

## Variability and character association in fennel (*Foeniculum vulgare* Mill.) germplasm

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### Abstract

Extent of variability was assessed in 50 accessions of fennel using augmented design. The analysis of variance indicated that the accessions differed significantly for most of the characters. The genotypic coefficient of variation was high for umbels plant<sup>-1</sup> (16.96%) followed by seed yield plant<sup>-1</sup> (13.13%), plant height up to primary umbel (11.08%) and length of first internodes (10.71%) indicating that these characters have high to moderate amount of genetic variability. The estimates of heritability (broad sense) was high for plant height up to primary umbel (78.66%), plant height up to top of the plant (72.61%), length of first internodes (64.07%) and days to 50% flowering (51.24%). The genetic advance as percentage of mean were high for umbels plant<sup>-1</sup> (22.75), plant height up to primary umbel (20.24), length of first internodes (17.66) and plant height up to top of the plant (16.43). On the basis of yield as well as other morphological characters the accessions namely JF-406, JF-533-2, NDF-45 and GK/RM/AK-7 were found superior as compared to checks.

**Key words :** Correlation, *foeniculum vulgare*, genetic advance, heritability, variability.

### Introduction

Spices occupy an important place in the lives of peoples since Vedic and Biblical times. They have been considered requisite in seasoning of foods, flavouring of beverages, perfumeries, cosmetics and medicines. The test and flavour of spices prompted explorers like Columbus and Vasco de Gamma to undertake dangerous sea journey to discover India "the land of spices" (Agrawal *et al.*, 2001).

Fennel (*Foeniculum vulgare* Mill.) belonging to family Apiaceae. It is a cross pollinated crop and diploid species. It is having chromosome number,  $2n = 22$  and is native of Europe and Meditarrean region. Fennel is cultivated throughout the temperate and sub-tropical regions in the world mainly in the countries like, Romania, Russia, Hungary, Germany, France, Italy, India, Sri Lanka, Malaysia, Japan, Argentina and U.S.A. In India, it is mainly grown in the states of Gujarat, Rajasthan, Haryana and some extent in West Bengal and Uttar Pradesh. Total area under the crop in India is about 74660 hectares with production of 124610 tonnes. In Rajasthan, it occupies an area of 30720 hectares with an annual production of 35500 tonnes (Spice Board India, 2016-17).

Steam distillation of dried fruit yields, an essential oil referred as 'Fennel oil', used for flavouring purposes in the many parts of the world (Abdallah *et al.*, 1978;

Lawrence, 1992). Fennel and its preparations are used to cure various disorders, and also act as a carminative, digestive and diuretic agent. Fennel increases elasticity of connective tissues and act as anti-aging agent (Arslan *et al.*, 1989). There are some commercial pharmaceuticals with formula based on fennel essential oil. The antimicrobial properties of the essential oil have been also recognized (Elagayyer *et al.*, 2001; Ruberto *et al.*, 2000). The essential oil from plants has usually been isolated by either steam distillation or solvent extraction (Sajid *et al.*, 2008). Information on extent of variation, estimates of heritability and expected genetic advance irrespective of yield and yield determining traits constitutes the basic requirement for a crop improvement programme that is lacking. Keeping in view above facts, the present investigation was aimed to evaluate variability, heritability and genetic advance of grain yield and its component characters fennel genotypes and correlation with yield contributed characters. Therefore this study was conducted and use of these traits for further breeding programme for improvement in fennel crop.

### Materials and methods

The present study was carried out in *Rabi* season of 2012-13 at the Research Farm of National Research Centre on Seed Spices, Ajmer (26° 27' 0" N, 74° 38' -1" E and 460 meter above sea level altitude). The region falls under

agro climatic zone III of Rajasthan. The soil of research farm is sandy loam, poor in fertility and water holding capacity, having pH 8 to 8.3, EC 0.07 to 0.12 and 0.15 to 0.23% organic carbon, available N 178.5 kg ha<sup>-1</sup> (low), P<sub>2</sub>O<sub>5</sub> 12 kg ha<sup>-1</sup> (medium), K<sub>2</sub>O 85 kg ha<sup>-1</sup> (low), Ca 214.7 kg ha<sup>-1</sup> (high), Mg 258 kg ha<sup>-1</sup> (medium), S 27 kg ha<sup>-1</sup> (high). The study was carried out with 50 diverse genotypes from different geographic and genetic origin with five checks in Augmented Block Design. Plot size was kept 6 × 2, row to row spaced 50 cm apart. Plant spacing within rows was maintained 20 cm. The recommended package of practices was adopted for raising healthy crop. Five plants were randomly selected from each line and observations were recorded on plant height, primary branches, umbel plant<sup>1</sup>, umbellate umbel<sup>1</sup>, seeds umbellate<sup>-1</sup>, test weight and yield (g). Data were analyzed through Indostate version 8.5 software.

## **Results and discussion**

In the present investigation, augmented block design was used to evaluate the accession along checks. This design is ideal when the number of lines to be evaluated is more and seed supply is less. The analysis of variance revealed that significant amount of variability was present in the germplasm for all the characters studied (Table-1), this suggests that the material used in the study had adequate variability and response to selection may be expected in breeding programme for seed yield plant<sup>1</sup>. An appreciable portion of this variability was contributed by genotypic variance which is in agreement with earlier reports of Ramanujam and Joshi (1966), Bhargava *et al.*, (1971), Shukla *et al.*, (2003), Lal (2007), Telci *et al.*, (2009), Meena *et al.*, (2009) and Malik *et al.*, (2009). The block (eliminating check + varieties) effects were also significant for duration of complete anthesis, 1000-seed weight (g), expected other characters indicating the sensitivity of genotypes to the environment. These observations are similar to that observed by Singh *et al.*, (2004). The check varieties showed significant difference for most the characters. It indicated that checks themselves were diverse. Estimate of genotypic and phenotypic coefficient variance (GCV & PCV) indicated that in general phenotypic variance were higher than genotypic coefficient variances indicating the role of environmental factors on the characters expression. The variances of characters were compared on basis of coefficient of variation. The range and coefficient of variation indicated that the variability was high for seed yield plant<sup>1</sup>, umbel plant<sup>1</sup>, moderate for plant height up to primary umbel, 1000-seed weight (g), length of first internodes (cm), it was low for plant height up to top of the plant, seeds umbellates<sup>-1</sup>, umbel plant<sup>1</sup>,

days to 50% flowering duration of complete anthesis, primary branches plant<sup>1</sup>, secondary branches plant<sup>1</sup> (Table-2).

It indicated that simple selection for seed yield plant<sup>1</sup>, umbel plant<sup>1</sup>, umbellets umbel<sup>1</sup> and secondary branches plant<sup>1</sup>, might be advantageous as compared to other characters under study. Similar pattern of variability for different characters among accession have earlier been reported by Bhargava *et al.*, (1971), Kathiria (1980), Agnihotry *et al.*, (1997), Shukla *et al.*, (2003), Rajput *et al.*, (2004), Meena *et al.*, (2009) and Singh *et al.*, (2009). With the help of PCV and GCV alone it is not possible to determine the amount of variation which is heritable. In the present investigation, broad sense heritability was observed to be high for plant height up to primary umbel (cm), plant height up to top of the plant (cm), length of first internodes (cm), days to 50% flowering and similar results were also obtained by Agnihotry *et al.*, (1997), Jain *et al.*, (2002), Lal (2007), Singh and Choudhary (2008), Meena *et al.*, (2010) and Yogi *et al.*, (2013). Moderate heritability (30-60 per cent) was observed for secondary branches plant<sup>1</sup>, duration of complete anthesis, diameter of primary umbel (cm), umbels plant<sup>1</sup>, primary branches plant<sup>1</sup>. Which is in agreement with earlier report of Kathiria (1980), Bhandari and Gupta (1991) in coriander. Low heritability was observed seed yield plant<sup>1</sup>, 1000-seed weight, umbellets umbel<sup>1</sup>, seeds umbellets<sup>-1</sup>.

Genetic advance as percentage of mean for the characters ranged from 2.43 % (days to 50% flowering) to 22.75 % (umbels plant<sup>1</sup>). High magnitude of genetic advance as percentage of mean was estimated for umbels plant<sup>1</sup>, plant height up to primary umbel (cm), length of first internodes, which is in agreement with earlier reports of Mathur *et al.*, (1971) in coriander, Agnihotri (1990), Agnihotri *et al.*, (1997) and Meena *et al.* (2010). Moderate genetic advance as percentage of mean was observed for plant height up to top of the plant (cm), secondary branches plant<sup>1</sup>, seed yield plant<sup>1</sup> (g), diameter of primary umbel, umbellets umbel<sup>1</sup>, which is in agreement with earlier reports of Ramanujam *et al.*, (1964), Kathiria (1980), Bhandari and Gupta (1991) and Singh *et al.*, (2004). Low genetic advance as percentage of mean was observed for 1000-seed weight (g), seeds umbellets<sup>-1</sup>, primary umbel plant<sup>1</sup>, duration of complete anthesis, days to 50% flowering. These results are in accordance with the early reports of Jain and Dubey (1972) in coriander, Bhandari and Gupta (1991).

The plant height up to primary umbel (cm), length of first internodes (cm), plant height up to top of the plant (cm) had higher magnitude of heritability and genetic advance as percentage of mean.

**Table 1.** Analysis of variance for different characters

Sources of Variation	Df	days to 50% flowering	length of first internodes	primary branches plant <sup>-1</sup>	diameter of primary umbel	duration of complete anthesis	secondary branches plant <sup>-1</sup>	umbels plant <sup>-1</sup>	umbellets umbel <sup>-1</sup>	seeds umbellets <sup>-1</sup>	plant height up to primary umbel (cm)	plant height up to top of the plant (cm)	1000-seed weight (g)	Seed yield per plant (g)
Block (ignoring Treatments)	4	38.553**	2.704**	0.926	14.658**	24.540**	9.514	59.596	18.191	12.481*	784.195**	1134.654**	1.025	6.071
Treatment (eliminating Blocks)	54	5.415	1.086**	0.524	2.653	0.612	9.055*	79.695	14.231	6.687	187.249**	172.638**	0.641	84.482**
Checks	4	18.040**	2.420**	0.726	0.582	0.742	20.539**	38.698	11.880	10.606	211.184**	180.493*	1.024	388.425**
Checks + Var vs. Error	50	4.405	0.979**	0.508	2.818	0.602	8.136	82.974	14.419	6.373	185.334**	172.009**	0.611	60.167**
Error	16	2.890	0.308	0.334	2.037	1.149	3.847	41.131	10.414	3.813	35.273	52.286	0.383	17.701
Block (eliminating check + Var)	4	4.740	0.328	0.246	5.628	6.678**	4.691	49.044	4.696	9.014	98.886	80.336	1.481*	3.231
Entries (ignoring blocks)	54	7.950	1.217**	0.575	3.321	1.920	9.412*	80.476	15.230	6.944	238.012**	250.698**	0.608	84.692**
Checks	4	18.040**	2.420**	0.726	0.582	0.742	20.539**	38.698	11.880	10.606	211.184**	180.493*	1.024	388.425**
Varieties	49	7.228*	1.091**	0.567	3.608	2.056	8.256*	84.431	15.814	5.754	221.042**	250.320**	0.548	21.726
Checks vs. Varieties	1	2.940	4.205**	0.336	0.231	0.007	21.584*	53.820	0.054	50.576**	1176.840**	550.084**	1.855*	195.5104**
Error	16	2.890	0.308	0.334	2.037	1.149	3.847	41.131	10.414	3.813	35.273	52.286	0.383	17.701

**Table 2.** Overall mean value of genotypes, their range, genotypic and phenotypic coefficient of variation, heritability in broad sense, and genetic advance as percent of mean for different characters in fennel.

Characters	Mean	Range	Genotypic coefficient of variation(GCV) %	Phenotypic coefficient of variation (PCV)	Heritability in broad sense (%)	Genetic Advance % of mean
Days to 50% flowering	105.58	100.96-110.16	1.65	2.30	51.24	2.43
Length of first inter nodes	6.91	4.70 - 9.55	10.71	13.38	64.07	17.66
primary branches per plant	7.410	6.03 – 8.99	5.45	9.51	32.89	6.44
Diameter of primary umbel (cm)	14.68	10.22 – 19.43	7.14	12.05	35.07	8.71
Duration of complete anthesis	13.61	10.80 -15.40	5.84	9.80	35.58	7.18
secondary branches per plant	17.95	9.16 – 25.18	9.78	14.66	44.52	13.44
Umbels per plant	32.45	13.64 – 63.74	16.96	26.04	42.43	22.75
Umbellets per umbel	22.51	15.86 – 36.34	8.63	16.73	26.63	9.17
Seeds per umbellets	23.66	18.05 – 28.97	4.92	9.60	26.28	5.20
Plant height up to primary umbel (cm)	102.90	74.92 – 137.10	11.08	12.49	78.66	20.24
Plant height up to top of the plant (cm)	125.25	101.11 – 162.55	9.36	10.98	72.61	16.43
1000-seed weight (g)	5.86	4.02 – 8.97	5.80	12.03	23.26	5.76
Seed yield per plant (g)	12.77	5.16 – 32.23	13.13	35.44	13.73	10.02

**Table 3.** Correlation coefficient on the basis of genotypic level and phenotypic level between different characters of fennel

Characters	PI/G	Length of first inter nodes	primary branches plant <sup>-1</sup>	Diameter of primary umbel	Duration of complete anthesis	secondary branches plant <sup>-1</sup>	Umbels plant <sup>-1</sup>	Umbellets umbel <sup>-1</sup>	Fruits umbellets <sup>-1</sup>	Plant height up to primary umbel (cm)	Plant height up to top of the plant (cm)	1 000- seed weight (g)	Seed yield plant <sup>-1</sup> (g)
Days to 50% flowering	G	-0.358**	0.055	0.0755	0.2936*	0.338**	-0.175	-0.17	0.1825	0.1843	0.294'	0.0324	-0.053
Length of first inter nodes	P	-0.44**	0.1346	0.2814*	0.5982**	0.358**	-0.11	-0.0591	0.04'	0.317*	0.44**	0.018	-0.041
	G	1	-0.031	-0.372**	-0.186	-0.22	-0.075	0.0608	-0.24**	0.1249	-0.054	-0.092	-0.151
primary branches per plant	P	1	-0.0737	-0.384**	-0.3231*	-0.246*	-0.044	-0.0299	-0.064	-0.103	-0.236*	-0.155	-0.189
	G	1	1	0.0635	0.0527	-0.03	0.333**	0.31'	0.0637	0.0777	0.217	-0.191	0.0867
Diameter of primary umbel	P	1	1	0.1604	0.0862	0.018	0.2824*	0.3655**	0.093	0.177	0.255'	-0.129	0.048
	G	1	1	1	0.2817*	0.173	0.267*	0.0248	0.0894	0.144	0.344**	0.0394	0.173
Duration of complete anthesis	P	1	1	1	0.135	0.127	0.1257	0.1542	-0.008	0.308*	0.429**	-0.058	0.173
	G	1	1	1	1	0.599**	0.251*	0.0523	0.0835	0.4492**	0.525**	0.1018	-0.083
secondary branches per plant	P	1	1	1	1	0.375**	0.0234	-0.0163	-0.159	0.233*	0.347**	0.009	-0.09
	G	1	1	1	1	1	0.338**	0.0638	-0.081	0.4458**	0.484**	0.2195	0.0358
Umbels per plant	P	1	1	1	1	0.3364**	0.1636	0.1636	-0.2	0.386**	0.414**	0.122	0.125
	G	1	1	1	1	1	1	0.1508	-0.37**	0.0422	0.129	-0.111	0.1985
	P	1	1	1	1	1	1	0.1683	-0.48**	0.053	0.043	-0.102	0.179

Characters	P/G	length of first inter nodes	primary branches plant <sup>-1</sup>	diameter of primary umbel	duration of complete anthesis	secondary branches plant <sup>-1</sup>	umbels plant <sup>-1</sup>	umbellets umbel <sup>-1</sup>	fruits umbellets <sup>-1</sup>	plant height up to primary umbel (cm)	plant height up to top (cm)	1000-seed weight (g)	seed yield plant <sup>-1</sup> (g)
Umbellets per umbel	G						1	0.0985	0.2931*	0.275*	0.275*	-0.01	0.0098
seeds per umbellets	P						1	0.185	0.382**	0.366**	0.366**	-0.076	-0.015
	G						1	1	0.172	0.237*	0.237*	-0.038	-0.137
	P												
Plant height up to primary umbel (cm)	G								1	0.229	0.183	-0.039	-0.106
	P								1	0.862**	0.862**	-0.006	0.336**
Plant height up to top of the plant (cm)	G								1	0.907**	0.907**	-0.245*	-0.327*
1000-seed weight (g)	P											-0.119	-0.257*
	G											-0.174	-0.261*
Seed yield per plant (g)	P											1	0.0874
	G											1	0.223
	P											1	1
	P											1	1

Genotypic and phenotypic correlations were worked out among different characters including seed yield. The phenotypic correlation coefficients were generally higher than the respectively genotypic correlation coefficients. Phenotypic correlation coefficient provides a measure of genetic association between characters and thus, help in identifying the traits which need to be considered for improvement of yield. Since, suitable test for significance of phenotypic correlations is not available so their main usefulness is in strengthening the interpretations based on genotypic correlations.

At genotypic level, seed yield per plant had significant negative correlation with plant height up to primary umbel (cm), plant height up to top of the plant (cm), where, it had non significant negative correlation with days to 50% flowering, length of first internodes, duration of complete anthesis, umbellets umbel<sup>-1</sup>, seeds umbellets<sup>-1</sup> and non significant positive correlation with 1000-seed weight (g), umbel plant<sup>-1</sup>, secondary branches plant<sup>-1</sup>, diameter of primary branches per plant (Table-3).

Plant height up to primary umbel (cm) had positive correlation plant height up to top of the plant (cm), plant height up to primary umbel (cm) with significant negative correlation, seed yield plant<sup>-1</sup>, 1000-seed weight are in agreement with earlier reports of Jindla and Rang (1986), Agnihotry *et al.*, (1997) and Idhol *et al.*, (2009).

Primary branches plant<sup>-1</sup> had significant positive correlation umbel plant<sup>-1</sup>, umbellets umbel<sup>-1</sup>, plant height up to top of the plant (cm), are in agreement with earlier reports of Kathiria (1980) and Singh *et al.* (2004). Days to 50% flowering had significant positive correlation with secondary branches plant<sup>-1</sup>, plant height up to primary branches plant<sup>-1</sup> (cm), are in agreement with earlier reports of Ali *et al.*, (1993), Godara (1995) and Agnihotri *et al.* (1997).

Genotypic level of significant and positive correlation with secondary branches plant<sup>-1</sup> with umbels plant<sup>-1</sup>, plant height up to primary umbel (cm), plant height up to top of the plant which earlier reports of Singh *et al.* (2009). The association of seeds per umbel with plant height up to top of the plant had significant and positive.

It is also noticed that character which exhibited negative association with seed yield plants<sup>-1</sup> have also exhibited negative association among themselves. Thus, these characters could be simultaneously improved to increase the seed yield. On the basis of yield as well as other morphological characters the accessions namely JF-406, JF-533-2, NDF-45 and GK/RM/AK-7 were found superior as compared to checks.

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