

# Estimation of Genetic Variability, Heritability and Genetic Advance for Essential Oil Yield and Related Traits in Genus *Ocimum*

#### Smita<sup>\*</sup>S<sup>1,2</sup> and Kishori RL<sup>2</sup>

<sup>1</sup>Academy of Scientific and Annovation Reasearch (AcSIR), CSIR-Central Institute of Medicinal and Aromatic Plants, Lucknow, India

<sup>2</sup>Department of Genetics and Plant Breeding, India

\*Corresponding author: Smita S, Academy of Scientific and Annovation Reasearch (AcSIR), CSIR-Central Institute of Medicinal and Aromatic Plants, Lucknow, India, Tel: 9450629069; E-mail: cimap.smita@gmail.com

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## Abstract

Assessing variability is fundamental to identify the most important traits in *Ocimum* improvement program. The objective of the present study was to estimate variability, heritability and genetic advance based on twelve morphological characters of *Ocimum*. The experiment was conducted in 2015/16 main cropping season at one location by using randomized complete block design with three replications. The results revealed highly significant differences (P<0.01) among genotypes for all characters considered. The phenotypic coefficient of variation (PCV) for all character was higher than genotypic coefficient of variation (GCV). The estimation of broad sense heritability  $\hat{h}2(BS)$ % was observed to be lower than those of broad sense heritability  $\hat{h}2(BS)$ % for all characters. Genetics advance was recorded as maximum for eugenol content (96.65). Highest heritability in broad sense  $\hat{h}2(BS)$ % recorded for days to maturity (99.46). On the basis of study accessions G-4, G-7, G-9, G-11, G-18 and G-25 were identified high oil of better quality. These accessions may be exploited for commercial cultivation.

**Keywords:** GCV; Genetic advance; Heritability; *Ocimum*; PCV; Variability

## Introduction

Ocimum (family-Lamiaceae) is a genus of about 200 species of annual and perennial aromatic herbs and shrubs. Most species are native to the tropical and warm temperate regions [1]. The dry herb (leaves), Ocimum leaf tea, essential oil and its chemical derivatives (eugenol, methyl-eugenol, linalool etc) are exported to European countries in sizable quantity every year [2,3] Basil has several medicinal properties. It is rich in carbohydrates, fibre, phosphorous, calcium, protein, iron, beta-carotene, vitamins B1 and B2 and in aromatic oils [4]. It is good for colds and coughs, indigestion, stomach pain, diarrhoea, nausea, ulcers, ringworm and asthma. It is said to lower blood sugar and increase lactation. The essential oil is also used as anti-perspirant and as fly and mosquito repellent [5]. Available genetic stocks at CSIR-CIMAP, 180 genetic stocks belonging to 5 Ocimum species-Ocimum sanctum-Krishna and Shyam tulsi, O. kilimandscharicum, O. africanum, O. gratissimum and O. basilicum including 8 varieties: CIM Ayu, CIM Angana, CIM Saumya, CIM Jyoti, CIM Sharada, CIM Kanchan, Kushmohak, Vikarsudha used in this investigation.

For any crop improvement program, germplasm collection and assessment of genetic variability is an important step. Hence, it is very essential to partition the observed variability into heritable and non-heritable components measured as genotypic coefficient of variation (GCV), phenotypic coefficient of variation (PCV), broad sense heritability  $h^2(BS)$ %, genetic advance (GA), and genetic advance expressed as percent mean (GAM%) [6]. Surveys of genetic variability with the help of suitable parameters such as GCV, heritability estimates, and GA are very necessary to start an efficient breeding program [7]. Heritability in associated with genetic advance over

means (GAM%) is more effective and reliable in predicting the resultant effect of selection [8,9]. Thus, the present study was conducted in 60 genotypes of *Ocimum* to evaluate genetic variation among 12 quantitative characters.

## **Materials and Methods**

The present investigation comprising different experiments was carried out at the Research Farm of CSIR-Central Institute of Medicinal and Aromatic Plants, Lucknow, Uttar Pradesh (INDIA) located 26.9°N latitude 80.5°E longitude with an elevation of 120 m above the mean sea level having the subtropical climatic conditions during Kharif season.

180 collections were assembled from different state of India including few exotic from 6 countries (Tanzania, Thailand, Singapore, Slovak Republic, USA and South Africa). Removing duplicates, sixty genetic stocks were examined for high herb and essential oil. These sixty genetic stocks of *Ocimum* are belonging to five species: 46-*Ocimum basilicum* (French basil, Sweet basil, Zanzibar basil, Indian basil, and Thai basil); 5-*Ocimum sanctum* (Krishna/holy basil, Shyam tulsi and Scare basil); 2-*Ocimum kilimandscharium* (Champhor tulsi); 5-*Ocimum africanum* (Hoary basil); 2-*Ocimum gratissimum* L. (African basil/Van tulsi/Tree basil/clove basil) (Table 1).

## Observations

Twelve morph-metric observations were recorded on the following economic traits as below:

Days to flowering (50%), Plant height (cm), Number of branches/ plant, Fresh herb yield/ plant (g), Oil content (%), Oil yield/plant (g), Inflorescence length (cm), Days to maturity, Methyl chavicol content (%), Linalool content (%), Citral content (%), Eugenol content (%). Citation: Smita S and Kishori RL (2018) Estimation of Genetic Variability, Heritability and Genetic Advance for Essential Oil Yield and Related Traits in Genus Ocimum . Adv Crop Sci Tech 6: 350. doi:10.4172/2329-8863.1000350

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S.No	Accessions/Genotypes/cultivar	Botanical name	Origin				
1	CIM-Ayu	Ocimum sanctum	CSIR-CIMAP, Lucknow U.P. (India)				
2	Vikarsudha	Ocimum basilicum	CSIR-CIMAP, Lucknow U.P. (India)				
3	CIM-Angana	Ocimum sanctum	CSIR-CIMAP, Lucknow U.P. (India)				
4	French basil	Ocimum basilicum	Bangalore, Karnataka (India)				
5	French basil	Ocimum basilicum	Mangalore, Karnataka (India)				
6	CIM-Jyoti	Ocimum africanum	CSIR-CIMAP, Lucknow U.P. (India)				
7	Sweet basil	Ocimum sanctum	Gandhi Nagar Gujarat (India)				
8	Selection I	Ocimum africanum	Chennai, A.P. (India)				
9	Sweet basil	Ocimum basilicum	Singapore				
10	Sweet basil (Kushmohak)	Ocimum basilicum	CSIR-CIMAP, Lucknow U.P. (India)				
11	CIM-Kanchan	Ocimum santum	CSIR-CIMAP, Lucknow U.P. (India)				
12	Indian basil	Ocimum basilicum	CSIR-CIMAP, Lucknow U.P. (India)				
13	French basil	Ocimum basilicum	Mangalore, Karnataka (India)				
14	Indian basil	Ocimum basilicum	Muzaffarpur, Bihar (India)				
15	Indian basil (CIM Saumya)	Ocimum basilicum	CSIR-CIMAP, Lucknow U.P. (India)				
16	Sweet basil	Ocimum basilicum	Udaipur, Rajasthan (India)				
17	Zanzibar basil	Ocimum basilicum	Tanzania				
18	Indian basil	Ocimum basilicum	Bareilly, Uttaranchal, (India)				
19	French basil	Ocimum basilicum	CSIR-CIMAP, Lucknow U.P. (India)				
20	Indian basil	Ocimum basilicum	Lucknow, U.P. (India)				
21	Indian basil	Ocimum basilicum	Lakhimpur (Kheri), U.P. (India)				
22	Kapoor/camphor tulsi	Ocimum kilimandscharicum	CSIR-CIMAP, Lucknow U.P. (India)				
23	French basil	Ocimum basilicum	Nasik, Maharashtra (India)				
24	French basil	Ocimum basilicum	Lucknow, U.P. (India)				
25	Indian basil (sel-2)	Ocimum basilcum	CSIR-CIMAP, Lucknow U.P. (India)				
26	Sweet basil (Selection-1)	Ocimum basilicum	CSIR-CIMAP, Lucknow U.P. (India)				
27	Sweet basil	Ocimum basilicum	Trivandrum, Kerala (India)				
28	French basil	Ocimum basilicum	Lucknow, U.P. (India)				
29	CIM- Surabhi	Ocimum basilicum	CSIR-CIMAP, Lucknow U.P. (India)				
30	French basil	Ocimum basilicum	Haridwar, Uttaranchal (India)				
31	Sweet basil	Ocimum basilicum	CSIR-CIMAP, Lucknow U.P. (India)				
32	Thai basil	Ocimum basilicum var. thyrsiflora	Thailand				
33	Shaym tulsi	Ocimum sanctum	Puralia, W.B. (India)				
34	Camphor basil	O. Kilimandscharicum	CSIR-CIMAP, Lucknow U.P. (India)				
35	Sweet basil	Ocimum basilicum	Jammu (J. and K. (India)				

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36	Lemon basil	Ocimum africanum	Phagwara, Punjab (India)		
37	Indian basil	Ocimum basilicum	Barabanki U.P., (India)		
38	Clove basil	Ocimum gratissimum L	Shillong, Meghalaya (India)		
39	Indian basil	Ocimum basilicum	Razaganj, U.P. (India)		
40	Indian basil	Ocimum basilicum	Rishikesh, Uttaranchal (India)		
41	French basil	Ocimum basilicum Bangalore, Karnataka (India)			
42	French basil	Ocimum basilicum Mangalore, Karnataka (India)			
43	French basil	Ocimum basilicum	Chandigarh		
44	Sweet basil	Ocimum basilicum	CSIR-CIMAP, Lucknow U.P.		
45	Sweet basil	Ocimum basilicum	Singapore		
46	Sweet basil	Ocimum basilicum	Singapore		
47	Sweet basil	Ocimum basilicum	Singapore		
48	Sweet basil	Ocimum basilicum	CSIR-CIMAP, Lucknow U.P. (India)		
49	Sweet basil	Ocimum basilicum	Košice, Slovak Republic		
50	Lemon basil	Ocimum africanum	Gandhi Nagar Gujarat (India)		
51	CIM-Sharada	Ocimum basilicum	CSIR-CIMAP, Lucknow U.P. (India)		
52	Lemon basil	Ocimum africanum	Mangalore, Karnataka (India)		
53	Sweet basil	Ocimum basilicum	Košice, Slovak Republic		
54	Sweet basil	Ocimum basilicum	Gandhi Nagar Gujarat (India)		
55	Sweet basil	Ocimum basilicum	Singapore		
56	French basil	Ocimum basilicum	Chennai, A.P., (India)		
57	Clove basil	Ocimum gratisimum	CSIR-CIMAP, Lucknow U.P. (India)		
58	Sweet basil	Ocimum basilicum	Gandhi Nagar Gujarat (India)		
59	Sweet basil	Ocimum basilicum	Gandhi Nagar Gujarat (India)		
60	Sweet basil	Ocimum basilicum	Singapore		

Table 1: Accessions of Genus Ocimum their botanical name and geographic origin.

#### **Essential oil analysis**

The fresh 100 gm aerial part of *Ocimum* spices were collected from plants of the field of CSIR-CIMAP, were processed by hydrodistillation for 3 hrs in a Clevenger apparatus to obtain the essential oil [10]. Identification of the essential oil composition was analyzed by gas chromatography (GC) and (MS).

#### **Statistical Analysis**

Data were tabulated and mean data were subjected to statistical analysis using computer program package developed at the Data Processing Division of the CIMAP, Lucknow, which based on Singh and Choudhary [11].

#### Analysis of variance (ANOVA)

The analysis of variance was based on the model; Xijk=m+gi+rj +eijk

Where, Xijk=effect of kth observation of ith genotype in jth replication, m=general mean, gi=effect of i<sup>th</sup> genotype, rj=effect of jth replication, eijk=environmental effect associated with ijk<sup>th</sup> observation.

Sources of Variation	df	Mean sum of product (MSS)	Expectation
Replications	r-1	Rms	
Genotypes	g <sup>-1</sup>	Gms	
Error	(r <sup>-1</sup> ) (g <sup>-1</sup> )	Ems	

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Total	(rg <sup>-1</sup> )		]

#### Parameters of genetic variability

 Table 2: Break up of variance components.

The ANOVA based on this model led to the following break up of variance components (Table 2).

In order to evaluate and quantify the genetic variability among the genotypes for the characters and study the following parameters were estimated (Table 3).

Genetic parameters	Days to flowering (50%)	Plant height (cm)	Number of branches	Inflorescence length (cm)	Fresh herb yield/plant (g)	Oil content (%)	Oil yiel/ plant (g)	Days to maturity	Methyl chavicol content (%)	Linalool content (%)	Citral content (%)	Eugenol content (%)
Genotypic variance (σ <sup>2</sup> g)	74.05	371.97	6.15	72.33	98350.78	0.048	11.597	764.39	243.16	135.69	59.71	6.58
Phenotypic variance $(\sigma^2g)$	79.799	523.31	15.74	76.23	216420.4	0.0594	19.63	768.47	247.86	137.14	60.45	12.27
Genotypic coefficient of variation (G.C.V.) %	5.47	18.05	22.05	13.16	43.57	34.64	23.93	74.65	94.41	263.73	258.71	55.47
Phenotypic coefficient of variation (P.C.V.) %	5.68	21.41	35.28	13.51	64.63	38.37	31.14	74.85	95.32	265.12	260.32	75.74
Heritability (broad sense) ĥ2(BS)%	92.78	71.08	39.05	94.87	45.44	81.49	59.07	99.46	98.11	98.94	98.77	53.63
Genetic advance over mean (GAM %)	25.63	26.42	17.72	86.43	40.79	57.81	89.89	36.01	85.21	51.03	63.59	94.65

**Table 3:** Genotypic ( $\sigma^2$ g) and phenotypic ( $\sigma^2$ p) variance, Genotypic (GCV) and phenotypic (PCV) coefficient of variation, heritability  $\hat{h}^2$ (BS) and genetic advance (GA as % of mean) of twelve quantitative traits of 60 genotypes of Genus *Ocimum*.

## Estimation of variance components

Genotypic and phenotypic coefficient of variability: Genotypic and phenotypic coefficients of variability were computed according to Burton and Devane [12].

Genotypic coefficient of variability (*GCV*) =  $\frac{\sqrt{\sigma^2 g}}{x} \times 100$ 

Phenotypic coefficient of variability (*PCV*) =  $\frac{\sqrt{\sigma^2 p}}{x} \times 100$ 

Environmental coefficient of variability (*ECV*) =  $\frac{\sqrt{\sigma^2 e}}{x} \times 100$ 

Where,  $\sigma 2g$ =Genotypic variance,  $\sigma 2p$ =Phenotypic variance and  $\sigma 2e$ =Environmental variance,  $\overline{X}$ =General mean of character

The PCV and GCV values are ranked as low, medium and high [13] as follows: 0-10%-Low; 10-20%-Moderate; >20%-High.

Heritability: Broad sense heritability was estimated based on the ratio of genotypic variance to the phenotypic variance and was expressed in percentage [14].

$$\hat{h}^2(BS)\% = \frac{V_g}{V_p} \times 100b$$

Where  $h_2$  (BS)%=heritability in broad sense; Vg=Genotypic variance; Vp=Phenotypic variance. Heritability values are categorized as low, moderate and high [15] as follows: 0-30%: Low; 30-60%: Moderate; 60% and above: High.

Genetic advance: The extent of genetic advance is expected by selecting certain proportion of the superior progeny was calculated by using the following formula [15] given by Robinson et al.

Genetic advance (GA)=k. σp.h<sup>2</sup>

Where: k=selection intensity at 5% (k=2.06);  $\sigma$ p=Phenotypic standard deviation;  $\hat{h}2(BS)$ %=Heritability in broad sense.

Genetic advance expressed as percentage over Mean (GAM%):

$$GAM(\%) = \frac{GA}{\overline{X}} \times 100$$

Where: GAM (%)=Genetic advance over mean;  $\overline{X}=$  General mean of the character

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# **Results and Discussion**

The present investigation was carried out to assess the genetic variability, path analysis between traits as related by genotypic and phenotypic relationship (correlations) and character contribution in 180 genetic stocks of Ocimum [16]. All the twelve traits were highly significant (p<0.01%). The phenotypic variances are always greater than the genotypic variances. The maximum amount of genotypic coefficient of variation (GCV) in percent recorded for the traits was linalool content (263.73%) followed by citral content (258.71%), methyl chavicol content (94.41%), date of maturity (74.65%), eugenol content (55.47%) and fresh herb yield (43.57%) was found suitable for favourable selection for further genetic improvement. Other traits like, oil content (34.64%), oil yield/plant (23.93%), number of branches (22.05%), plant height (18.05%), inflorescence length (13.16%) were expressed medium GCV. The days of flowering (50%) was recorded least (5.47%) genotypic coefficient variance. The phenotypic coefficient of variation (PCV) values were higher than their corresponding GCV for all traits under study, which reflects that the apparent variations were not only due to genotypes alone but environment also. The maximum amount of PCV was displayed by these characters namely, linalool content (265.12), chiral content (260.32%), methyl chavicol content yield (95.32%), eugenol content (75.74%), days to maturity (74.85%), fresh herb yield (64.63%), oil content (38.37%), number of branches (35.28%), oil yield/ plant (31.14%), plant height (21.41%) and inflorescence length (13.51%). The lowest estimate of PCV (5.68%) was observed for days of 50% flowering respectively [17].

## Estimates of heritability $\hat{h}^2$ (BS)% in broad sense

Information on heritability in broad sense h2(BS) and genetic advance of yield attributing traits and their association helps plant breeder to identify characters for effective selection [18]. The concept of heritability explains whether differences observed among individuals rose as a result of differences in genetic makeup or due to environmental forces. According to Robinson et al. [15], heritability values are categorized as low from 0-30%, moderate from 30-60% and 60% and above are high. Considering this benchmark, heritability estimate of this study are described as follows.

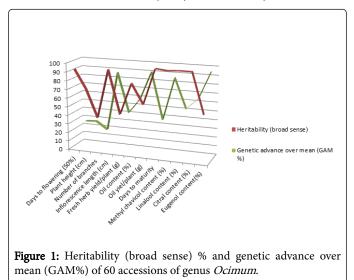
Heritability in broad sense  $h^2$ (BS)% parameter also played a major role in unison with character association and path analysis in genetic improvement program of *Ocimum*. The high heritability in broad sense  $h^2$ (BS)% was observed for date of maturity (99.46%) followed by linalool content (98.94%), citral content (98.77%), methyl chavicol content (98.11%), inflorescence length (94.87%), days to flowering 50% (92.78%) and oil content (81.49%) in order. Hence, the selections of above mentioned traits are easy in this crop. Moderate heritability was observed for plant height (71.08%) followed by oil yield/plant (59.07%), eugenol content (53.63%) and fresh herb yield (45.44%) respectively. Low heritability was observed in branches/ plant (39.05%). Therefore, selections of these traits are difficult due to high environmental influences [19].

#### Genetic advance over mean (GAM%)

The expected genetic advance (GA) values for 12 characters of the genotypes evaluated are presented in Table 2. High genetic advance over mean were observed for the traits eugenol content (94.65) followed by oil yield/plant (89.89), inflorescence length (86.43) and methyl chavicol content (85.21). The medium genetic advance was noticed in citral content (63.59), oil content (57.81), linalool content

(51.03), fresh herb yield/plant (40.79) and days to maturity (36.01). Remaining three characters exhibited low genetic advance for example plant height (26.42), days of flowering (50%) (25.45) and number of branches (17.72) respectively (Figure 1).

The estimates of genetic advance help in understanding the type of gene action involved in the expression of various polygenic characters. High values of genetic advance are indicative of additive gene action whereas low values are indicative of non-additive gene action [20]. Accordingly, Heritability and genetic advance are important selection parameters. The estimate of genetic advance is more useful as a selection tool when considered jointly with heritability estimates [21].



#### Conclusion

Analysis of variance disclosed highly significant difference among the accessions for all the parameters studied. In this research the phenotypic and genotypic coefficient of variation was found to be high for citral content and methyl chavicol content while, moderate GCV and PCV was noticed for days to maturity and eugenol content. Days to flowering (50%) recorded least phenotypic and genotypic coefficient of variation. The heritability estimates was noticed to be high for all characters studied. The characters viz. Days to maturity, linalool content, citral content, methyl chavicol content, inflorescence length and days to flowering (50%) exhibited high heritability coupled with a high genetic advance representing that simple selection scheme would be good enough for these traits to enhance genetic improvement in desired direction. On the basis of study accessions G-4, G-7, G-9, G-11, G-18 and G-25 were identified high oil of better quality. These accessions may be exploited for further crop improvement for the development of high essential oil yielding Ocimum cultivars of better quality.

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