of lettuce population have high chances to be improved; since they maintained relatively high variability in the original population. The obtained results were generally with those of Kumar *et al.* (2010), Tashi *et al.* (2010) on lettuce and Surlan-Mo *et al.* (1997) on cabbage; who found wide range of variation in most of the studied characters. In the same context Damarany (1989a) recorded such wide ranges of variation in most of the characters in cabbage and concluded that the characters could be improved through selection method.

Likewise, Solieman (1992) reported that Balady local cultivar of cabbage can be considered as a rich source of variation and can be used as main genetic material in breeding program to improve the characteristics of this crop.

Table 1: Estimates of variability parameters; range, mean \bar{X} , standard deviation and coefficient of variation (C.V. %), for the studied important characters in the original population of "Balady" cultivar.

Parameters	Rang		Mean \bar{X}	Standard deviation	Coefficient of variation (C.V.%)
	Maximum	Minimum			
Characters					
Maturity	74.00	54.00	65.04	5.61	8.62
Non-wrapper leaves No.	7.00	2.00	4.99	1.51	30.21
core length (cm)	10.00	2.00	6.09	2.36	38.66
core diameter (cm)	6.50	2.00	3.65	1.19	32.48
Head length (cm)	35.50	18.30	27.20	3.35	12.32
Head diameter (cm)	20.00	10.50	13.13	1.36	10.36
Head weight (g)	1020	137.2	520.46	153.99	29.59
Leaves dry mater (%)	7.85	5.18	6.39	0.68	10.65
Bitterness (scale)	3.00	1.00	1.73	0.79	45.66
Firmness (scale)	5.00	1.00	3.85	1.20	31.22

Evaluation the two selection methods:

Means of the different studied characters of the seven different populations; i.e., the original population (C_0), a population derived from the second cycle of mass selection (C_2) and the second selfed progenies of the five selected individual plants (S_{2-1} , S_{2-2} , S_{2-3} , S_{2-4} and S_{2-5}) are listed in Table 2. In all studied characters, the differences between the mean values of all selected populations and that of the original population appeared to be significant; but, with different magnitudes. The results, concerning the general performances of all studied characters, illustrated that the mean values of the five characters core diameter, head length, head diameter, head weight, and leaves dry matter were noticed to be increased (a desirable effect) after practicing the two cycles of mass selection or individual plant selection. But, in the case of the three characters core length, maturity and non-wrapper leaves number; their mean values were reduced (also, a desirable effect) after the two cycles of the selection methods. Concerning the remaining two characters i.e., bitterness and firmness; favorable performances, relative to the original population, were recorded for the two studied selection methods. These obtained results seemed to be compatible with those of Soliman (1992) on cabbage, who illustrated that all studied characters were improved through two practiced breeding systems; i.e., recurrent selection and individual plant selection; but, with different rates for the selected characters and the used breeding program. Also, Koutsos *et al.* (2001) recorded a desirable increment on the mean value of the yield character; which was estimated by 36%, without any undesirable changes in dry matter and soluble solids contents, as a result of applied three cycles of mass selection on cabbage. In this concern, there is a little published articles on emphasized selection of genotypes with lettuce breeding practices.

Estimates of heritability:

High heritability values in broad sense (Table, 3) were generally observed for all studied

characters; which ranged from 79.18% in diameter character to 99.66% in head diameter character. Similar findings were reported by *et al.* (2008) on lettuce; Shweta *et al.* (2011) Meena *et al.* (2014) on cabbage; who reported heritability estimates for their most characters. Generally, it may be stated that studied characters could be improved through selection, but with varying degrees, according to the amount of variation present in the population and the heritability of the concerned character. Therefore, most of the studied characters seem to have high chances to be improved through selection, since they appeared to maintain low variability in the original population characterized with high heritability percentages.

Estimates of phenotypic correlation coefficient

Concerning the phenotypic correlation coefficients, the obtained results (Table, 4) showed significant and desirable positive correlations between characters pairs head weight with core length and head diameter; and maturity with each of core length and bitterness. Also, significant positive correlations were detected between leaves dry matter with each of core diameter, head length and head diameter and head weight. On the other hand, significant negative correlations were detected between head weight with each of core length and bitterness, which are also desirable relationships. Results of correlation coefficients in the present study, appeared to be in a harmony with those of Rai and Asati (2005); Sharma (2005); Meena *et al.* (2010); Singh *et al.* (2011) on cabbage. From the previous results of correlation coefficients it may be concluded that some inter-relationships among various characters might be used to help in the selection of difficult to measure characters through selection of particular correlated ones that are easily measured and recognized.

Generally, it could be concluded that improving production of lettuce can be achieved through purification of already established cultivars and introducing improved strains through *in situ* mass and individual-plant selection programs.

Table 3: Estimates of phenotypic variance (δ_p^2), genotypic variance (δ_g^2) and broad sense heritability (H_{bs}^2 , %) for the ten studied characters of the evaluated genotypes of lettuce.

Parameters	H_{bs}^2 , %	δ_p^2	δ_g^2
Characters			
Maturity	94.09	62.15	58.48
Non-wrapper leaves No.	99.23	4.17	4.14
Core length (cm)	99.62	6.32	6.29
Core diameter (cm)	79.18	0.36	0.28
Head length (cm)	97.43	24.46	23.83
Head diameter (cm)	98.78	11.39	11.26
Head weight (g)	99.66	71364.29	71118.25
Leaves dry matter (%)	93.29	1.48	1.38
Bitterness (scale)	82.32	0.66	0.54
Firmness (scale)	92.62	1.34	1.24

1—Yield/vine:

Data in Table (1) clearly show that spraying clusters of Early sweet grapevines with GA₃ at 10 to 40 ppm or Sitofex at 2.5 to 10 ppm was significantly effective in improving the yield relative to the check treatment. The promotion on the yield was accompanied with increasing concentrations of each plant growth regulator. Using GA₃ at 10 to 40 was significantly preferable than using Sitofex at 2.5 to 10 ppm in improving the yield. A slight and insignificant promotion on the yield was attributed to increasing concentrations of GA₃ from 20 to 40 ppm and Sitofex from 5 to 10 ppm. The maximum yield was produced on the vines that received one spray of GA₃ at 40 ppm but the best treatment from economical point of view was the application of GA₃ at 20 ppm (since no measurable promotion on the yield was recorded between 20 and 40 ppm of GA₃). Under such promised treatment, yield/ vine reached 13.6 and 14.0 kg during both seasons, respectively. The control vines produced 9.1 and 9.6 kg during 2013 and 2014 seasons, respectively. The percentage of increase on the yield due to application of GA₃ at 20 ppm over the check treatment reached 49.5 and 45.8 % during both seasons, respectively. The beneficial effects of GA₃ on the yield might be attributed to their positive action on increasing cluster weight. The promoting effects of GA₃ on the yield was supported by the results of Dimovska *et al.*, (2011) and Abu Zahra and Salameh (2012) on different grapevine cvs.

The results regarding the beneficial effects of Sitofex on enhancing the yield are in harmony with those obtained by Juan *et al.* (2009); Abdel Fattah *et al.*, (2010) and Al Obeed (2011).

2—Harvesting date:

It is clear from the data in Table (1) that all GA₃ and Sitofex treatments had significantly delayed on the harvesting date of Early Sweet grapevines rather than the control treatment. The degree of delayness on harvesting date was correlated to the increase of the concentrations of both GA₃ and Sitofex. Using GA₃ significantly delayed harvesting date comparing with using Sitofex. Increasing concentrations of GA₃ from 20 to 40 ppm and Sitofex from 5 to 10 ppm failed to show significant delay on harvesting date. A considerable advancement on harvesting date was observed on untreated vines the great delay on harvesting date was observed on the vines that received GA₃ at 40 ppm during both seasons. GA₃ and Sitofex were shown by many authors to retard the release of ethylene and the disappearance of pigments such as chlorophylls and carotenoids and onset of maturity start. Also they were responsible for prolonging pre-maturity stages Nickell (1985). These results regarding the delaying effect of GA₃ and Sitofex on harvesting date were in harmony with those obtained by Wassel *et al.*, (2007), Kassem *et al.*

(2011), Abu Zahra and Salameh (2012) and *et al.* (2012).

3—Cluster weight and dimensions:

It is evident from the data in Table (1) treating clusters with GA₃ at 10 to 40 ppm or Sitofex at 2.5 to 10 ppm was significantly accompanied with enhancing weight, length and width of cluster relative to the control treatment.

The promotion was significantly associated with increasing concentrations of GA₃ and Sifofex. GA₃ was significantly favourable than using Sifofex in this respect. The maximum values were recorded on the vines that received one spray of GA₃ at 40 ppm. Meaningless promotion was detected with increasing concentrations of GA₃ from 20 to 40 ppm and Sifofex from 5 to 10 ppm. The untreated vines produced the minimum values during both seasons. The positive action of GA₃ on cluster weight and dimensions might be attributed to its essential role on stimulating cell division and enlarging cells, the water absorption and the biosynthesis of proteins which will lead to increase berry weight (Dimovska *et al.*, (2011); Abu Zahra and Sifofex (2012) and Dimovska *et al.*, (2014).

The previous essential role of CPPU on cluster weight was attributed to its higher concentration of cytokinin when applied to plants (Nickell, 1994).

4- Shot berries %:

Data in Table (2) obviously revealed that the percentage of shot berries in the clusters of Sweet grapevines was significantly controlled by spraying GA₃ at 10 to 40 ppm or Sifofex at 2.5 to 10 ppm relative to the check treatment. Using GA₃ was preferable than using Sifofex in reducing the percentages of shot berries. There was a reduction in the percentage of shot berries with increasing concentrations of GA₃ and Sifofex. A slight reduction in such an unfavourable phenomenon with increasing concentrations from 20 to 40 ppm and Sifofex from 5 to 10 ppm. The minimum values of shot berries (7.3 and 7.5%) during both seasons, respectively) were recorded in the clusters harvested from vines treated with GA₃ at 40 ppm. The maximum values of shot berries (12.0 & 12.5 %) during both seasons were recorded on the untreated vines during both seasons. The reducing effect of GA₃ on shot berries might be attributed to its important role on enhancing cell division and the biosynthesis of proteins (1985). These results were supported by the work of Wassel *et al.* (2007) and Abu Zahra and Sifofex (2012).

5- Fruit quality:

Data in Tables (2, 3 & 4) clearly showed that spraying clusters with GA₃ at 10 to 40 ppm or Sifofex at 2.5 to 10 ppm significantly accompanied with enhancing weight, long and equatorial diameter of berry, total acidity%, pro-

~~and percentages of P, K and Mg and T.S.S. %, reducing sugars %, T.S.S. / acid and total carotenoids relative to the check treatment. The effect either increase or decrease was associated with increasing concentrations of each auxin. Using GA₃ significantly changed these parameters than using Sitofex. A slight effect was recorded on these quality parameters with increasing concentrations of GA₃ from 20 to 40 ppm and Sitofex from 5 to 10 ppm. From economical point of view, the best results with regard to fruit quality were observed due to treating clusters with GA₃ at 20 ppm. Untreated vines produced unfavourable effects on fruit quality. These results were true during both seasons. The effect of GA₃ on increasing berry weight and dimensions might be attributed to its effect in promoting cell division and enlargement of cells, water uptake and the biosynthesis of proteins Nickell (1985). These results were in concordance with those obtained by Williams and Ayars (2005) and Dimovska *et al.*, (2014).~~

~~The higher content of Sitofex from cytokinins surly reflected on enhancing cell division and the elongation of berries Nickell (1985). These results were in agreement with those obtained by Abu Zahra (2013) and Retamales *et al.* (2015).~~

CONCLUSION

~~Treating Early Sweet grapevines once when the average berries reached 6mm with GA₃ at 20 ppm was responsible for promoting yield and fruit quality.~~

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الملخص العربي

الانتخاب الاجمالي وانتخاب النباتات الفردية كطريقتي تربية لتحسين نباتات الخس

انتصار ابراهيم مسعود راغب

قسم الخضر - كلية الزراعة - جامعة الإسكندرية

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~~قسم بحوث العنب - معهد بحوث البساتين - مركز البحوث الزراعية - الجيزة - مصر~~

ت تجارب حقلية وذلك بهدف تقدير حجم الأختلافات الموجوده؛ وكذلك تقدير كفاءة دورتين من الانتخاب ، وانتخاب النباتات الفردية لمدة جيلين علي بعض الصفات الهامة في الصنف البلدي لمحصول الخس، وذلك ة إلي تقدير كفاءة التوريث فى المعنى الواسع وكذلك معامل الارتباط المظهري بين أزواج الصفات المختلفة، وقد هذه الدراسة بمحطة البحوث الزراعية بأبيس التابعة لكلية الزراعة - جامعة الإسكندرية- خلال المواسم الشتوية لأعوام 2011/2010، 2012/2011، 2013/2012. وأظهرت النتائج بصفة عامة وجود اختلافات كبيرة تات وذلك فى الصفات المختلفة المدروسة في العشيرة الأصلية للخس(الصنف البلدى) ولكن بدرجات متفاوتة، كست أهمية البدء بأحد برامج التربية لتحسين الخس لانتاج تركيب وراثى أفضل فى صفاته مقارنة بالعشيرة . وأظهرت النتائج أيضا أن استخدام كل من طريقتي التربية المستخدمتين قد أدى إلي تحسين ملحوظ ومعنوي

لصفات المدروسة مقارنة بالعشيرة الأصلية، وذلك بدرجات مختلفة والتي تعتمد على برنامج التربية المستخدم صفات موضع الدراسة. وعكست الدراسة أيضا قيمة مرتفعة نسبيا لنسب كفاءة التوريث في المعنى الواسع المدروسة. ولقد أوضحت تقديرات معامل الارتباط المظهري بين الأزواج المختلفة للصفات المدروسة وجود معنوية وعالية المعنوية منها من يسلك الاتجاه الموجب ومنها من يسلك الاتجاه السالب (لكنها علاقات مرغوبة) ب أخذها في الاعتبار عند اجراء الانتخاب كأحد الوسائل (الادوات) الهامة التي يمكن أن يستفيد بها مربي في برامج الانتخاب. بناءا على ما سبق، فإنه يمكن تحسين انتاجية الخس من خلال تنقية الأصناف الموجودة ق تطبيق برنامج الانتخاب الاجمالي أو انتخاب النباتات الفردية للحصول على سلالات محسنة.

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