# Evaluation of Arabica Coffee (*Coffea arabica* L.) Genotypes for Biometrical Traits

### S. Muthuramalingam\*, K. Rajamani and C. Krishnamoorthy

Department of Spices and Plantation Crops, Horticultural College and Research Institute Tamil Nadu Agricultural University, Coimbatore - 641 003

An experiment on arabica coffee (Coffea arabica L.) genotypes for biometrical traits evaluation was carried out at Chettalli in Karnataka during the period 2004 - 2007 to study the variation in the vegetative and yield component traits. Eighty five genotypes exhibited significant variation for all the traits in three seasons of the study. The per se performance of present study revealed that the traits viz., plant height, bush diameter, stem girth, number of primary branches, length of primary branches, internodal length, leaf area, number of flowers per axil, number of flowers per node, fruit length, fruit width, number of fruits per node, fruit set percentage, weight of fruits per plant and percentage of fruit fertility were high in Tafarikela, Maragogipe, Cauvery x SIn.9 and S.12, Kaffa. For the trait viz., percentage of A grade beans and weight of 100 A grade beans, the genotypes Cioccie, Agaro, Maragogipe and Taferikela recorded the highest per se values. Mean percentage of caffeine content was higher in the genotype Kents, closely followed by Agaro, Maragogipe, Tafarikela and Cauvery x Sln. 9. Hence the genotypes can be exploited in selection for further breeding programmes for the respective traits. The genotypes Taferikela, Maragogipe, Cauvery x Sln.9, S12, Kaffa and Agaro recorded higher fruit length and fruit width. Mean performance of clean coffee per hectare showed that the genotype Tafarikela registered the highest value followed by Maragogipe, Cauvery x SIn.9, S12, Kaffa and Agaro.

Key words: Genotypes, biometrical traits, per se performance, caffeine, grade, clean coffee

Coffee is an important beverage crop used all over the world. It is native of Ethiopia and occupies a place of pride in international trade next to petroleum. Major coffee producing countries are Brazil, Cameroon, Congo, Costa Rica. Ethiopia, Guatemala, Guinea, Kenya, Puerto Rico and Sudan. India has emerged as a quality coffee producer with a total production of 288000 MT from 328157 hectares during 2006-07 accounting for a world share of 4.45 per cent. In India, it is mainly cultivated in Karnataka, Kerala, Tamil Nadu, Andhra Pradesh and North East regions. The stimulative effect of coffee is due to the presence of caffeine content. Coffee consumption has been associated with reduced risk of several diseases including certain cancers, Parkinson's disease, hepatic diseases, kidney stones and also for increased mental alertness, reduction of fatigue and improvement in the performance on vigilance tasks.

However in the export trade, India does not hold any key place in realization of its potential which necessitates improvement in yield and quality. In view of breeding constraints and complexities, the identification of potential genotypes has been considered very important. In genus coffea, only two species *Coffea arabica* L. and *C. canephora* are commercially cultivated in India. *C. arabica* is self

\*Corresponding author email: muthuramhort@yahoo.co.in

compatible tetraploid while *C. canephora* is selfincompatible diploid species (Srinivasan *et al*, 1979). In order to increase the yield, productivity and quality components of this crop, crop improvement research becomes inevitable. Selection of promising types from the gene pool is the most useful method of crop improvement in this crop.

The primary objective of any crop improvement programme is identification of promising genotypes for which knowledge on variability existing in a population is imperative. Studies on variability helps in locating the desirable plant type either for direct introduction or to use in the further breeding programme. The yield being a complex character, knowledge on the optimal plant characters, the extent of variability available and character association with yield, is essential to improve the vield level in coffee. The genetic variability and association of different characters help us to develop our effective selection of suitable parents in breeding. Hence identification of coffee genotype that exhibits a consistently higher rate of transmission of their desirable characters to the progeny would be highly useful for improvement of yield.

#### **Materials and Methods**

The experiment was conducted at Coffee Board, Coffee Research Sub Station, Chettalli, North Coorg Dist, Karnataka which is situated at  $12^{\circ}N$  latitude  $75^{\circ}E$  longitude and at an elevation of 1038 m above mean sea level. The study was carried out during three seasons viz., season I (2004 –2005), season

II (2005-2006) and season III (2006 –2007). Eighty five genotypes maintained in the arabica coffee germplasm collection at Coffee Board, Coffee Research Sub Station, Chettalli, Coorg dist, Karnataka, were selected for this study. The details of the genotypes are presented in Table 1.

The experimental plot size was 30 m x 4 m with spacing of 2m x 2m accommodating 30 plants per plot. All the genotypes taken for study were collected from different countries. The plants were allowed to grow on topped single stem system under a mixed shade canopy and all plants were sprayed with Bordeaux mixture twice in a year and standard cultural practices were adopted. Uniform fertilizer application at the rate of 40:30:40 NPK kg/ha twice a year (pre monsoon and post monsoon) was given to all the plots. The experiment was laid out in a randomized block design with three replications. Five randomly selected plants in each genotype per replication were tagged for recording observations on plant characters and the mean values were subjected to statistical scrutiny. The selection of plants within genotypes and within field was done with care to see that there was no field and plot variation. Recommended package of practices were followed for all the genotypes (Anonymous, 2003). The observations recorded on eighty five coffee (Coffea arabica L.) genotypes for twenty characters viz., plant height, bush diameter, stem girth, number of primary branches, length of primary branches, internodal length, leaf area, number of flowers per axil, number of flowers per node, fruit length, fruit girth, number of fruits per node, fruit setting per cent, weight of fruits per plant, yield of clean coffee per hectare, percentage of fruit fertility (100 well developed fruits in hard endosperm stage were transversely cut and the fruit fertility was recorded), percentage of 'A' grade beans, weight of 100 A grade beans, caffeine content and per cent rust incidence were recorded.

## **Results and Discussion**

The data on mean performance, range and variability of genotypes were observed for twenty characters for three seasons (2004-2005, 2005-2006 and 2006- 2007).

Among the eighty five genotypes studied, the difference in morphological traits were noticeable among arabica coffee varieties. In all the three seasons, a wide range of variation was exhibited for plant height, bush diameter and stem girth in the Taferikela, Maragogipe, Cauvery x Sln.9 and S.12 Kaffa genotypes (Table 2). The increase in the plant height and yield of clean coffee across the genotypes were relatively uniform. The variation among the genotypes of coffee for the morphological attributes

Table 1. Genotypes and source	of genotypes of
arabica coffee	

arabica coffee								
S.No.	Genotype	Name	Source					
1	S.1495	Tafarikela	Ethiopia					
2	S.881	Rume Sudan	Sudan, Kenya					
3	S.1593	Jimma Galla Sidamo	Kenya					
4	S.1779	Limma Gimma Kaffa	Ethiopia					
5	S.1784	Misan Tafarikela	Ethiopia					
6	S.1783	Babaca Kaffa	Ethiopia					
7	S.1497	S. 12 Kaffa	Ethiopia					
8	S.1785	Ainamba Kaffa	Ethiopia					
9	S.1782	Badabuna Jimma Kaffa	Ethiopia					
10	S.882	Barbuk Sudan	Sudan					
11	S.1700	COF297 Zambeia Mocambic	Portugal					
12	S.1591	Dalle Melville	Kenya					
13 14	S.1590 S.1580	Dalle Mixed Dele CHO	Kenya					
			Kenya					
15 16	S.1582 S.1581	S-16 Wollamo 2 B Enneria	Portugal Portugal					
17	S.1583	S-17 Yrgalem	Portugal					
18	S.2130	Dilla and Alghe	Kenya					
19	S.2129	(110/2) S 4 Agaro	Portugal					
20	S.2128	(110/5) Agaro	Portugal					
21	S.2126	(113/1)S/6 Cioccie	Portugal					
22	S.2127	(113/2)S/6 Cioccie	Portugal					
23	S.1659	Kouti	Paris					
24	S.1699	Mbirizi de Mulungu	Portugal					
25	S.1698	Local Bronze De Mulungu	Congo					
26	S.1664	Nicaragua	Paris					
27	S.1726	Cundina Marca	Surinam					
28	S.1661	Mkoungan	Paris					
29	S.1663	Reunion	Paris					
30	S.1656	Salvador	Guinea					
31	S.1466	Leroy Laurina	Gautemala					
32	S.1742	Maragogipe	Tanganyika					
33	S.1493	Laurina Maragogipe	USDA					
34 35	S.1476 S.1768	Kona Murta	Gautemala Gautemala					
36	S.1708 S.1417	San Ramon	Guatemala					
37	S.1417	S12 Kaffa	Ethiopia					
38	S.1743	Giesha	Tanganyika					
39	S.1470	Blue Mountain	Gautemala					
40	S.1467	Sumatra	Gautemala					
41	S.1472	Phillipinean	Gautemala					
42	S.1648	Costa Rica	Guinea					
43	S.1694	Puerto Rico	Congo					
44	S.1471	Padang	Gautemala					
45	S.1468	Surinam	Gautemala					
46	S.1494	Mundo Novo	USDA					
47	S.1463	Туріса	Gautemala					
48	S.1464	Bourbon	Gautemala					
49	S.1490	Bourbon Amarelo	USDA					
50	S.1491 S.1492	Bourbon Vermelho	USDA					
51 52	S.1492 S.1504	Caturra Amarelo Mysore	USDA					
52 53	S.1304 S.1483	K.P. 423 (Kents)	Congo Tanganyika					
53 54	S.1483 S.1586	K.P. 423 ( Kenis ) K - 7	Tanganyika Kenya					
54 55	S.1905	S.L. 34	Burma					
56	S.1660	Kenya	Paris					
57	S.1693	C. Arabica578	Paris					
58	S.1735	Darnimin	Srilanka					
59	S.1465	Pache	Gautemala					
60	S.1696	Granja Bloomay	Congo					
61	S.1655	Mokablac	Guinea					
62	S.1695	Antigua	Congo					

Table 1. continued . . .

S.No.	Genotype	Name	Source	
63	S.1692	Barbirina	Congo	
64	S.1734	Martinique	Surmarno	
65	S.1652	Macenta	Guinea	
66	S.1697	Tumbadir	Congo	
67	S.1662	Foumbon	Paris	
68	S.1667	Mazeron	Paris	
69	S.1691	Las Palmas	Congo	
70	S.1690	Green Tipped	Congo	
71	S.1591	Delle Melville	Kenya	
72	S.1653	Togo Kouma	Guinea	
73	S.1473	Goudaluope	Gautemala	
74	S.1737	Madagaskar	Portugal	
75	S.1502	Bourbon Mayagese	Congo	
76	S.1726	Cundina Marca	Surinam	
77	S.2790	SIn.9	India	
78	S.4349	Cauvery	India	
79	S.4850	SIn.9 X Cauvery	India	
80	S.4843	Cauvery X Sln.9	India	
81	S.4854	S.881 X Cauvery	India	
82	S.4844	Cauvery X S.881	India	
83	S.4846	Taferikela X Cauvery	India	
84	S.4855	Cauvery X Taferikela	India	
85	S.4847	Cauvery X Cauvery	India	
may	be due	e to their diversified	origin and	

acclimatization of the particular genotype as

ical location

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morphotype in their specific geographical locations. Similar kind of report on evaluation of genotypes for growth and yield of clean coffee was made by Srinivasan and Subbalakshmi (1981). It clearly depicts that the expression of enhanced plant height among genotypes may be attributed to the differential ability of genotypes for the synthesis of phytohormones and nutritional factors, involved in the growth mechanism as reported by Santaram *et al* (1986), Ahmed and Sreenivasan (1988) and Ram *et al.* (1994) in coffee.

Taferikela, Maragogipe, Cauvery x Sln.9 and S.12 Kaffa were better performers in their growth characters viz., plant height, bush diameter and stem girth which indicated that these genotypes showed higher vegetative vigour which might be attributed due to better adaptability of the genotypes under Chettalli, Coorg dist, Karnataka condition than other genotypes. The best performance of the genotypes may be due to higher efficiency of genes in combination with environment made higher synthesis of carbohydrates due to increased photosynthetic efficiency would have resulted in better partitioning in reserved food. The results are in concordance with the earlier work of Sreenivasan

Table 2. Mean performance of arabica coffee genotypes for morphological characters (mean of three
seasons)

S.No.	Genotype	Plant height	Bush diamete	r Stem girth	No.of primary	Length of primary	Internodal
		(cm)	(cm)	(cm)	branches	branches (cm)	length (cm)
1.	S.1495	172.14	186.01	30.29	9.40	115.20	10.25
2.	S.881	123.95	123.89	26.67	6.22	91.70	8.95
3.	S.1593	93.88	119.20	21.91	4.68	77.94	6.25
4.	S.1779	80.50	122.31	26.07	4.15	71.38	6.52
5.	S.1784	104.71	139.59	27.57	4.68	79.94	7.93
6.	S.1783	104.01	138.74	26.07	3.78	82.44	8.18
7.	S.1497	143.91	175.74	28.61	7.44	105.98	10.49
8.	S.1785	93.36	138.66	27.30	4.35	93.39	7.60
9.	S.1782	80.47	144.67	27.21	4.08	82.09	8.96
10.	S.882	88.36	129.26	25.78	3.68	67.09	9.23
11.	S.1700	85.88	121.96	26.61	4.68	83.61	8.60
12.	S.1591	85.12	117.23	26.52	3.81	72.88	7.73
13.	S.1590	77.06	112.27	26.38	4.15	75.53	8.17
14.	S.1580	83.14	134.31	26.59	3.88	88.04	7.10
15.	S.1582	86.74	135.92	26.43	3.95	85.38	6.78
16.	S.1581	96.22	116.26	25.79	3.75	77.19	7.16
17.	S.1583	88.86	130.43	25.98	3.61	80.27	7.77
18.	S.2130	98.66	119.27	25.61	4.38	76.08	7.65
19.	S.2129	91.15	127.36	28.42	5.35	72.76	8.26
20.	S.2128	138.96	172.29	28.35	7.32	102.01	9.36
21.	S.2126	116.78	157.35	26.26	5.32	83.19	9.03
22.	S.2127	91.67	132.36	25.86	5.82	85.72	8.43
23.	S.1659	93.68	106.32	25.25	4.23	71.24	7.67
24.	S.1699	81.94	118.93	25.60	4.25	71.33	7.25
25.	S.1698	93.30	117.79	25.61	4.01	78.92	7.08
26.	S.1664	97.86	109.03	25.95	4.81	76.96	7.11
27.	S.1726	82.81	123.75	26.21	4.68	77.82	7.33
28.	S.1661	82.38	123.94	25.83	4.48	76.36	7.74
29.	S.1663	96.99	129.13	27.06	4.08	68.32	7.71
30.	S.1656	88.63	96.52	25.52	3.95	74.27	7.98

S.No.	Genotype	Plant height (cm)	Bush diameter (cm)	Stem girth (cm)	No.of primary branches	Length of primary branches (cm)	Internodal length (cm)
31.	S.1466	69.99	82.49	26.12	3.21	48.78	4.46
32.	S.1742	156.02	188.99	29.41	8.20	123.94	10.89
33.	S.1493	106.18	106. 59	27.16	3.71	73.55	9.21
34.	S.1476	65.70	115.78	23.37	2.41	80.37	4.13
35.	S.1768	67.56	76.87	22.79	2.60	44.22	3.48
36.	S.1470	85.95	74.05	21.18	4.81	40.17	3.04
37.	S.1497	88.11	97.52	24.25	3.81	67.03	7.95
38.	S.1743	115.44	153.19	26.45	6.09	88.56	9.05
39.	S.1470	98.53	132.50	27.05	4.55	72.48	7.12
40.	S.1467	97.56	109.30	25.51	4.01	76.27	5.78
41.	S.1472	109.00	136.26	25.23	4.31	76.80	8.24
42.	S.1648	83.27	106.02	27.13	5.18	75.86	7.19
43.	S.1694	67.73	78.59	23.67	3.14	47.95	4.13
44.	S.1471	77.52	108.83	25.75	4.95	68.44	8.36
45.	S.1468	92.91	104.51	25.42	4.08	78.57	7.72
46.	S.1494	107.33	106.56	24.79	4.55	78.06	7.30
47.	S.1463	79.23	114.78	25.58	3.95	71.39	7.19
48.	S.1464	83.77	126.22	25.13	5.19	71.03	7.81
49.	S.1490	93.35	119.53	25.20	5.05	74.89	7.29
50.	S.1491	97.72	111.13	24.82	5.42	59.64	8.13
51.	S.1492	93.39	100.63	25.19	4.88	78.44	6.79
52.	S.1504	99.66	113.84	25.21	5.32	71.64	6.34
53.	S.1483	134.06	162.03	28.41	7.17	100.28	9.85
54.	S.1586	90.97	105.49	25.57	4.68	88.21	7.36
55.	S.1905	87.75	116.39	25.65	3.81	65.22	7.23
56.	S.1660	96.38	128.43	25.55	5.35	79.60	7.61
57.	S.1693	84.10	108.73	26.64	5.12	81.78	7.35
58.	S.1735	106.11	113.95	27.47	4.65	81.81	7.33
59.	S.1465	72.88	86.13	23.72	3.31	49.03	5.41
60.	S.1696	104.78	112.15	25.14	5.08	83.84	8.12
61.	S.1655	93.33	100.33	25.16	4.51	76.39	7.17
62.	S.1695	103.90	130.43	26.79	4.51	80.41	8.28
63.	S.1692	91.74	110.19	25.71	4.08	69.80	8.17
64.	S.1734	99.99	126.15	24.93	4.68	83.01	7.04
65.	S.1652	94.43	123.48	26.00	5.16	73.50	6.80
66.	S.1697	97.11	115.38	25.21	4.18	75.95	7.59
67.	S.1662	93.31	121.00	25.39	4.95	85.62	6.86
68.	S.1667	83.11	129.00	26.19	5.15	76.15	7.16
69.	S.1691	94.86	118.23	25.82	3.68	70.63	7.09
70.	S.1690	85.15	110.30	24.88	5.69	82.37	7.29
71.	S.1591	103.29	106.59	25.09	4.82	68.09	7.36
72.	S.1653	100.20	128.39	24.63	5.25	77.75	7.32
73.	S.1473	100.21	100.94	25.75	4.31	86.59	7.06
74.	S.1737	93.32	123.54	26.60	4.35	80.60	7.11
75.	S.1502	91.57	115.07	25.76	4.75	81.74	6.79
76.	S.1726	86.82	121.47	25.36	4.01	80.80	7.41
77.	S.2790	125.55	143.14	27.02	6.02	87.36	6.52
78.	S.4349	99.35	109.23	27.02	6.57	72.34	8.63
70. 79.	S.4349 S.4850	110.65	155.36	27.36	6.52	75.54	6.80
79. 80.	S.4850 S.4843	149.73	181.33	28.74	7.61	137.14	9.98
80. 81.	S.4854	127.90	131.34	25.29	5.99	77.22	9.98 8.40
82.						93.78	8.40 10.37
	S.4844	111.01	162.11 102.64	27.86	5.91		
83.	S.4846	120.27	102.64	25.35	4.92	81.41	8.09
84. 95	S.4855	125.55	141.87	27.43	6.02	104.99	9.33
85.	S.4847	100.67	116.27	25.05	6.02	90.27	6.78
GM		98.1070	122.8442	25.9565	4.8382	78.9681	7.5819
SEd		8.2467	10.3763	1.1545	0.6686	7.3130	0.7326
CD (.0	JO)	16.2810	20.4853	2.2792	1.3199	14.4377	1.4463

Table 2. continued . . .

*et al,* (1993), Sadananda and Azizuddin (1996) and Arunkumar (2003) in turmeric.

The number of primary branches per plant was statistically significant among the genotypes of arabica coffee (Table 2). The findings of the present study revealed that the genotypes Tafarikala, Margogipe and Cauvery  $\times$  SIn.9 were superior in their performance to number of primary branches per plant which might be due to the vigour of the concerned genotypes. This indicates the suitability of these genotypes to Karnataka conditions than other genotypes. This is in consonance with the earlier work of Ram *et al.* (1992) in coffee.

Table 3. Mean performance of arabica coffee genotypes for leaf area and yield characters (mean of three seasons)

S.No.	Genotype	Leaf area	Number of	Fruit length	Fruit width	Fruit set	No.of fruits
	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	(cm₃)	flowers per axil	(mm)	(mm)	percentage	per node
1.	S.1495	101.18	10.90	23.93	17.78	88.30	23.48
2.	S.881	72.95	9.16	17.17	11.73	73.94	16.26
3.	S.1593	49.51	7.03	14.99	4.03	28.26	5.04
4.	S.1779	66.29	7.93	15.79	12.71	53.73	4.47
5.	S.1784	54.62	7.63	15.58	12.66	60.47	8.80
6.	S.1783	53.13	6.21	15.69	13.06	52.71	6.59
7.	S.1497	93.72	10.03	18.53	15.87	80.03	17.05
8.	S.1785	43.44	5.96	16.05	12.81	37.59	6.72
9.	S.1782	47.40	7.83	15.88	12.71	40.29	5.64
10.	S.882	49.29	5.69	15.76	13.36	72.58	6.60
11.	S.1700	43.50	6.93	16.06	12.91	47.99	5.78
12.	S.1591	52.93	5.69	16.08	13.02	52.17	5.41
13.	S.1590	46.43	6.69	16.18	14.21	37.47	7.20
14.	S.1580	42.39	7.02	16.56	13.46	46.83	7.63
15.	S.1582	42.17	7.36	16.79	13.20	34.11	5.20
16.	S.1581	45.08	6.69	16.52	12.86	68.63	5.85
17.	S.1583	43.31	7.86	16.14	13.10	60.09	6.04
18.	S.2130	45.75	6.68	15.54	12.55	28.38	5.57
19.	S.2129	51.18	6.52	15.58	12.00	62.40	8.53
20.	S.2128	88.42	10.03	18.00	15.12	80.01	16.55
21.	S.2126	66.12	8.36	16.40	13.76	65.25	15.75
22.	S.2127	63.58	8.47	16.50	13.10	69.57	6.95
23.	S.1659	48.38	7.69	16.76	14.06	24.10	4.69
24.	S.1699	48.19	7.36	16.71	13.59	38.80	7.48
25.	S.1698	43.25	5.69	15.91	13.56	39.72	6.31
26.	S.1664	52.39	7.03	16.54	14.32	35.66	4.48
27.	S.1726	50.17	7.86	16.95	12.70	37.47	5.77
28.	S.1661	44.77	7.03	14.68	11.39	46.48	7.05
29.	S.1663	42.23	6.02	14.82	13.01	23.76	5.63
30.	S.1656	46.85	7.03	15.10	14.46	27.42	4.32
31.	S.1466	36.82	4.64	13.14	10.64	19.07	3.87
32.	S.1742	101.18	10.54	20.06	17.01	84.72	17.65
33.	S.1493	55.01	6.22	14.65	14.46	55.09	4.29
34.	S.1476	30.38	3.01	10.88	9.70	16.24	3.50
35.	S.1768	32.83	4.85	11.76	10.03	17.61	3.75
36.	S.1470	51.15	8.70	15.51	13.66	49.17	12.66
37.	S.1497	56.08	8.19	17.06	11.73	63.40	10.12
38.	S.1743	80.29	7.86	16.91	14.36	70.85	11.34
39.	S.1470	52.40	6.35	16.09	12.78	62.31	6.33
40.	S.1467	50.98	6.36	16.09	13.59	26.76	5.97
41.	S.1472	62.42	5.68	16.50	12.78	26.65	6.38
42.	S.1648	60.92	6.36	16.90	11.73	43.13	5.93
43.	S.1694	34.11	4.46	12.63	11.86	18.35	4.06
44.	S.1471	64.99	7.83	16.14	12.11	51.50	15.22
45.	S.1468	56.99	6.68	15.99	12.77	62.51	4.91
46.	S.1494	57.64	7.83	609.48	12.65	25.58	5.50
47.	S.1463	57.92	5.95	16.14	12.20	38.29	5.66
48.	S.1464	50.15	5.41	16.14	13.43	46.14	6.21
49.	S.1490	61.05	6.56	16.79	12.14	48.72	6.83
50.	S.1491	50.43	7.69	16.74	12.14	26.29	6.77

Table 3. continued
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S.No.	Genotype	Leaf area (cm₃)	Number of flowers per axil	Fruit length (mm)	Fruit width (mm)	Fruit set percentage	No.of fruits per node
51.	S.1492	64.68	6.68	16.18	12.82	36.79	6.48
52.	S.1504	56.72	7.36	16.28	14.01	29.32	6.63
53.	S.1483	85.23	8.86	17.84	13.72	80.92	16.25
54.	S.1586	52.62	8.70	16.72	12.59	44.89	5.91
55.	S.1905	54.52	7.86	16.66	13.76	41.55	5.45
56.	S.1660	60.43	8.03	16.61	12.54	59.78	4.85
57.	S.1693	49.33	5.81	16.84	13.17	38.79	4.59
58.	S.1735	68.81	6.88	16.92	12.77	23.76	7.75
59.	S.1465	37.95	4.86	13.54	10.92	19.39	4.50
60.	S.1696	57.92	7.45	16.74	12.20	23.50	6.38
61.	S.1655	58.84	7.46	16.50	12.33	42.94	4.79
62.	S.1695	57.81	7.63	16.22	12.02	51.94	5.02
63.	S.1692	52.52	7.55	16.12	12.38	30.24	5.32
64.	S.1734	51.85	6.76	16.79	13.93	34.75	6.38
65.	S.1652	50.49	7.44	16.41	12.64	30.16	7.02
66.	S.1697	58.48	6.89	16.35	12.58	29.21	5.20
67.	S.1662	56.60	6.88	15.79	11.96	44.21	4.61
68.	S.1667	52.58	8.63	15.89	11.84	49.13	5.83
69.	S.1691	58.62	8.33	16.28	12.57	36.32	5.70
70.	S.1690	58.75	8.16	16.06	12.11	25.09	6.90
71.	S.1591	60.47	7.42	16.16	12.51	27.30	7.85
72.	S.1653	66.98	7.09	15.66	12.58	45.26	6.83
73.	S.1473	61.80	6.55	16.72	12.82	48.21	5.56
74.	S.1737	63.48	7.09	15.69	12.18	64.69	6.19
75.	S.1502	53.35	8.16	16.12	12.11	58.87	6.52
76.	S.1726	50.68	7.90	16.51	12.56	43.09	7.72
77.	S.2790	52.23	6.02	15.06	13.90	71.24	14.05
78.	S.4349	59.61	5.86	16.44	12.20	72.33	15.05
79.	S.4850	64.45	6.42	16.52	11.96	71.56	12.98
80.	S.4843	94.13	10.06	18.94	16.26	81.68	17.49
81.	S.4854	60.30	8.56	15.67	12.74	75.59	10.89
82.	S.4844	100.82	8.46	16.55	13.55	69.56	10.61
83.	S.4846	60.75	8.03	15.89	12.70	69.56	7.62
84.	S.4855	71.28	8.03	17.59	14.99	76.59	15.36
85.	S.4847	60.18	9.16	15.43	13.58	64.22	8.16
GM		57.3218	7.2537	23.1820	12.8744	47.9885	7.8612
SEd		8.3806	0.7868	NS	0.8777	6.5802	1.2761
CD (.	05)	16.5453	1.5533	NS	1.7328	12.9910	2.5192

Regarding the length of primary branches, the eighty five genotypes showed noticeable variations among themselves. The longest primary branch was recorded in Cauvery x SIn.9 (137.14 cm), followed by Tafarikala, Maragogipe in the present study. The length of primary branches decides the number of secondary and tertiary branches. Higher number of branches observed with respective genotypes would have resulted in high vegetative vigour and ultimately resulted in higher yield as reported by Srinivasan (1988), Selvakumar et al. (1992) Sadananda and Azizudin (1996), Nair and Nampoothiri (1993) and Ratnambal et al. (1995). The genotype San Ramon exhibited shorter length of primary branches (40.17 cm) followed by Murta, Puerto Rico and Leroy Laurina. Ram et al. (1992) also reported variations for this character among different arabica coffee populations.

Among the different arabica genotypes studied, the character internodal length showed notably high variation. The longest internode was recorded in Maragogipe (10.89) where as the genotype San Ramon exhibited the shortest internodal length (Table 2). Shorter internodal length gives more number of nodes per branch which facilitates higher yield in coffee. Narasimhaswamy (1965), Ahmed and Srinivasan (1988) and Kumar *et al.* (1999) also reported variations for this character among different coffee populations.

Mean leaf area, showed significant difference among the coffee genotypes studied. The leaf area is an important character, since it decides the ability of the leaf to support the yield of fruit and also the photosynthetic efficiency as reported by Srinivasan (1972), Orozco and Nieto (1972) and Selvakumar *et al.* (1992), Choudhuri and Hore (2004). The mean

S	60	
2	09	

S.No.	Genotype	Weight of fruits per	Yield of clean coffee	Percentage of fruit		Weight of 100 'A' grade	Caffeine content	% of rust disease
		plant (kg)	(kg ha₁)	fertility	beans	beans (g)	(%)	incidence
1.	S.1495	7.01	2827.79	97.78	76.51	18.86	2.80	12.17
2.	S.881	3.59	1975.55	88.93	73.07	15.64	1.98	17.59
3.	S.1593	1.30	809.33	68.55	58.19	16.14	0.76	29.26
4.	S.1779	1.72	889.67	77.24	60.20	14.79	0.71	17.49
5.	S.1784	1.49	752.89	78.60	65.79	14.14	0.34	25.55
6.	S.1783	3.01	1471.55	74.48	67.23	14.90	1.74	36.91
7.	S.1497	4.97	2244.78	95.68	75.87	19.02	2.68	16.06
8.	S.1785	1.47	634.78	77.25	59.67	15.11	1.60	29.26
9.	S.1782	1.80	896.33	70.90	54.86	13.93	1.29	26.01
10.	S.882	3.68	1762.56	75.91	59.77	15.28	0.95	28.14
11.	S.1700	2.51	1913.11	73.58	55.20	16.38	0.67	30.76
12.	S.1591	2.78	1591.89	73.78	62.22	14.79	0.53	30.20
13.	S.1590	2.58	1156.56	78.60	48.47	14.39	0.57	28.09
14.	S.1580	3.34	1471.56	73.12	46.60	14.31	0.75	26.37
15.	S.1582	2.22	1040.11	81.49	52.71	14.89	1.20	28.42
16.	S.1581	2.53	1150.44	73.69	51.76	16.34	0.93	31.77
17.	S.1583	1.79	893.00	79.90	45.01	13.82	0.87	51.12
18.	S.2130	1.38	759.11	70.90	43.27	14.36	0.92	38.38
19.	S.2129	3.35	1636.66	92.63	57.03	17.81	2.57	15.67
20.	S.2128	4.68	2182.56	94.93	78.50	19.69	2.87	15.93
21.	S.2126	3.33	1906.33	90.92	80.36	20.07	2.57	18.66
22.	S.2127	3.01	1902.55	91.53	61.54	17.62	2.41	18.12
23.	S.1659	1.87	1102.78	67.22	59.20	14.72	1.32	28.10
24.	S.1699	3.85	1337.56	80.60	38.12	15.42	0.52	30.11
25.	S.1698	1.95	908.00	71.90	34.52	13.86	0.78	38.93
26.	S.1664	1.59	709.00	79.01	58.47	15.40	0.72	42.84
27.	S.1726	1.73	923.00	70.67	50.67	16.06	1.24	43.18
28.	S.1661	2.49	7870.67	69.00	41.71	16.04	1.29	27.52
29.	S.1663	3.05	1424.78	71.57	40.13	14.32	0.77	31.69
30.	S.1656	1.67	816.00	80.50	58.05	15.24	1.09	38.67
31.	S.1466	1.10	583.56	63.69	30.13	11.60	0.53	43.03
32.	S.1742	6.02	2658.89	97.09	77.22	19.28	2.82	13.41
33.	S.1493	2.54	1200.78	83.08	66.72	17.71	2.65	24.66
34.	S.1476	1.02	485.89	63.48	28.27	11.26	1.09	34.41
35.	S.1768	1.06	530.22	63.55	30.68	11.51	1.52	39.47
36.	S.1470	1.59	1419.00	85.29	57.34	15.28	2.03	31.69
37.	S.1497	2.68	1311.67	90.03	67.57	17.80	2.59	20.61
38.	S.1743	4.18	1651.67	76.68	52.85	17.76	2.38	17.74
39.	S.1470	2.31	1070.33	78.93	38.13	14.41	0.79	30.96
40.	S.1467	2.54	1103.67	71.90	38.80	15.16	0.92	24.45
41.	S.1472	2.04	943.22	78.60	37.38	13.54	0.69	31.91
42.	S.1648	2.12	552.56	63.62	33.81	15.52	1.99	26.52
43.	S.1694	1.07	969.89	72.45	29.10	11.49	1.25	37.35
44.	S.1471	2.95	1384.56	83.62	40.47	15.56	0.94	37.55
45.	S.1468	3.27	1444.89	85.62	40.14	12.84	0.90	39.11
46.	S.1494	2.24	1083.56	81.14	44.48	15.22	1.51	32.24
47.	S.1463	3.02	1424.78	80.27	33.45	14.69	1.44	28.57
48.	S.1464	2.51	1157.22	74.57	43.66	13.99	1.64	41.81
49.	S.1490	2.76	1103.78	69.90	36.12	15.29	1.82	38.87
-50.	S.1491	2.91	1296.00	78.60	37.47	15.42	1.94	36.22
50. 51.	S.1491 S.1492	1.63	949.78	77.15	37.12	17.40	1.81	33.97
52.	S.1492 S.1504	2.95	1358.00	75.59	56.19	13.56	1.37	28.93
52. 53.	S.1304 S.1483	4.52	2108.78	89.99	75.14	18.74	2.93	26.93
53. 54.	S.1465 S.1586	4.52 2.51	1097.11	89.99 75.24	66.89	15.31	2.93 1.37	25.75
54. 55.	S.1566 S.1905	1.69	842.89	75.24 70.00	60.59 60.51	15.31	0.91	28.83 29.76

Table 4. Mean performance of arabica coffee genotypes for yield characters, quality characters and percentage of rust disease incidence (mean of three seasons)

Table 4. continued . . .

S.No.	Genotype	Weight of	Yield of clean coffee	Percentage of fruit	Percentage of 'A' grade	Weight of 100 'A' grade	Caffeine	% of rust disease
		fruits per			0	0	content	
		plant (kg)	(kg ha₁)	fertility	beans	beans (g)	(%)	incidence
56.	S.1660	1.55	776.00	71.57	42.81	14.48	2.08	39.55
57.	S.1693	1.48	762.55	71.90	41.14	16.78	1.74	30.94
58.	S.1735	2.30	1057.00	74.24	53.52	15.89	0.93	46.83
59.	S.1465	1.10	607.11	63.76	26.66	11.03	0.54	41.28
60.	S.1696	1.88	903.00	76.93	43.14	14.59	0.89	15.37
61.	S.1655	2.39	1063.55	77.14	36.99	15.98	0.92	28.52
62.	S.1695	1.68	1003.33	70.24	43.15	14.67	1.33	27.85
63.	S.1692	2.44	1070.33	78.05	37.12	14.41	1.01	33.36
64.	S.1734	1.55	769.22	76.92	62.68	15.25	0.91	43.39
65.	S.1652	1.38	763.55	68.22	40.45	14.17	0.72	23.89
66.	S.1697	2.93	1398.00	73.58	40.13	14.61	1.75	20.57
67.	S.1662	2.03	1016.66	73.71	53.86	15.49	1.85	27.88
68.	S.1667	1.85	1006.00	73.80	44.16	14.86	1.68	30.38
69.	S.1691	1.51	781.00	71.35	43.48	16.42	1.21	28.43
70.	S.1690	2.94	1384.56	78.93	43.59	16.09	0.97	21.30
71.	S.1591	2.14	1036.89	75.24	44.90	17.14	2.11	30.22
72.	S.1653	1.97	968.22	73.58	38.12	15.85	1.95	39.38
73.	S.1473	2.08	876.33	70.42	37.79	15.41	1.75	26.46
74.	S.1737	2.02	988.89	78.93	46.24	13.22	2.19	22.20
75.	S.1502	2.01	1003.33	70.24	55.52	15.84	0.93	26.39
76.	S.1726	1.78	903.00	68.07	43.37	15.44	1.49	18.29
77.	S.2790	4.11	1220.78	82.95	67.90	15.09	2.11	46.82
78.	S.4349	3.81	1287.67	84.96	62.88	15.39	2.08	50.33
79.	S.4850	3.90	1144.56	82.18	73.83	17.24	2.51	8.46
80.	S.4843	5.11	2374.56	96.23	61.53	18.21	2.73	15.25
81.	S.4854	3.17	1443.00	90.20	79.44	17.70	2.21	9.02
82.	S.4844	2.64	1711.23	90.87	68.73	17.17	1.70	15.14
83.	S.4846	2.14	769.89	86.29	67.62	17.34	C2.51	7.36
84.	S.4855	4.39	2029.00	87.09	71.91	16.43	2.63	25.00
85.	S.4847	2.01	859.22	76.92	74.37	17.18	2.42	37.97
GM		2.5560	1290.2587	77.8292	52.0156	15.5057	1.5191	29.0422
SEd		0.5808	1038.6230	6.4860	5.0012	1.3806	0.2915	5.8017
CD (.0	)5)	1.1466	2050.4943	12.8049	9.8735	2.7257	0.5755	11.4540

leaf area was highest in the genotype Taferikela and Maragogipe (101.18 cm<sub>3</sub>) (Table 3). As there was higher leaf area, the total photosynthetic surface on per plant basis was higher which in turn increased the total quantity of photosynthates. Similar results were also reported by Selvakumar *et al.* (1992). The genotype Kona showed the lowest leaf area (30.38 cm<sub>3</sub>) and also the reduction of leaf growth in these genotypes might be due to inhibition of auxin synthesis, inhibition in the rate of assimilation consequent to changes in the specific activity of enzyme and delay in the onset of first mitosis as reported by Arunkumar (2003) in turmeric.

The number of flowers per axil was the highest in the genotypes Taferikala (10.90) (Table 3). This might be due to genetic makeup of these genotype to produce more number of flowers per axil. This is in confirmation with the previous findings of Vishveshwara and Srinivasan (1977) and Thimma Reddy and Srinivasan (1979).

Wide range of variations among the genotypes were observed for the traits *viz*., fruit length and fruit

width. The highest mean values for fruit length and fruit width were recorded in Taferikela (23.93 mm) and Maragogipe (17.01 mm). Whereas the genotypes Kona and Murta exhibited the lowest values for fruit length and fruit width. Better fruit characters might be attributed by absorption of nutrients without any hindrance and the nature of genotypes. These results are in corroboration with the findings of Narasimhaswamy (1965) and Srinivasan (1972) in coffee.

Fruit setting percentage and number of fruits per node are important yield attributing characters and have direct influence on yield of coffee (Table 3). These characters could be considered as important traits for selection of genotypes for high yield. Variation in number of fruits per node and percentage of fruit setting were also observed over different seasons by Kumar *et al.* (1994). In the present study, the highest fruit setting percentage was observed in Taffarikela (88.30), Maragogipe (84.72) and Cauvery × SIn.9 (81.68). This might be due to high pollen germination percentage, higher number of flowers produced and size of the flower. Minimum level of fruit setting percentage and number of fruits per node were recorded in Kona followed by Murta, Puerto Rico. Similar results of variability were obtained by Orozco and Nieto (1972), Muluk (1987) and Renuga (1999) in coconut and Srinivasan (1988) in coffee.

The results of the present study revealed that the genotypes Taferikala followed by Maragogipe, Cauvery X Sln.9 and S<sub>12</sub> Kaffa exerted the highest weight of the fruits per plant and yield of clean coffee (Table 4). This might be attributed to higher bearing area through vegetative growth and number of fruits of the particular genotypes.

From the above results, it could be inferred that the genotypes which recorded the highest yield per plant might be attributed due to the said vegetative frame of these genotypes. The highest branches per plant and length of branches resulted in more vigour of plants which would have resulted in higher number of leaves, leaf area and leaf area index and inturn increases the efficiency of photosynthesis and photosynthetic rate for higher assimilation. Further the genotypes also registered higher number of flowers per plant and fruit setting per cent ultimately increased the yield (Table 4) . Conducive microclimate and soil fertility status are the major determining factors in the potential expression of a particular genotype for growth, development and yield. Of which the relative humidity, light intensity and leaf temperature are the important features which enhances the enzymatic processes in the plant growth and development phase. These high yielding genotypes have the capacity to utilize the environment. On the contrary, low yielding genotypes viz., Kona, Murta, Puerto Rica and Leroy Laurina were less vigorous in growth with poor expression of all the characters does not have the equilibrium with environment. The present finding is in conformity with the earlier results of Vishveshwara and Ahmed (1975), Sadananda and Azizudin (1996) in coffee.

In the present study, the genotypes with higher percentage of 'A' grade beans and weight of 100 'A' grade beans were observed in genotype Cioccie followed by Maragogipe in all the three seasons. Whereas, the lowest percentage of 'A' grade beans and weight of 100 'A' grade beans were recorded in the genotypes Pache, Kona and Puerto Rico (Table 4). Similar relationship was reported by Ahamed and Srinivasan (1988) and Srinivasan and Subbalakshmi (1984) in coffee.

There was significant variation for caffeine content, Among all the genotypes studied, Higher caffeine content was observed in the genotype Kents followed by Agaro and Maragogipe whereas, the lowest caffeine content was recorded in the genotypes Sumatra and Phillipinean (Table 4). The variation for caffeine content among the genotype might be due to the inherent genome constitution and environmental conditions provided for the particular genotype where it is grown. The dry weather or drought situation may increase the synthesis of the secondary metabolites. The variation in caffeine content might also be due to the effect of temperature, relative humidity and soil factors. This is in concordance with earlier works of Selvakumar and Sreenivasan (1996).

The percentage of rust disease incidence was statistically significant among the genotypes of arabica coffee. The finding of the present study revealed that the genotype Pache exerted the highest level of rust disease incidence followed by Barbuk Sudan, Phillipinean and Murta. The genotype Taferikala registered lesser percentage of rust disease incidence followed by Maragogipe, Cauvery X Sln.9, and S.12 Kaffa (Table 4). This might be due to the differential levels of sensitivity and tolerance possessed by individual genotypes to biotic and abiotic stress factors. These findings are in line with Vishveswara (1971) and Ram *et al.* (1994) who reported that certain genotypes possessed tolerance to diseases in coffee.

## Conclusion

From the study on mean performance of genotypes, based on various growth, yield and quality attributing traits, the genotypes *viz.*, Tafarikela, Maragogipe, Cauvery X Sln.9, S.12 Kaffa and Agaro showed high *per se* performance for high yield. These genotypes can be successfully used in breeding programmes for further exploitation.

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