

# Hybrid Vigor And Combining Ability Of Faba Bean Genotypes And Their Crossbreeds As Affected By Arginine Spraying

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## Abstract

*This study was conducted during the 2022/2023 and 2023/2024 agricultural seasons at the Zafaraniya Research Station of the Horticulture Department. The aim was to evaluate the genetic diversity of 13 broad bean cultivars and assess the performance of their individual hybrids under the influence of arginine spraying at two concentrations (0 and 250 mg/L<sup>-1</sup>). The first season included molecular analysis using SSR and RAPD techniques. Based on these results, six cultivars were selected that showed clear genetic divergence and low tannin and vicine contents. The selected cultivars were introduced into a crossbreeding program to produce 15 first-generation (F<sub>1</sub>) hybrids, which were studied with their parents in the second season. The experiment was implemented in a split-plot design within a randomized complete block design with three replicates. Data were statistically analyzed using SAS software and Duncan's multiple range test at the 0.05 probability level. Genetic components were estimated using the Griffing (1956) method (second method) within the fixed model. The results showed significant variation in general and specific combining ability, with parents 4 and 5 showing the best positive general combining ability under the arginine treatment, while parent 6 recorded the lowest general combining ability for most traits. Several hybrids, including 5x4, 6x5, and 2x1, showed significant specific combining ability. Dominant genetic variances exceeded additive genetic variance for most traits under treatments A<sub>0</sub> and A<sub>1</sub>, indicating the importance of dominance gene action. Genotypic values also exceeded environmental variance for most traits, indicating the presence of a genetic basis that can be exploited in faba bean genetic improvement programs.*

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## THE INTRODUCTION

Broad beans (*Vicia faba* L.) are a strategic legume crop, characterized by their rich nutritional content, including protein (22–38.2%), fiber (~12%), carbohydrates (57.3%), vitamins, minerals, and important medicinal and environmental properties (Labba et al., 2021; Jayakodi et al., 2023). They belong to the legume family (Fabaceae), and their cross-pollination rates range between 8% and 84%, with an average of 30% to 60% depending on environmental conditions and genetic variations among cultivars (Suso et al., 2001; Suso & Maalouf, 2010). Broad beans are the fourth most important legume crop after peas, chickpeas, and lentils, and the seventh most produced globally (Kaur et al., 2014; Oliveira et al., 2016). Estimating heritability is a crucial step in predicting the potential for genetic progress through selection programs. High values of additive action reflect promising opportunities for genetic improvement, while low values indicate the dominance of non-additive action (Narayanan, 1993). Diallel crosses are an effective tool for analyzing genetic variance, as they allow for the estimation of general combining ability (GCA) associated with additive genetic action and specific combining ability (SCA) associated with non-additive genetic action. Several studies, including Alghamdi (2009) and Al-Jubouri (2014), have demonstrated significant GCA and SCA squares for traits such as plant height, number of branches, flowering date, and pod filling duration, reflecting clear genetic differences between the studied populations. Al-Shakirji (2011) also noted that several hybrids exhibited high combining abilities for important quantitative traits. In this context, other studies have confirmed the presence of wide genetic variation in mineral content (Baloch et al., 2014, 2017), and high heritability ratios for traits such as pod number, 100-seed weight, and grain yield (Bakhiet et al., 2015; Sheelamary and Shivani, 2015; Sharifi, 2015; Hamza et al., 2017; Mesfin et al., 2021).

A high coefficient of genetic variation indicates a low environmental influence on phenotypic expression, enhancing the efficiency of genetic selection (Ejigu et al., 2016), while a high phenotypic variance relative to genetic variation indicates the dominance of environmental factors and the difficulty of genetic improvement (Mesfin, 2019). Therefore, combining genetic variance analysis with heritability and genetic progression estimates is essential for effective decision-making in broad bean improvement programs

## MATERIALS AND WORKING METHODS

Seeds of selected broad bean varieties were sown on October 1, 2023, at the Zafaraniya Research Station, affiliated with the Horticulture Department / Spring Palm Station, to study the genetic and productive performance of genotypes under the influence of arginine spraying. A factorial experiment was conducted according to a randomized complete block design (RCBD) and a split plot design with three replicates. It included two factors: the first was the genotypes, which included six parents and fifteen single hybrids resulting from cross-breeding. The second factor was foliar spraying with arginine at two levels (no spraying and spraying at a concentration of 250 mg/L<sup>1</sup>). Spray treatments were placed in the main plots, and genotypes in the secondary plots. Seeds were sown in lines parallel to drip irrigation pipes, with a distance of 0.75 m between lines, 1 m between replicates, and 0.2 m between plants. Each replicate contained 42 genotypes (21 genotypes × two spray treatments), resulting in a total of 126 experimental units. All agricultural maintenance operations, including irrigation, weeding, hoeing, and pest control, were carried out as needed. Based on the results of genetic fingerprinting using the SSR and RAPD techniques, six cultivars were selected from among 13 cultivars that showed clear genetic divergence. These cultivars were Histal, Claro ED luna, Sakiz Bakala, Broad Beans, Kawadlji, and Iraqi Kurtana, and were designated (V1-V6), respectively. These cultivars were entered into a half-crossing program according to the first method and the fixed model of Griffing's (1956) methods, to produce first-generation hybrids (F<sub>1</sub>). Subsequent measurements and data related to morphological, physiological, and production traits were conducted to analyze variance and estimate genetic traits

Characteristics of the product and its components

The study included the evaluation of several traits related to yield and its components, which were measured according to the following methods.

Number of pods per plant (pod/plant<sup>-1</sup>): The number of pods was counted in five randomly selected plants from each experimental unit, and the average was calculated

Average pod weight (g pod<sup>-1</sup>): Green pods were collected from the same five plants used for counting and weighed to extract the average pod weight

Number of seeds per pod (seed/pod<sup>-1</sup>): The seeds were counted in ten randomly selected pods from the plants, and the average was calculated

Weight of 100 seeds (g): Several samples of 100 whole seeds were taken and weighed to extract the overall average in grams

Green pod yield (g plant<sup>-1</sup>): was calculated by multiplying the average number of pods by the average pod weight for each individual plant

Total yield (tons ha<sup>-1</sup>): was calculated using the equation The following The total yield of the experimental unit in hectares was calculated using the following equation: Yield of the experimental unit (g) × Area of the experimental unit (m<sup>2</sup>) × 1000/10000

Table No.

(1) shows the names and origins of the fava bean varieties

Origin	Item name	T
Spain	Luz DE otono	1
Spain	Histal	2
Spain	Claro ED luna	3
Iraqi	Local variety (Kartania)	4
Netherlands	Broad bean Quadrilogy	5
New Zealand	AGuadulce	6
California	Basic	7
Turkey	Bakla Tohumu	8
Turkey	Bakla Sevilla	9
Turkey	Sakiz Bakala	10
Moroccan	Monarch	11
Netherlands	Quads SN23	12
America	Broad Beans	13

## 3-RESULTS AND DISCUSSION

1-3-Hybrid vigor based on the deviation of the first generation from the best parents at the two levels of arginine spraying.

Average weight of the pod

The study showed significant differences in heterosis for pod weight between parents and their hybrids under the influence of arginine spraying. Six hybrids achieved significant positive heterosis, with the (4x5) hybrid having the highest percentage, exceeding 7% in the unsprayed treatment and 6.5% with spraying, indicating its genetic superiority in this trait. In contrast, some hybrids recorded negative heterosis, such as the (1x5) hybrid, which reached -31%, reflecting poor genetic performance. These results reflect the importance of genetic compatibility between parents in improving pod weight, and the effect of arginine spraying on enhancing heterosis for some combinations

Table 2: Hybrid vigor and pod weight calculated on the basis of the deviation of the first generation from the best parents under the effect of spraying with arginine.

Hybrid	Arginine spray treatments		Hybrid power of coefficient averages
	A2	A1	
-0.526**	-0.501*	-0.554**	1x2
-19.185**	-18.748**	-19.642**	1x3
0.889**	0.850**	0.931**	1x4
-29.576**	-28.173**	-31.125**	1x5
-0.421	-0.401	-0.443*	1x6
-29.080**	-28.263**	-29.951**	2x3
-10.769**	-10.301**	-11.281**	2x4
-4.387**	-4.176**	-4.621**	2x5
-3.250**	-3.086**	-3.432**	2x6
6.233**	5.694**	6.838**	3x4
-1.227**	-1.479**	-0.933**	3x5
2.707**	2.303**	3.165**	3x6
6.754**	6.461**	7.076**	4x5
2.740**	2.621**	2.870**	4x6
1.177**	1.120**	1.239**	5x6
0.287	0.293	0.282	

### Number of horns

The study showed significant differences in the heterosis for the number of pods between parents and their hybrids under the effect of arginine spraying. Six hybrids achieved significant positive heterosis in the unsprayed treatment (A1), with the highest positive effect reaching 11.69% in the hybrid (2x3). In the sprayed treatment (A2), five hybrids achieved positive heterosis, the highest of which was 6.99% in the hybrid (4x6). The average of the two treatments also showed positive heterosis in five hybrids, with the best percentage being 6.62% for the hybrid (4x6). However, some hybrids recorded negative heterosis, such as the hybrid (2x3), which reached -12.97%, indicating poor genetic performance of these combinations

Table 3: Hybrid vigor. Number of horns calculated on the basis of the deviation of the first generation from the best parents under the influence of spraying with arginine

Hybrid	Arginine spray treatments		Hybrid power of coefficient averages
	A2	A1	
-4.858**	-4.545**	A1	1x2
5.062**	4.158**	-5.216**	1x3
3.749**	3.406**	6.036**	1x4
-6.959**	-6.497**	4.106**	1x5
-0.735**	-0.684**	-7.489**	1x6
-1.486**	-12.967**	-0.793*	2x3
-7.430**	-6.951**	11.690**	2x4
-10.260**	-9.598**	-7.977**	2x5
-7.430**	-6.951**	-11.015**	2x6
2.974**	1.509**	-7.977**	3x4
-2.831**	-2.643**	2.086**	3x5
-0.275	-0.455	-3.046**	3x6
2.624**	2.698**	-0.297	4x5
6.619**	6.992**	2.507**	4x6

-2.418\*\*      -2.257\*\*      6.185\*\*      5×6  
0.386   0.306   -2.602\*\*

### Number of seeds in a pod

The results showed significant differences in the heterosis for number of seeds per pod between the parents and their hybrids. Eight hybrids achieved positive heterosis in the unsprayed treatment (A1), with the highest positive effect reaching 15.37% in the hybrid (5×6). In the sprayed treatment (A2), eight hybrids also achieved positive heterosis, with the highest reaching 12.95% in the same hybrid (5×6). The average of the two treatments also showed positive heterosis in the eight hybrids, with the best percentage reaching 14.05% in the hybrid (5×6). In contrast, some hybrids recorded negative heterosis, with the lowest value reaching -19.88% in the hybrid (1×3), indicating differences in the genetic performance of these combinations

Table 4: Hybrid vigor. Number of seeds in pods calculated on the basis

of the deviation of the first generation from the best parents under the effect of spraying with arginine

Hybrid Arginine spray treatments Hybrid power of coefficient averages

A2	A1		
3.486**	3.246**	3.765**	1×2
-18.736**	-17.713**	-19.884**	1×3
2.275**	2.132**	2.439**	1×4
-1.534**	-1.428**	-1.656**	1×5
-12.831**	-11.948**	-13.855**	1×6
-12.309**	-11.637**	-13.063**	2×3
-3.539**	-3.317**	-3.794**	2×4
0.646**	0.595**	0.706**	2×5
2.990**	2.721**	3.319**	2×6
2.614**	2.471**	2.774**	3×4
-5.991**	-5.664**	-6.358**	3×5
-5.337**	-5.046**	-5.664**	3×6
6.321**	5.924**	6.775**	4×5
2.275**	2.132**	2.439**	4×6
14.054**	12.946**	15.371**	5×6
0.343	0.313	0.374	

### Weight of 100 seeds

The differences between parents and their hybrids showed positive and negative heterosis in 100-seed weight, depending on the best parent. In the treatment without arginine spray (A1), 15 hybrids achieved significant positive heterosis, with the highest positive effect reaching 30.60% in the hybrid (1×4). Similarly, in the treatment with arginine spray (A2), the same 15 hybrids achieved positive heterosis, with the highest percentage reaching 30.44% in the hybrid (1×4). As for the average effect of the two treatments, 15 hybrids also showed positive heterosis, with the best reaching 28.34% in the hybrid (1×4). Table5:Hybrid vigor, weight of 100 seeds, calculated on the basis of the deviation of the first generation from the best parents under the effect of spraying with arginine

Hybrid Arginine spray treatments Hybrid power of coefficient averages

A2	A1		
27.619**	26.030**	29.240**	1×2
14.386**	15.756**	13.021**	1×3
28.338**	30.444**	30.597**	1×4
16.574**	22.783**	20.888**	1×5
12.780**	15.066**	10.464**	1×6
20.564**	22.847**	18.271**	2×3
23.688**	24.521**	22.844**	2×4
7.766**	13.936**	11.335**	2×5
22.297**	21.901**	22.703**	2×6
16.415**	18.246**	14.589**	3×4

9.595**	9.588**	9.602**	3×5
19.218**	20.260**	18.168**	3×6
12.329**	13.669**	21.148**	4×5
16.948**	16.507**	17.402**	4×6
7.027**	13.993**	9.595**	5×6
0.534	0.209	0.860	

### Yield of one plant

The differences between parents and their hybrids on plant yield resulted in varying effects of heterosis, both positive and negative, compared to the best parents. In the treatment without arginine spray (A1), eight hybrids showed positive effects on heterosis, but only two of them were significant, with the hybrid (4×5) recording the highest positive percentage of 14.606%. In the treatment with arginine spray (A2), nine hybrids showed positive heterosis, of which six were significant, with the highest being 15.661% in the same hybrid (4×5). The average effect of the two treatments was that eight hybrids showed positive heterosis, with the best reaching 15.191%. In contrast, the remaining hybrids showed negative heterosis, with the lowest being -35.203% in the hybrid (1×5)

**Table 6: Hybrid vigor, yield per plant, calculated on the basis of the deviation of the first generation from the best parents under the effect of spraying with arginine.**

Hybrid Arginine spray treatments				Hybrid power of coefficient averages
A2	A1			
4.378	4.075*	4.768	1×2	
-15.093**		-15.390**		1×3
4.720	4.375**	5.150	1×4	
-33.318**		-28.540**		1×5
1.675	1.554	1.830	1×6	
-24.565**		-33.532**		2×3
-10.309**		-9.096**		2×4
-11.421**		0.750	-12.212**	2×5
-3.945	-3.824	-4.195	2×6	
9.531**		7.269**		3×4
1.004	-0.744	3.343	3×5	
3.713	1.871	6.174	3×6	
15.191**		15.661**		4×5
10.886**		12.058**		4×6
-1.265	6.975**		-1.394	5×6
3.613	2.412	4.814		

### Total Yield

The study showed that differences between parents and their hybrids significantly affected the total plant yield, with both positive and negative heterosis appearing compared to the best parents. In the first treatment (A1), only one hybrid achieved a significant positive heterosis of 14.606% in the (1×3) hybrid. In the second treatment (A2), two hybrids achieved significant positive heterosis, the highest of which was 37.829% in the same hybrid (1×3). Averaged across the two treatments, two hybrids achieved a maximum positive heterosis of 32.578% in the (1×3) hybrid. In contrast, the other hybrids showed a negative heterosis of -47.232% in the (2×5) hybrid. These results reflect a clear genetic variation among the genotypes, suggesting the potential for improving faba bean productivity through the selection of hybrids with positive heterosis

**Table 7: Hybrid vigor of the total plant, calculated on the basis of the deviation of the first generation from the best parents under the effect of spraying with arginine**

Hybrid Arginine spray treatments				Hybrid power of coefficient averages
A2	A1			
-10.837**		-10.819**		1×2

32.578**	37.829**	14.831**	1×3
-25.934**	-25.757**	-26.164**	1×4
-17.927**	-17.425**	-18.561**	1×5
-6.913**	-6.598**	-7.341**	1×6
11.598**	12.048**	-2.052**	2×3
-22.443**	-22.504**	-22.357**	2×4
-45.278**	-43.719**	-47.232**	2×5
-2.590**	-2.426**	-2.800**	2×6
-33.570**	-32.507**	-34.894**	3×4
-27.302**	-26.337**	-28.507**	3×5
-10.591**	-10.100**	-22.097**	3×6
-7.783**	-7.519**	-8.111**	4×5
-5.307**	-5.079**	-5.589**	4×6
-3.734**	-3.109**	-4.520**	5×6
1.173	1.007	1.340	

### Effect of General Combining Ability

Table 8 Estimation of the effect of general damage susceptibility of parents on the studied traits at the level of arginine spraying of 0 and 250 mg L<sup>-1</sup> for fava beans.

SE(ĝi)	parents Transactions						
0.096	6	5	4	3	2	1	Weight Qurna
0.032	5.030*	-1.557*	-0.596*	-0.058	-1.726*	-1.091*	0
0.106	-0.729*	-1.583*	2.641*	7.005*	-4.692*	-2.640*	250
0.070	-0.316*	-0.149*	0.145*	0.213*	0.693*	-0.586*	0
0.094	0.073*	0.192*	0.300*	-0.218*	-0.071*	-0.277*	250
0.056	-0.902*	-0.371*	0.590*	1.127*	-0.539*	0.095*	0
0.196	-0.845*	-0.314*	0.646*	1.184*	-0.653*	-0.018	250
0.048	-2.004*	-2.389*	1.775*	2.994*	0.175	-0.551*	0
100							
seeds							
1.902	-1.337*	-3.908*	0.536*	3.574*	0.090*	1.043*	250
0.953	-118.4*	68.068*	32.964*	25.414*	53.569*	-10.877*	0
0.305	-104.129*		42.657*	58.335*	26.237*	26.770*	-49.872*
0.230							
	-1.542*	0.515*	12.273*	-0.898*	-6.791*	-3.557*	0
Total result							
0.096	-0.779*	1.645*	14.499*	-5.082*	-7.231*	-3.050*	250

The results of the study showed that the six parents showed different general abilities for the studied traits under the two treatments (A0: no spray and A1: arginine spray). Parent 1 had a positive and desirable general ability for the number of seeds per pod and 100-seed weight traits under treatment A1. Parent 2 recorded a positive and significant desirable general ability in yield per plant under both treatments. It also showed positive performance in number of pods and 100-seed weight under treatment A1, but showed negative general ability in the remaining traits. Parent 3 achieved a positive and significant desirable general ability in number of seeds per pod, 100-seed weight, and yield per plant under both treatments, in addition to a positive ability in pod weight under treatment A1, but showed negative general ability in the remaining traits. Parent 4 showed significant and positive general potency in most of the studied traits, including pod number, number of seeds per pod, 100-seed weight, yield per plant, and total yield under both treatments, with a positive potency in pod weight under treatment A1. Parent 5 outperformed in significant and positive general potency in yield per plant and total yield under both treatments, in addition to a positive performance in pod number under treatment A1, but recorded negative general potency in the remaining traits. Parent 6 showed a positive and desirable general potency in pod number only, with negative potency in the remaining traits.

High values of general potency for each parent indicate effective transfer of genetic traits to hybrid progeny, while low values indicate difficulty in transferring desired traits. This is consistent with genetic

principles in hybrid breeding and is consistent with the results of previous studies such as (Soliman et al., 2023) and (Ghannam et al., 2024)

### Effect of Specific Combining Ability

Table 9 Estimation of the effect of the specific ability on combining for individual hybrids for the studied traits at the level of arginine spraying of 0 and 250 mg L<sup>-1</sup>

studied characteristics			Transactions					
Total result		Yield of one plant		Weight				
100								
seeds	Number of seeds in pods			Number of horns			Weight	
Qurna	Arginine spray level							
1.758*	-0.059	0.741*	14.856*	97.886*	-2.957*	0	1×2	
4.697*	0.759*	0.839*	11.286*	108.507*	-5.161*	250		
0.131*	0.623*	-0.885*	4.597*	-154.14*	21.503*	0	1×3	
-3.280*	1.775*	-0.958 *	4.350*	-132.989*	33.499*	250		
1.299*	1.041*	0.282*	15.087*	10.650*	-7.993*	0	1×4	
1.562*	0.206*	0.208*	18.864*	-4.313*	-11.520*	250		
1.230*	-0.957*	0.213*	13.805*	44.164*	-1.566*	0	1×5	
-9.442*	-1.245*		0.140	14.335*	54.029*	-3.138*	250	
-6.167*	0.050	-0.065	-2.109*	196.85*	-5.942*	0	1×6	
0.773*	-0.286*		-0.138	1.235*	136.865*	-8.013*	250	
1.356*	4.649*	0.339*	11.473*	134.163*	10.582*	0	2×3	
-6.308*	-2.456*		0.266*	15.585*	149.152*	18.445*	250	
1.474*	-1.691*	0.457*	9.743*	-327.95*	-0.643	0	2×4	
-1.235*	-0.718*		0.383*	11.673*	-299.386*	-2.984*	250	
1.035*	-2.387*	0.018	-0.158	69.348*	-24.591*	0	2×5	
1.609*	-1.600*		-0.054	3.018*	23.141*	-29.227*	250	
-6.272*	-1.230*		-0.170	13.986*	114.548*	0.988*	0	2×6
-0.454*	-0.491*		-0.243*	11.677*	98.032*	0.484	250	
1.020*	-0.372*	0.003	4.537 *	-41.782*	-20.100*	0	3×4	
4.486*	0.267*	0.032	8.470*	-52.588*	-18.543*	250		
1.191*	-0.358*		0.174	1.485*	118.548*	-13.334*	0	3×5
4.881*	0.095	0.203*	0.365*	133.448*	-11.374*	250		
-5.336*	-0.600*	0.765*	13.503*	110.018*	-11.461*	0	3×6	
6.047*	-0.195		0.794*	13.264*	118.076*	-9.274*	250	
1.509*	1.460*	0.492*	11.831*	244.37*	7.476*	0	4×5	
2.945*	1.516*	0.521*	2.205*	226.911*	8.223*	250		
-5.398*	1.428*		0.703*	5.099*	138.508*	12.258*	0	4×6
0.421*	1.825*	0.732*	3.738*	146.866*	13.918*	250		
-5.467*	0.312*	0.634*	-0.385	89.834*	9.283*	0	5×6	
-0.273*	-0.056	0.663*	4.700*	123.025*	11.064*	250		
0.265		0.293	0.259	0.539	5.226	0.840	0	SE(sij)
0.090		0.192	0.155	0.132	2.619	0.631	250	

The results of the Specific Combining Ability (SCA) analysis showed significant positive and negative variances in most of the studied hybrids for the different productive traits under the influence of the two treatments (A0 without spraying and A1 with arginine spraying), reflecting the presence of non-additive genetic interaction between parents in the formation of hybrids. For pod weight, the hybrid (1×5) recorded the highest positive value of 6.047 under treatment A1, while the hybrid (1×3) recorded the lowest negative value of -9.442. As for the number of pods, the hybrid (4×6) achieved the highest value of 1.825 under A1, while the lowest was -2.456 in the hybrid (2×3). No significant differences were found in some hybrids, such as (3×5), (3×6), and (5×6) under A1, and (1×6) under A0. For number of seeds per pod, hybrid (3×6) had the highest positive value of 0.794 under A1, while hybrid (1×3) had the lowest negative and significant value of -0.958. No significant differences were found in several hybrids including (1×6), (2×5), and (3×4) in both treatments. In 100-seed weight, hybrid (1×4) had the highest positive value of 18.864 under A1, while hybrid (1×6) recorded the lowest value of -2.109 under A0. Likewise, the

differences were not significant for the (2×5) and (5×6) hybrids at A0. As for the per-plant yield, all hybrids showed significant differences, with the (5×4) hybrid recording the highest positive value of 244.37 at A0, while the lowest value was -327.95 in the (2×4) hybrid at A0. In the total plant yield, the highest positive value was for the (1×3) hybrid at A1, reaching 33.499, and the lowest negative value was -29.227 in the (2×5) hybrid under the same treatment. No significant differences were recorded for the (2×4) hybrids at A0 and (2×6) at A1. These results indicate that hybrids that showed significant and positive specific aptitude, especially in the desired direction, can contribute to improving production traits by increasing available genetic variation. Such high values are often associated with the presence of parents with high general aptitude. These results are of practical importance in breeding programs, in line with the findings of Heiba et al. (2023) and Jasim et al. (2023)

## REFERENCES

1. Alghamdi, S. S. (2009). Heterosis and combining ability in diallel cross of eight faba bean (*Vicia faba* L.) genotypes. *Asian Journal of Crop Science*, 1(2), 66
2. AlJubouri, Hatim Mohammed Hassan. (2014). Effect of row spacing on yield and its components of some faba bean (*Vicia faba* L.) cultivars. M.Sc. Thesis, College of Agriculture, University of Kirkuk
3. Al-Shukri, Weam Yahya Rasheed. (2011). General and specific combining ability and diallel cross analysis for yield and its components in F2 hybrids of faba bean. *Al-Rafidain Agriculture Journal*, 3(2), 41
4. Bakhiet MA, Rania AR, El-Said Raslan MA, Abdalla NG, 2015. Genetic variability, heritability and correlation in some faba bean genotypes under different sowing dates. *World Appl Sci J*, 33: 1315-1324
5. Bakhiet MA, Rania AR, El-Said Raslan MA, Abdalla NG, 2015. Genetic variability, heritability and correlation in some faba bean genotypes under different sowing dates. *World Appl Sci J*, 33: 1315-1324
6. Baloch FS, Karaköy T, Demirbaş A, Toklu F, Özkan H, 2014. Variation of some seed mineral contents in open pollinated faba bean (*Vicia faba* L.) landraces from Turkey. *Turk J Agric For*, 38: 591-602.
7. Baloch K, Rizwan S, Mahmood K, Jan MH, Hussain J, 2017. Biochemical and trace elements composition of faba bean (*Vicia faba* L.) cultivated in Panjgur and Kech districts of Balochistan. *Pure Appl Biol*, 6: 981-988.
8. Ejigu E, Wassu M, Berhanu A, 2016. Genetic variability, heritability and expected genetic advance of yield and yield related traits in common bean genotypes (*Phaseolus vulgaris* L.) at Abaya and Yabello, Southern Ethiopia. *Afr J Biotechnol*, 17(31): 973-980
9. Georgieva N, Nikolova I, Kosev V, 2016. Evaluation of genetic divergence and heritability in pea (*Pisum sativum* L.). *J Biosci Biotechnol*, 5: 61-67.
10. Ghannam, H. A., RA, I., Ibrahim, M. A., Mousa, M. I., & Soliman, A. A. (2024). Genetic Analysis for Some Faba Bean Agronomic Traits. *Scientific Journal of Agricultural Sciences*, 6(4), 72-91.
11. Heiba, H., Mahgoub, E., Mahmoud, A., Ibrahim, M., & Mansour, E. (2023). Combining ability and gene action controlling chocolate spot resistance and yield traits in faba bean (*Vicia faba* L.). *Journal of Agricultural Sciences*, 29(1), 77-88.
12. Jayakodi, M., Golicz, A.A., Kreplak, J., et al., 2023. The giant diploid faba genome unlocks variation in a global protein crop. *Nature* 615, 652–659.
13. Jasim, E. A. A., Esho, K. B., & Salim, N. S. (2023). Study the Genetic Performance of Some Faba Bean Genotypes Under Mosul Condition, Iraq. *Bionatura*, 1(8), 1-12.
14. Kaur, S., Cogan, N. O., Forster, J. W., & Paull, J. G. (2014). Assessment of genetic diversity in faba bean based on single nucleotide polymorphism. *Diversity*, 6(1), 88-101.
15. Labba, I.C.M., Frøkiær, H., Sandberg, A.S., 2021. Nutritional and antinutritional composition of fava bean (*Vicia faba* L., var. minor) cultivars. *Food Res. Int.* 140, 110038.
16. Mesfin T, 2019. Breeding achievements of faba bean (*Vicia faba* L.) and its impact in the livelihood of Ethiopian farmers. *Int J Agric Biosci*, 8(5): 263-269.
17. Mesfin T, Wassu M, Mussa J, 2021. Variation in genetic variability and heritability of agronomic traits in Faba bean (*Vicia faba* L.) genotypes under soil acidity stress evaluated with and without lime in Ethiopia. *Afr J Agric Res*, 17(2): 355-364.
18. Narayanan, S. S. (1993). Biometrical techniques in plant breeding. Kalyani Georgieva N, Nikolova I, Kosev V, 2016. Evaluation of genetic divergence and heritability in pea (*Pisum sativum* L.). *J Biosci Biotechnol*, 5: 61-67.
19. Oliveira, H. R., Tomás, D., Silva, M., Lopes, S., Viegas, W., & Veloso, M. M. (2016). Genetic diversity and population structure in *Vicia faba* L. landraces and wild related species assessed by nuclear SSRs. *PloS one*, 11(5), e0154801
20. Sharifi P, 2015. Genetic variation for seed yield and some of agro morphological traits in faba bean (*Vicia faba* L.) genotypes. *Acta Agric Slov*, 105(1): 73-83.
21. Sharma PP, Vyas M, Meghwal DR, 2017. Estimation of genetic variability and correlation analysis in field pea (*Pisum sativum* L.) genotypes. *J Plant Dev Sci*, 9: 53-56.
22. Sheelamary S, Shivani, 2015. Genetic variability, heritability and correlation of faba bean (*Vicia faba* L.) grown in New Delhi. *Int J Adv Technol Eng Sci*, 3: 48-51.
23. Soliman, A. A., Ibrahim, M. A., Mousa, M. I., Mansour, E., He, Y., & Yu, H. (2024). Genetic potential and inheritance pattern of agronomic traits in faba bean under free and infested *Orobanche* soil conditions. *BMC Plant Biology*, 24(1), 301.
24. Suso, M. J., & Maalouf, F. (2010). Direct and correlated responses to upward and downward selection for outcrossing in *Vicia faba*. *Field Crops Research*, 116(1-2), 116-126.
25. Suso, M. J., Pierre, J., Moreno, M. T., Esnault, R., & Le Guen, J. (2001). Variation in outcrossing levels in faba bean cultivars: role of ecological factors. *The Journal of Agricultural Science*, 136(4), 399-405.