Estimation some genetic parameters of five tomato cultivars under the influence of Nanofertilizers addition

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ABSTRACT

for Breeding genetic structures adaptive for the country's atmosphere and to give a high yield, must the work on diffused varieties locally cultivation to determine the traits the most correlated with yield and The most inheritability and the affected by nano-Fertilization ,Which is one of the most important tools of modern agriculture to increase the productivity of tomatoes and reduce nutrient losses in rhizosphere area. However ,little is known about responding different genotypes of tomatoes nanofertilizers under field conditions in Babylon .Thus, using nanofertilizers may increase tomato growth productivity and decreasing the loss of nutrient in Sustainable agriculture .Here, we planted tomato cultivars (Red rock, Castel rock, Rock star, Early pearson, and Super queen) and sprayed them with three rates of nanofertilzers (0,1,2 g L-1) the factorial experiment was conducted in Complete Randomized Blocks Design (RCBD), The varieties differed among themselves in the characteristics of plant growth and under the effect of the three treatments, and the superiority of the variety early pearson on the other varieties in The qualities of vitamin C and under the influence of all the treatments and in the content of chlorophyll content on all varieties and under the influence of all treatments and recipe for the hight of the first inflorescence and acidity under the influence of the first treatment Recipe for percentage of potassium under the treatment of the first and second and The genetic variability values and the of expected genetic improvement were high in the percentage of nitrogen and potassium under the first treatment and Under the second treatment the genetic variability values and the of expected genetic improvement were high for the height of the first inflorescence, the percentage of potassium and phosphorus, low for the number of branches and the hardness and medium for the acidity and vitamin C and Under the influence of the third treatment, the genetic variance values and the of expected genetic improvement were high for the hight of the first inflorescence and vitamin C and the percentage of phosphorus.

Introduction

There is a need to increase the productivity of this crop through appropriate breeding programs that meet the requirements of local and export markets. Plant breeders have sought to produce Genetic variations . They are the basis for successful selection of the traits they are looking for ,Locally grown varieties possesses Genetically diverse Much more than modern or hybrid varieties , they are among the most important sources of genetic variation of breeders, and a large number of locally grown species and species

collected (Chen have been et al., 2009). Varieties vary in their growth and potential productivity and Which are controlled by the interaction of both genetic and environmental variability. This diversity can be attributed to adaptability of genes, to characteristics. morphological and physiological factors during crop growth. This diversity can be attributed to adaptability of genes, morphological characteristics, and physiological factors during crop growth (Olaniyi et al., 2010(.result increase in the

world's population led to increased demand for food and due to high food production prices; it necessary to use technology was in agriculture. Nanotechnology could be used as an alternative in a broad scientific field. Nanotechnology was described as materials, systems and processes that operate on a range of 100 nm or less(Ditta, 2012). Nanotechnology is a modern technology that has made a big breakthrough in various fields of science ,Progress Nanoparticles in agriculture For the purpose of minimizing nutrient loss to soil, water and air through direct internalization of crops and avoid

MATERIALS AND METHODS

A field experiment was carried out in Babylon province in March / 2018, where five cultivars (Red rock, Castel rock, Roke star, Early pearson, Super queen) were planted in glass dishes in the glass house and were transferred to the field on 20/3/2018. Plants were sprayed with nanofertilizer in 2018-4-2 with three batches separated by one month from one batch to another (0, 1,2 g L-1 macro and micro elements). All different service operations were carried out by irrigating, combing and plants fertilizing according to the recommended fertilizer by (200N + P 60 + 120K kg ha-1 (Faraj et al., 2012) Use the RCBD design and analyze the results by testing the least significant difference below the 5% probability level and the three replicates.

studied the traits: Number of branches ,chlorophyll content spad, Percentage of contract, height of the first inflorescence cm, potassium% and nitrogen% for all treatments, the number of days of planting until the flowers bloom 50% of plants experimental nutrient interactions with soil, microorganisms and air (Servin et al., 2015). This research aims to:-

*Evaluation of the performance of five local varieties of Tomato in the conditions of the province of Babylon.

*Estimation of some genetic parameters of these varieties under the influence of manure.

*Effect of nano-Fertilisers in the performance of these varieties.

*Interaction of varieties and manure in some growth characteristics of Tomato

unit, percentage of phosphorus, the percentage of acidity, vitamin C mg.100gm-1, Hardness cm2.

The genetic parameters were studied and included first ;the components of phenotypic variance were estimated to be based on the absence of an overlap between the genetic and environmental structure and the absence of a link between genetics and the environment according to the following equation: VP = VG+ VE

VG: Genotypic variance, variance Environmental VE: Variance Phenotypic: VP According to the genetic and environmental variance of the mean squares of the table for the analysis of variance for the design of complete random sectors, Table (1) according to the fixed model provided by (singh and ceccarelli, 1996(.

The analysis of genetic variance, phenotypic, environmental, and qualities Common was estimated according to the method explained by Walter (1975). VG=(msg-mse)/R

Average
squares
expected
Ve+g θ^2 R
Ve+r θ^2 G
Ve+ θ^2 GR

Table 1. ANOVA for complete randomized segments design with mean squares expected by constant model

Where R is the number of replicates = three, G is the number of Varieties = five.

where: MSe = VE, VG = (MS2-MS1)/R

M2: Average squares for varieties, M1: Mean squares of experimental error (VE = M1) R: number of replicates.

Determination of genotypic coefficients of variation and phenotypic coefficients of variation (panse and sukhatme, 1984) using the following The equation :

II. Estimation of inheritance in the broad sense: Broad Sense Heritability (H2 (b.s().

Falconer and Mackay (1996). H2(B.S)=VG/VP*100

The values of inheritance in the broad sense were expressed within the limits outlined by Ali (1999) within the following limits.

(Less than 40% low, between 40-60% medium, more than 60% high)

GCV%=(VG/ \tilde{Y}) ×100

PCV%=(VP/ \tilde{Y}) ×100

Note that \tilde{Y} is the arithmetic mean of the characteristic.

The values of the genetic and phenotypic coefficient were expressed using the scales used by Agarwal and Ahmed (1982) and in the following ranges

(Less than 10% low, between 10-30% medium, more than 30% high).

The percentage of contamination in the broad sense was estimated in the equation presented by

III. Estimation of Expected Genetic Improvement Expectant Genetic Advance (EGA:(

The expected genetic improvement as the percentage of the mean of the characteristic in the way explained (Kempthorne, 1969)

E.G.A% = [(K H2(b.s) VP)/ \tilde{Y}] ×100 where :

E.G.A: Genetic improvement expected as a percentage of the overall mean.

K: 2.06 The intensity of the selection for 5% of plants (Allard, 1960)

H2 (b.s): inheritance in the broad sense

 $\sqrt{VP}\,$: square root of phenotypic variation, $\tilde{Y}\colon$ average trait.

Results and discussion

Evaluation of the genotypes under the influence of the three factors flowering

The results in Table 2 indicate that the mean of the varieties of most studied traits under the influence of all the treatments were significant differences. Red rock was significantly superior to all varieties

In the vitamin C under the influence of all treatments and in the characteristics of early and percent of phosphorus under the influence of the third treatment and in the rate of hardness under the treatment of the first treatment. The superiority of the castel rock cultivar significantly in the rate of early flowering and the percent of phosphorus under The effect of the first and second treatment and the percent of the contract under the treatment of the first treatment. The superiority in the percent of nitrogen of the Rock star under the first treatment and the percent of potassium under the treatment of the third treatment. And the superiority of the category Early person in the content of chlorophyll on all varieties and under the influence of all treatment and the recipe for the hight of the first inflorescence and acidity under the influence of the first treatment and recipe percent of potassium under the treatment of the first and second. And the superiority of the phosphorus and low for the number of branches and hardness and medium for acidity and vitamin C The percentage of genetic improvement expected gave the highest rate of the hight of the first inflorescence and the percentage of potassium and phosphorus and the lowest rate of the number of branches and category Super queen on all classes in the number of branches under the influence of all treatment In the percentage of nitrogen under the influence of the second treatment and in the percent of the contract under the influence of the third treatment.

That the existence of moral differences between these varieties is necessary to continue to study the genetic behavior of these qualities to improve them (Hamdani, 2013.(

Genetic variability, inheritance ratio and expected genetic Advance The results of Table (3) indicate the general average and some genetic indicators of vegetative and syphilis growth characteristics and the qualitity characteristics of the tomato plant she was The percentage of inheritance ratio and the expected percentage of genetic improvement were different between the varieties. The inheritance rate was high for the characteristics of early flowering, chlorophyll content and the percentage of the contract under the influence of all the treatments and high in the percentage of nitrogen and potassium while it was low for acidity and vitamin C and medium for the number of branches and hardness and percentage of phosphorus The percentage of genetic improvement expected was high percentage of nitrogen and potassium and low for acidity and vitamin C for the first treatment. Under the influence of the second treatment, the inheritance ratio was high for the hight of the first inflorescence For the percentage of potassium and hardness, and under the impact of the third treatment was the inheritance rate high to the hight of the first inflorescence, Vitamin C, phosphorus percentage, low acidity, hardness, nitrogen percentage, average number of branches and percentage of potassium. The expected genetic improvement rate gave the highest rate of the hight of the first inflorescence and vitamin C and the percentage of phosphorus and the lowest rate of acidity and hardness and percentage of nitrogen.

Varie	Num	Chlor	High	Early	Fruit	Acidit	Hardness	Vit	%N	%P	%K
ties	ber of	ophyl	of	in	set	y%	kg.cm2	С			
	branc	l(spad	first	floweri	%	•		mg.1			
	hog)	inflor		70			111g.1			
	nes)	Inffor	ng				UUg			
			scenc					m ⁻¹			
			es cm								
	Under	the influ	ence of t	he first tr	ansactio	n	I		1		1
Red	8.33	37.9	11.00	42.33	28.17	0.907	2.10	12.6	1.369	0.380	1.369
rock								6			
Caste	8.00	33.7	11.67	39.3	36.31	0.763	1.51	9.09	1.577	0.423	1.577
l rock	0.00	447	15.00	10.67	24.00	0.770	1.57	10.5	2.072	0.000	2.072
KOCK star	8.00	44.7	15.00	43.67	24.89	0.770	1.57	10.5	2.073	0.330	2.073
Early	8.00	53.9	15.33	49.00	35.68	0.787	1.91	9.66	1.867	0.327	1.867
perso											
n											
Super	13.33	35.4	11.00	46.67	33.02	1.087	1.64	7.73	1.590	0.400	1.590
queen	0.015	10.01	0.554	4.207	7 400	NG	0.050	1.00	0.000	NG	0.000
L.S.D	3.315	10.21	3.556	4.307	5.499	N.S	0.953	4.09	0.299	N.S	0.2993
U.U5 Vorio	Undon	tha influ	onco of t	ha cacand	trancaa	tion		/	3		
v al le ties	Under	ule illiu		ne seconu	u ansac	uon					
Red	13.33	40.8	13.67	39.67	38.68	0.720	2.75	17.6	1.919	0.577	2.6400
rock								3			
Caste	14.33	51.5	12.67	35.00	36.08	1.170	2.74	15.9	2.127	0.777	3.1470
l rock								0			
Rock	12.00	35.8	16.33	39.33	29.13	0.933	2.07	16.1	1.827	0.613	3.880
star	11.67	560	10.00	47.00	20.27	1.077	1.00	2	2.077	0.502	4.220
Early	11.67	56.3	18.33	47.33	38.37	1.377	1.89	12.5	2.077	0.503	4.330
perso								0			
Super	16.67	39.6	12.67	43.00	22.62	1.373	2.91	11.5	2.223	0.480	3.673
queen								0		SN 2072-3	857
L.S.D	5.484	12.60	3.446	4.429	5.564	0.458	N.S	3.86	0.266	0.1248	0.4568
0.05						7/2		1	2		
Varie	Under	the influ	ence of t	he third t	ransactio	on –					
ties		1						1			
Red	13.33	38.9	11.67	33.00	31.89	1.133	1.90	23.9	2.243	0.823	2.96

rock								5			
Caste	11.67	38.4	18.33	35.67	36.13	0.670	1.97	11.6	2.033	0.607	4.01
l rock								7			
Rock	11.67	56.7	15.67	35.33	29.85	1.183	2.42	21.2	2.530	0.450	4.86
star								2			
Early	9.00	58.6	20.00	46.33	38.46	0.980	2.27	15.8	2.313	0.740	4.24
perso								0			
n											
Super	13.33	37.6	15.00	37.00	34.26	0.827	1.60	21.0	2.200	0.610	4.27
queen								8			
L.S.D	3.113	14.68	2.254	5.532	4.942	0.423	0.894	3.50	0.599	0.1467	1.061
0.05						7		1	6		

Table (2) The average studied traits of the tomato plant under the influence of the treatments Statistical and genetic analyzes

The high values of inheritance ratios of some traits give an indication of the high values of genetic variability and low environmental variability values, The traits are genetically controlled and The environment has slightly effect on them (Welsh, 1981). The values of genetic variation were high for the percentage of nitrogen and potassium, while low for acidity, vitamin C, medium for the number of hardness and branches, percentage of phosphorus. The values of environmental variability were high for acidity and vitamin C and low for the percentage of nitrogen and potassium and medium for the number of branches, Percent of the phosphorus for the first treatment. Under the second treatment, the genetic variability values were high for the height of the first bulb and the percentage of potassium and phosphorus and low for the number of branches and the hardness and medium for the acidity and vitamin C While she was the values of the environmental variability were high for the number of branches and the hardness and low for the hight of the first inflorescence and the

percentage of potassium and phosphorus and the medium of the acidity and vitamin C, but under the treatment of the third was the values of genetic variance high for the hight of the first inflorescence and vitamin C and the percentage of phosphorus and low for the acidity and hardness and percentage of nitrogen And the mean of the number of branches and the percentage of potassium. The values of environmental variation were high acidity, hardness and percentage of nitrogen and low for the hight of the first inflorescence and vitamin C and the percentage of phosphorus and medium for the number of And potassium percentage. These results are consistent with Hasan et al., 2016 (Kumar et al., 2013), Shashikanth et al., 2010 (Kaushik et al., 2011) (Kumar and Bhardwaj, 2014.(The high variance coefficient of some

characteristics means that dispersion exists between the studied traits. Thus, the selection of such qualities is easy and the program is effective for their improvement (Singh and Chaudhary, 1985.(

Varieties	Nu	Chlor	Hig	Early	Frui	acidit	Hardnes	Vit C	N%	P%	K%
	mbe	ophyl	h of	in	t set	у	s cm.k ⁻ g ⁻				
	r of	1	first	floweri			1				
	bra	(spad	inflo	ng							
	nch)	rsce								
			nces								

Table (3) The general average, variance phenotypical, genetic, environmental, inheritance and expected genetic improvement of the studied yields of the tomato plant under the influence of the treatments

Under the influence of the first transaction												
General	9.13	41.1	12.8	44.20	31.6	0.863	1.75	9.93	1.69	0.372	0.5071	
mean			0		1				5			
Genotypic	4.5	59.10	3.56	12.4	21.5	0.005	0.0231	1.767	0.06	0.003	0.4490	
variance		3	6		51	76			74	9		
variance	3.10	29.38	3.56	5.233	8.52	0.074	0.2563	4.734	0.02	0.004	0.0725	
Environmen	0		7		9	54			528	302		
tal												
Variance	7.6	88.48	7.13	17.633	30.0	0.080	0.2794	6.5	0.09	0.008	0.525	
Phenotypic		11.00	3	- 0 (8	3	1.00	4 . 4 .	26	2	122.12	
genotypic	23.2	14.38	14.7	7.96	14.6	27.80	1.32	17.16	15.1	16.78	132.13	
coefficients	3		5		8				3			
of variation	20.1	22.00	20.0	0.50	17.2	22.02	20.20	25 (7	17.0	24.24	242.40	
phenotypic	30.1 0	22.00	20.8 6	9.50	17.5	32.83	30.20	23.07	17.9	24.34	242.40	
of variation	9		U		5				U			
Broad Sonso	50.2	66 76	10 0	70.32	70.3	7 17	8 26	27.18	727	17 56	86.00	
Heritability	3 <i>7.2</i>	00.70	9	10.32	2	/.1/	0.20	27.10	2.	77.50	00.07	
Expectant	36.8	31 48	21.4	13 762	137	4 851	5 1 3 9	14 37	26.9	23 34	39.84	
Genetic	2	51.40	8	15.702	6	4.001	5.157	14.57	057	20.04	57.04	
Advance	-		Ŭ		Ū				007			
Under the infl	uence	of the se	cond tr	ansaction				1				
General	13.6	44.8	14.7	40.87	32.9	1.115	2.47	14.75	2.03	0.590	3.214	
mean	0		3		8		-		4			
Genotypic	1.25	60.64	5.13	19.3	45.4	0.062	0.234	5.288	0.01	0.012	0.4108	
variance					2	0			89	33		
variance	8.48	44.77	3.35	5.533	8.73	0.059	1.330	4.204	0.01	0.004	0.05886	
Environmen	3		0		2	37			999	393		
tal												
Variance	9.73	105.4	8.53	24.833	54.1	0.121	1.564	9.492	0.03	0.016	0.4697	
Phenotypic	3	1	3		52	3			889	723		
genotypic	8.22	17.38	15.3	10.74	20.4	22.33	19.58	15.59	6.75	18.82	18.13	
coefficients			8		3							
of variation	22.0	22.01	10.0	10.10		21.22	=0. (2	20.00	0.60	0 1.01	10.20	
phenotypic	22.9	22.91	19.8	12.19	22.3	31.23	50.63	20.88	9.69	21.91	19.39	
coefficients	3		3		1							
Of variation	12.0	57.50	60.1	77 71	02.0	51 11	14.06	55 71	10 5	72 72	07 16	
Horitability	12.0	57.54	00.1 5	//./1	03.0 7	51.11	14.90	55./1	40.5	15.15	0/.40	
Fynectant	-	27.15	3 24 5	10 51	38.5	32.88	15.60	23.07	9 70	33.20	3/ 03	
Genetic	0.00 75	27.13 4	24.3 7	17.51	5	52.00	13.00	23.91	9.70	55.29	54.75	
Advance	15	-	,		5							
Under the infl	uence	of the th	ird trai	nsaction								
General	11.8	46.0	16.1	37.47	34.	0.959	2.03	18.70	2.26	0.646	4.07	
mean	0		3		12				4			
Genotypic	2.23	92.51	9.83	23.76	9.2	0.028	0.0276	23.91	0.00	0.018	0.3772	
variance					3	6			106	36		

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variance	2.73	60.76	1.43	8.633	6.8	0.050	0.2256	3.458	0.10	0.006	0.3178
Environmen	3		3		90	65			14	072	
tal											
Variance	4.96	153.2	11.2	32.393	16.	0.079	0.2532	27.36	0.10	0.024	0.695
Phenotypic	3	7	63		12	25		8	24	43	
genotypic	12.6	20.90	19.4	13.008	8.9	17.63	8.18	26.14	1.43	20.97	15.09
coefficients	5		3		0						
of variation											
phenotypic	18.8	26.91	20.8	15.18	11.	29.35	24.78	27.97	14.1	24.19	20.48
coefficients	7		0		76				3		
of variation											
Broad Sense	44.9	60.35	87.2	73.34	57.	36.08	10.90	87.36	1.03	75.15	54.27
Heritability	3		7		25						
Expectant	17.4	33.45	37.4	22.94	13.	21.81	5.565	50.34	0.29	37.45	22.89
Genetic	7		0		87				9		
Advance											

Conclusion

Different varieties among them under the influence of all varieties and superiority of the variety Early person in most of the studied traits.

The values of genetic improvement were high for most studied traits and under the influence

REFRENCES

Agarwal, V. and Z. Ahmad .1982. Heritability and genetic advance in triticale . Indian J. Agric. Res. 16: 19-23.

AL-Hamdani, Sh. Y.H.2013. genetic variability, correlation and expectant genetic advance for yield and its combonents in melon (Cucumis melo L.) .Tikrit Journal of Agricultural Sciences.13(2): :227-236.

Ali, Abdo Al-Kamil Abdullah.1999. Hybrid force and gene action in yellow maize. PhD thesis, Faculty of Agriculture and Forestry, University of Mosul.

Allard , R.W. 1960.Principles of Plant Breeding.John willey and Sons.Inc. New York,USA. of all studied traits. The values of genetic improvement were high in the characteristics of early flowering, percentage of the fruit set and the chlorophyll content in the leaves under the influence of all the treatments.

AL-mfargy,O.K.A.,F.M.M. AL-mfargy.2015. Effect of fertilizer type on quantitate and qualitative yield characters of tomato (Lycopersecun esculantum Mill.) and some estimate genetic parameters . Dep. of Hort. and landscaping - College of Agri.- University of Diyala - Republic of Iraq.

Chen, J., H. Wang, H.L. Shen, M. Chai, J.S. Li, M.F. Qi and W.C Yang. 2009. Genetic variation in tomato populations from four breeding programs revealed by single nucleotide polymorphism and simple sequence repeat markers. Sci. Hortic., 122(1): 6-16.

Ditta A., 2012. How helpful is nanotechnology in agriculture? Adv. Nat. Sci.: Nanosci. Nanotechnol, 3(3):1-10. Falconer, D.C.,

and

T.F.C.Mackay.1996.Introduction to quantitative genetic(4th edition).John Wiley and Sons.New York.

Faraj, A. H., Z. H. Mohammud., A. J. Gaber., H. S.Ibrihium.2012.Respond Plant Tomato Lycopersicon esculentum Application of Foliar Cycocel and NPK in Desert Soil. Kufa Journal of Agricultural Sciences,4(2):137-156. Hasan, M.d. Mehedi., A. M.d. Al-Bari and M. A. Hossain. 2016. Genetic Variability and Traits Association Analysis of Tomato (Lycopersicon esculentum L.) Genotypes for Yield and Quality Attributes .Universal Journal of Plant Science 4(3): 23-34.

Kaushik, S. K., D. S. Tomar. and A. K. Dixit. 2011. Genetics of fruit yield and it's contributing characters in tomato (Solanum lycopersicom).Journal of Agricultural Biotechnology and Sustainable Development 3(10): 209 -213.

Kempthorne , B. 1969. An Introduction to Genetic Statistics. Ames Iows State Univ. press.

Kumar,N., and M.L. Bhardwaj.2014 .studies on genetic variability in tomato (Solanum lycopersicum L.)genotypes for different horticultural traits. submitted in partial fulfilment of the requirements for the degree of master of science (horticulture) ,Nauni Solan - 173230 (HP), INDIA.

Kumar, V., R. Nandan., S.K. Sharma., K. Srivastava., R. Kumar and M.K. Singh .2013. Heterosis study for quality attributing traits in different crosses in tomato (Solanum Lycopersicum L.). Plant Archives, 13(1): 21-26.

Olaniyi, J. O., W. B. Akanbi, T. A. Adejumo and O. G. Akande.2010. "Growth, fruit yield and nutritional quality of tomato varieties," African Journal of Food Science, 4(6): 398– 3402.

Panse , V.G. and P.V. Sukhatme .1984. Statistical Methods for Agricultural Workers. ICAR, New Delhi, India.

Servin ,A., W. Elmer , A. Mukherjee , R. D. Torre-Roche, H. Hamdi .2015.A review of the use of engineered nanomaterials to suppress plant disease and enhance crop yield, Jason C. White . Prem Bindraban . Christian Dimkpa 17(92):1-22.

Shashikanth, B. N., B. C. Patil., P. M. Salimath., R. M. Hosamani . and P.U. Krishnaraj. 2010. Genetic divergence in tomato (Solanum lysopersicom Mill.) Wettsd.).Karnataka Journal of Agricultural Sciences, 23(3): 538-539.

Singh, M. and S. Ceccarelli .1996.Estimation of Heritability of Crop Traits.From Variety Trial Data. Technical Manual No. 21.

Singh, R.K. and B.D. Chaudhary .1985. Biometrical Methods in Quantitative Genetic Analysis. Kalyani Publishers. New Delhi., India.

Walter, A.B. 1975. Manual of Quantitative Genetics [3 rd edition], Washington State Univ. Press. U.S.A.

Welsh, J.R. 1981. Fundamentals of Plant Genetics and Breeding.John Wiley &Sons, Inc. New York U.S.A.