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# Field Evaluation of *in vitro* Derived Mutants of different Varieties of Banana on Fruit Quality Parameters

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ABSTRACT: A field experiment was conducted on field evaluation of *in vitro* derived mutants of different varieties of banana during 2018-19 and 2019-20 at the Department of Fruit Science, College of Horticulture, University of Horticultural Sciences, Bagalkot. The 138 *in vitro* derived banana mutants both physical and chemical mutagens treated plants along with check were planted. The experiment was laid out in Augmented block design with 6 blocks and 26 subplots each block having 23 *in vitro* mutant lines and three checks. The different parameters like pulp weight, peel weight, pulp to peel ratio, fruit firmness, flavour acceptance and shelf life were taken for quality parameters. Among 138 *in vitro* derived mutants in plant crop, there was no significant difference noted in pulp weight, pulp to peel ratio and fruit firmness. The maximum pulp weight was recorded in NR45Gy-09 (86.45 g), peel weight, RAJ45Gy-14 (20.00 g) pulp to peel ratio YB35Gy-05 (6.70) flavour acceptance, YB40Gy-03 (7.82) fruit firmness, YB40Gy-12 (18.82 N) and shelf life YB40Gy-14 (8.10 days)). In ratoon crop, the maximum pulp weight was recorded in NR45Gy-09 (86.45 g), peel weight, was recorded in NR45Gy-09 (86.45 g), peel weight was recorded in NR45Gy-09 (86.45 g), peel weight, RAJ45Gy-14 (25.00 g) pulp to peel ratio, YB35Gy- 24 (7.36) flavour acceptance, RAJ45Gy -03 (7.48) fruit firmness, YB40Gy-11 (16.50 N) and shelf life YB45Gy- 08 (9.00 days).

Keywords: Banana, in vitro, physical mutants, chemical mutants, pulp weight, peel weight, shelf life.

### **INTRODUCTION**

Banana is a monocotyledonous herbaceous plant belonging to the section *Eumusa* under the family Musaceae. All edible bananas are originated from two species namely *Musa acuminata* and *Musa balbisiana* and most of the cultivated cultivars are their hybrids. The basic chromosome number in banana varies from 7-11 (X = 7, 9, 10, 11). The cultivated edible bananas are mainly triploids. As triploid varieties are highly sterile, edible plants are typically propagated by asexual methods (Simmonds and Shepherd, 1955).

Yelakki bale (AB) is the choicest diploid cultivar of banana, which is under commercial mono cultivation on a large scale especially in Karnataka and Tamil Nadu. It is popularly known as Puttabale, Mitle bale in Karnataka. It is slender plant bearing bunches of 12-14

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kg after 13-14 months. Fruits are small, slender having a prominent beak and arranged around the axis in a way that they bear a wind-blown appearance (Daniells *et al.*, 2001). Fruits are very attractive due to their bright yellow color with ivory white pulp.

Rajapuri bale (AAB) is a popular cultivar of banana grown in northern parts of Karnataka. It is a dwarf variety growing up to 6-8 feet height with a very thick stem and stands up very well to wind. The leaves are wider than those of most bananas growing up to 3 feet wide. It is the best plant to grow in marginal areas or where a grower does not intend to put much care into cultivation of bananas. The bunch weigh about 10-15 kg with 8-10 hands and 90-100 fingers. Fruits have attractive yellow colour with thick skin and good blend of sweet and acidity.

The Nanjangud Rasabale (Musa spp., AAB, silk subgroup) known for its unique taste has a huge demand across the country. But, conditions are not favorable enough for growing the banana and to match the huge demand that it generates (Rangaswamy, 2011). The cultivar Nanjangud Rasabale has been geographical indication tagged and once leading cultivar of then Mysore province is under threat of extinction due to its susceptibility to panama diseases (Pooja et al., 2013). Earlier, the cultivar grown in around 600 acres of land in the district of Mysore, Karnataka, now confined to only 30 acres of land in isolated area (Khan, 2015) due to susceptibility to wilt. Hence, the variety now which is at the danger of extinct has to strictly breed by potential methodology to save and preserve as a germplasm.

Most of the cultivated Musa varieties and cultivars are triploids. As triploid varieties are highly sterile, edible plants are typically propagated by asexual methods. Genetic improvement in banana has been extremely complicated due to varied genomic constitutions, heterozygosity, polyploidy, and the sterility in edible cultivars. Creating genetic variability for economically important traits in banana would not be supported application of conventional breeding programs due to the complexity in genome of Musa species. In this context, induced mutation has a high potential for bringing genetic improvements of vegetatively propagated crops like banana. The main advantage of induced mutations in vegetatively propagated plants is the ability to change one or a few characters of an outstanding cultivar without altering the remaining genetic background (Kulkarni et al., 2007).

Mutation induction using gamma rays had been applied to *Musa* spp. for improving many desirable traits, such as early flowering (Novak and Micke, 1990), tolerance to aluminium (Matsumoto and Yamaguchi, 1990) and *Fusarium* resistance (Chai *et al.*, 2004). A number of chemicals are able to induce mutations in banana plants such as sodium azide, 2,4-D (2,4-dichlorophenoxy acetic acid) and 6-Benzylaminopurine (6-BA) (Bhagwat and Duncan, 1998). Further, induction of mutation through chemical mutagens such as EMS and Sodium azide has generated several mutants of banana cv. Nanjangud rasabale under *in vitro* conditions (Kishor *et al.*, 2017). Ahmed (2003) showed that irradiated Grand Naine (clone 1 and clone 2) had longer green life than that of untreated Willams. Veena (2017) studied to characterize gamma irradiated *in vitro* regenerated banana cv. Grand Naine under field conditions. Maximum Total Soluble Solids and shelf life were found at 40Gy dose of irradiation.

Mutation induction may uncover a recessive phenotype by mutating, inhibiting or deleting the corresponding dominant allele (Jain and Swennen, 2004). Almost all the edible banana and plantain varieties originate through spontaneous mutations. The best example is the spontaneous banana mutant 'Cavendish' originating from Vietnam, which is resistant to Fusarium wilt (race 1) and which replaced 'Gros Michel' in the 1950s and 60s. The discovery of this banana mutant saved the banana industry from collapsing. In vitro mutagenesis, therefore, provides the necessary tool for the development of new clones resistant to diseases and also possessing improving characters like Shelf life, Pulp weight, and Flavour acceptance. However, field evaluation of in vitro derived mutants is ultimate answer for performance of newly mutated lines of banana under field condition for morpho-agronomic traits and also quality parameters. In this regard, the study has been planned in which the mutated (using physical and chemical mutagens) in vitro regenerated lines of the popular cultivars of Karnataka, viz., Rajapuri bale, Yelakki bale and Nanjangud Rasabale are field evaluated for morpho-agronomic and quality parameters.

## MATERIAL AND METHODS

The present investigation entitled "Field evaluation of in vitro derived mutants of different varieties of banana (Musa spp.)" was carried out during two years 2018-19 and 2019-20 at the Department of Fruit Science, College of Horticulture, University of Horticultural Sciences, Bagalkot. The 138 in vitro derived banana mutants (both physical and chemical mutagens treated plants) along with check were planted following Augmented Block Design at Fruit Orchard University of Horticultural Sciences, Bagalkot. The experimental site was divided into 6 blocks and 26 sub plots. The uniform pits of 60 cm<sup>3</sup> were dug out according to the plan of layout and recommended spacing  $(1.5m \times 1.5m)$ . Each pit was filled with 20 kg well decomposed farmyard manure. The banana mutants were subjected quality parameters after harvesting.

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	Origin of <i>in vitro</i> mutant line							
Sr. No.	(a) Yelakki Bale	(b) Rajapuri Bale	(c) Nanjangud Rasabale					
1.	Control (06 plants)	Control (06 plants)	Control (06 plants)					
2.	25Gy (04 plants)	25Gy (02 plants)	25Gy (0 plants)					
3.	30Gy (11 plants)	30Gy (06 plants)	30Gy (02 plants)					
4.	35Gy (18 plants)	35Gy (08 plants)	35Gy (04 plants)					
5.	40Gy (14 plants)	40Gy (12 plants)	40Gy (05 plants)					
6.	45Gy (12 plants)	45Gy (10 plants)	45Gy (04 plants)					
7.	EMS- 0.60% (2 plants)	EMS- 0.60% (1 plants)	EMS- 0.60% (1 plants)					
8.	EMS -0.90% (4 plants)	EMS -0.90% (3 plants)	EMS -0.90% (1 plants)					
9.	SA- 0.02% (1 plants)	SA- 0.02% (1 plants)	SA- 0.02% (1 plants)					
10.	SA- 0.03% (1 plants)	SA- 0.03% (2 plants)	SA- 0.03% (2 plants)					
11.	BAP-15 mg/l (1 plants)	BAP-15 mg/l (1 plants)	BAP-15 mg/l (1 plants)					
12.	BAP -20 mg/ 1 (1 plants)	BAP -20 mg/ 1 (1 plants)	BAP -20 mg/1 (1 plants)					
	75	53	28					
Total lines used 156								

Table 1: In vitro mutants of three cultivars of Banana.

Note: EMS - Ethyl Methyl Sulphonate; SA - Sodium Azide; BAP - Benzyl Amino Purine; Gy- Gamma irradiation



Plate 1. General view of experimental plot at shooting stage of in vitro mutants of different varieties of banana.

## **RESULTS AND DISCUSSION**

There was no significant difference noted in plant crop and ratoon crop of pulp weight among different mutant lines of banana and data is presented in (Table 2). In plant crop, the maximum and minimum pulp weight was found in NR45Gy - 09 (86.45 g), RAJ40Gy - 23 (35.00 g) respectively. In ratoon crop, the maximum and minimum pulp weight was found in RAJ45Gy - 07 (68.00 g), RAJ45Gy - 3 (35.07 g) respectively. Significant difference was noted in plant crop of peel weight among different mutant lines of banana and data is presented in Table 2. The maximum peel weight was found in RAJ45Gy - 14 (20.00 g). The minimum peel weight was found in YB30Gy - 09 (8.47 g). There was significant difference recorded in ratoon crop among different mutant lines of banana (Table 2). The maximum peel weight was found in RAJ45Gy - 14 (25.00 g). The minimum peel weight was found in YB25Gy - 02 (7.21 g).

This might be due to genetic makeup of the plants and also to some extent fruit size which contributes to the more peel weight. There is no report on variations in peel weight in mutant lines in banana. Hence, the results of present study revealed that there is a possibility of variations in peel weight in mutants due to mutations. However, the results of the present study are in line with Sagar *et al.* (2018) who has carried out evaluation of banana genotypes for qualitative traits. These results are in line with Balesh (2020) studied evaluation of elite tissue culture raised banana varieties for growth, yield and quality parameters.

There was no significant difference noted in plant crop and ratoon crop in pulp to peel ratio among different mutant lines of banana and data is presented in Table 2. In plant crop, the maximum pulp to peel ratio was recorded in YB35Gy - 05 (6.70) the minimum pulp to peel ratio was recorded in RAJ45Gy - 14 (2.60).). In ratoon crop, the maximum pulp to peel ratio was recorded in YB35Gy - 24 (7.36) the minimum pulp to peel ratio was recorded in RAJ45Gy - 14 (2.20).

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Mutant lines	Pulp v	p weight (g) Peel weight (g)		ight (g)	Pulp to peel ratio		
Wittant mics	Plant crop	Ratoon crop	Plant crop	Ratoon crop	Plant crop	Ratoon crop	
YB25GY 02	54.00	46.50	10.00	7.21	5.40	6.44	
YB25GY 03	48.20	49.00	13.22	8.63	3.69	5.67	
YB25Gy 04	55.60	51.00	9.50	9.12	5.85	5.59	
YB25Gy 05	42.38	47.20	11.00	7.69	3.86	6.13	
YB30Gy 01	45.21	56.75	13.84	10.06	3.25	5.64	
YB30Gy 02	58.00	44.00	12.26	7.68	4.73	5.72	
YB30Gv 03	42.00	40.00	9 50	8.05	4 42	4 96	
VB30Gv 04	50.00	52.80	9.82	11.20	5.09	4 71	
VB30Gv 06	49.00	61.00	11.20	9.81	4 37	6.21	
VB30Cy 07	55.13	57.21	10.67	9.64	5.16	5.03	
VB30Cy 08	58.21	53.00	12.54	8.25	1.64	6.42	
VB30Cy 00	43.50	48.00	8.47	9.42	5.13	5.00	
VP 20Cy 10	43.50	40.00	14.25	9.90	1.15	4.08	
1 B30Gy 10 VD20C 11	62.45	43.80	14.23	0.00	4.40 5.29	4.98	
1 B30Gy 11 VD20C- 15	03.43 50.56	34.00	12.00	0.59	3.20	4.77	
1 D30Gy 15 VD25C 01	39.30	40.00	12.62	9.30	4.04	4.60	
YB35Gy01 XB25G= 02	44.27	39.68	10.56	8.50	4.19	4.63	
YB35Gy 02	38.00	45.00	11.03	8.12	3.45	5.54	
YB35Gy 03	59.00	42.83	10.17	8.70	5.80	4.92	
YB35Gy 04	04.00	51.27	9.62	9.48	6.65	5.40	
YB35Gy 05	66.00	60.21	9.85	12.34	6.70	4.87	
YB35Gy 06	45.25	56.00	8.96	12.00	5.05	4.66	
YB35Gy 07	52.00	49.00	11.02	9.05	4.71	5.41	
YB35Gy 09	54.00	57.03	10.06	9.27	5.36	6.15	
YB35Gy 10	58.00	61.00	11.20	10.60	5.17	5.75	
YB35Gy 11	47.30	56.00	10.65	10.27	4.44	5.45	
YB35Gy 12	60.00	53.70	12.00	9.78	5.00	5.48	
YB35Gy 14	54.00	49.00	9.85	10.00	3.85	4.90	
YB35Gy 15	58.47	46.00	12.00	8.65	4.87	5.31	
YB35Gy 18	52.00	49.00	9.50	8.92	5.47	5.49	
YB35Gy 20	67.60	64.00	10.41	13.27	6.49	4.82	
YB35Gy 23	52.00	61.50	11.00	12.34	4.72	4.98	
YB35Gy 24	64.00	58.00	10.04	7.88	6.37	7.36	
YB35Gy 25	59.00	66.00	13.00	10.24	4.53	5.07	
YB40Gy 01	53.00	48.00	8.50	9.64	6.23	4.97	
YB40Gy 02	48.25	51.00	9.65	10.14	5.00	5.03	
YB40Gy 03	50.12	63.28	11.02	12.27	4.54	5.15	
YB40Gy 04	54.30	49.26	10.52	7.86	5.16	6.26	
YB40Gy 05	62.60	51.00	11.67	9.58	5.36	5.32	
YB40Gv 06	58.00	53.70	10.00	8.64	5.80	6.48	
YB40Gy 07	52.00	47.00	14 00	813	3.71	5 78	
YB40Gy 08	54.00	49.00	16.00	8.91	3 37	5 49	
VB40Gy 10	58.27	61.00	12.04	10.00	4 84	6.10	
VB40Gy 11	49.00	46.00	9.50	8 20	5.15	5.61	
VR40Gv 12	50.00	58.00	8.64	12.03	5.78	4.82	
VR40Gv 13	64.00	56.50	11.33	10.84	5.64	5.72	
VR40Gv 14	53.00	46.00	9.20	8 53	5.76	5 39	
VB40Gy 15	46.00	44.00	14.00	10.00	4 60	4 40	
VR45Cv 02	53 58	58.00	10.69	9.52	5.01	6.09	
VB45Cv 03	67.00	64.00	11.85	11.40	5.65	5.61	
VB45Cv 04	58.00	62.00	10.40	8 75	5 57	7.08	
VR45C-v 05	55.00	59.00	876	8 05	6.20	6 50	
VR45C-v 06	48.00	55.00	8 05	9.53	5 36	5 71	
VR45C v 07	63.25	57.00	10.00	9.02	632	6.06	
1 D45Gy U/ VR45Cy 00	40.20	54.00	0.85	10.26	5.00	5 21	
1 D45Gy 00 VR45Cy 00	47.20	52.00	7.03	0.00	5.00	5.21	
VD45C 11	50.06	65.00	0.42	10 59	5.55	5.00	
1 D45Gy 11 VD45G- 12	57.00	62.00	7.42 0 25	12.30	6.27	5.10	
1 D45(5y 12 VD45(-, 12	51.19	50.51	0.03	0.21	5.00	J.12 6 16	
1 D45Gy 15 VD45G- 15	50.00	55.00	12.00	9.21	2.90	0.40	
1 D43Gy 13 D 4 125C 61	30.00	33.00	10.54	0.20	3.84	0.00	
KAJ23GY UI DA 1250 62	47.51	44.00	10.54	13.01	4.50	3.23	
KAJ23GY U2 DA 120C 02	46.40	39.03	12.03	10.00	4.02	3.90	
KAJJUGY U2	39.50	42.01	14.00	12.18	2.82	3.49	
KAJ30Gy 03	43.18	47.00	12.30	12.50	3.31	3.74	
KAJOUGY 05	44./5	43.54	15.28	14.21	3.5/	3.06	
KAJ30Gy 07	46.52	58.25	11.31	13.36	4.11	2.86	
RAJ30Gy 12	53.00	55.64	16.04	9.30	3.30	5.98	
RAJ30Gy 13	64.25	39.22	17.50	10.52	3.67	3.72	
RAJ35Gy 01	58.60	46.00	10.04	12.89	5.83	3.56	
RAJ35Gy 02	68.59	38.51	18.35	10.00	3.73	3.80	
RAJ35Gy 03	56.00	42.00	11.00	9.50	5.09	4.42	

# Table 2: Quality parameters of plant and ratoon crop of banana mutant lines.

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RAJ35Gy 06	64.00	41.00	13.00	10.00	4.92	4.10
RAJ35Gy 07	55.00	59.00	9.50	15.12	5.78	3.70
RAJ35Gy 09	45.00	49.00	10.00	9.00	4.50	5.44
RAJ35Gv 10	57.00	53.06	12.30	11.00	4.63	5.27
RA 135Gy 12	64.00	47.00	14.00	12.60	4 57	3 73
DA 140C v 01	43.24	48.00	14.00	10.00	2.01	4.80
RAJ40Gy 01	43.24	48.00	14.04	10.00	2.91	4.00
RAJ40Gy 02	40.00	57.00	15.27	12.00	2.62	4.75
RAJ40Gy 03	44.00	61.00	14.00	10.23	3.14	5.96
RAJ40Gy 04	54.50	43.00	17.00	12.00	3.20	3.58
RAJ40Gy 05	49.64	56.00	16.00	13.00	3.10	4.30
RA.140Gv 08	40.82	49.00	15.25	10 51	2.67	4 66
RA 140Cy 00	54.20	56.00	14.50	13.00	3.73	4.30
DA 140C- 11	45.10	45.00	12.00	10.00	2.76	4.50
RAJ40Gy 11	45.19	43.00	12.00	10.00	3.70	4.30
RAJ40Gy 15	55.00	59.00	14.94	12.00	3.68	4.91
RAJ40Gy 22	42.64	37.00	9.54	11.50	4.48	3.21
RAJ40Gy 23	35.00	54.00	11.00	10.22	3.18	5.28
RAJ40Gy 25	55.00	49.50	14.63	11.00	3.75	4.50
RAJ45Gv 03	53.18	35.07	12.00	9.02	4 4 3	3.88
RA 145Gy 04	57.00	39.00	15.00	10.00	3.80	3.90
DA 1450- 07	57.00	68.00	10.00	16.00	2.20	1.25
RAJ45Gy 0/	04.30	08.00	19.00	10.00	5.39	4.23
RAJ45Gy 08	52.00	49.00	16.50	12.10	3.15	4.05
RAJ45Gy 10	51.50	54.00	19.00	11.28	2.71	4.78
RAJ45Gy 13	57.00	53.40	18.00	14.00	3.16	3.81
RAJ45Gy 14	52.00	55.00	20.00	25.00	2.60	2.20
RAJ45Gv 15	58.00	65.00	17.04	22.00	3.40	2.95
PA 145Cv 10	54.00	61.00	10.58	13.00	5.10	1 60
RAJ450y 19	54.00	50.00	10.58	10.00	5.10	5.00
KAJ45Gy 20	62.00	59.00	9.52	10.00	0.52	5.90
NR30Gy 06	51.00	56.00	12.00	11.00	4.25	5.09
NR30Gy 10	63.00	44.00	15.00	10.55	4.20	4.17
NR35Gy 08	58.00	49.00	10.00	10.00	5.80	4.90
NR35Gv 11	64.00	59.00	16.00	13.00	4.00	4.53
NR35Gy 12	60.07	52.00	14.00	11.14	4 29	4 66
NR35Cy 10	55.00	48.00	11.00	10.00	5.00	4.80
ND 40C 05	92.00	<del>4</del> 0.00	15.46	12.60	5.00	2.07
NR40Gy 05	82.00	54.00	13.40	15.60	4.03	5.97
NR40Gy 07	75.21	62.00	12.05	14.00	5.39	4.42
NR40Gy 09	68.00	48.00	13.80	10.38	4.92	4.62
NR40Gy 12	53.50	44.36	14.00	11.00	3.82	4.03
NR40Gy 15	64.00	42.00	12.55	9.00	5.10	4.66
NR45Gy 01	56.00	54.00	16.00	13.00	3.50	4.15
NR45Gy 02	54.00	46.00	13.00	9.00	4.15	5.11
ND45C 04	61.00	52.00	12.00	10.21	5.09	5.10
NR45Gy 04	01.00	55.00	12.00	10.21	5.08	J.19
NK45Gy 09	86.45	55.00	17.89	12.50	5.75	4.40
YB EMS 0.6% 1	53.07	43.86	9.56	8.64	5.55	5.07
YB EMS 0.6% 4	56.00	54.00	12.03	11.30	4.65	4.77
YB EMS 0.9% 1	59.24	46.00	10.82	9.58	5.47	4.80
YB EMS 0.9% 2	64.00	51.27	14.00	9.48	4.57	5.40
YB EMS 0.9% 3	54.37	49.60	8.97	9.20	6.33	5.39
VB FMS 0.9% 4	54.12	60.21	11.00	12 34	4.92	4 87
VR SA 0.02% 1	63.02	47.82	12.35	8 80	5.10	5.43
1D SA 0.02701	40.45	47.82 52.06	0.05	0.00	5.10	1.43
YB SA 0.05% 0	49.45	52.06	9.05	11.50	5.40	4.00
YB BAP15-1	59.21	46.00	12.82	9.58	4.61	4.80
YB BAP 20-2	55.03	48.25	10.56	8.34	5.21	5.78
RAJ EMS 0.6% 3	56.04	44.12	11.00	9.50	5.09	4.64
RAJ EMS 0.9% 1	64.32	55.02	13.06	10.34	4.92	5.32
RAJ EMS 0.9% 2	80.00	48.05	16.00	10.12	5.83	4.74
RAJ EMS 0.9% 4	63.04	54.00	10.56	9.00	5 97	6.00
RAISA 0.02% 1	58.25	51.30	12.00	11.00	4.85	4.66
DALSA 0.020/ 1	64.02	47.14	12.00	12.60	4.05	2.74
RAJ SA 0.03% 1	04.03	4/.14	14.00	12.00	4.37	5.74
RAJ SA 0.03% 2	53.00	48.05	14.84	10.00	3.56	4.80
RAJ BAP 15- 3	60.00	57.00	12.84	9.86	4.67	5.78
RAJ BAP 20 1	58.06	42.00	11.00	9.50	5.09	4.42
NR EMS 0.6% 3	56.10	53.00	9.54	10.65	6.33	4.41
NR EMS 0.9% 5	62.45	57.12	11.00	9.67	5.63	5.90
NR SA 0 02% 1	56.00	48 30	9 31	10.32	6.01	4 68
ND CA 0.020/ 1	40.04	46.22	9.51 9.40	0.21	5 64	5.02
NK 5A U.U5% 1	49.04	40.33	0.09	9.21	5.04	5.05
NR SA 0.03% 2	58.16	47.02	9.56	11.36	6.08	4.13
NR BAP 15-1	52.14	49.00	10.24	9.86	5.09	4.78
NR BAP20-3	48.05	43.21	9.22	7.62	5.20	5.67
YB	54.14	47.50	10.35	8.48	5.30	5.62
RAJ	59.47	51.78	10.95	9.68	5.44	5.35
NR	54 41	46.76	10.32	9.28	5 32	5.21
S Em t	0 11	7 25	2 17	0.50	1 10	0.00
	9.11 NG	/.55 NG	2.17	0.39	1.12 NC	0.90
CD @ 5%	INS	INS	5.58	1.52	INS	INS

NS- indicates Non significant; YB – Yelakki bale; RAJ- Rajapuri bale; NR- Nanjangud rasabale

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There was no significant difference noted in plant crop and ratoon crop of fruit firmness among different mutant lines of banana and data is presented in Table 3. In plant crop, the maximum fruit firmness was recorded in YB40Gy - 12 (18.82 N). the minimum fruit firmness was recorded in YB35Gy - 12 (6.08 N). In ratoon crop, the maximum fruit firmness was recorded in YB40Gy -11 (16.50 N) the minimum fruit firmness was recorded in RAJ EMS 0.9%- 4 (8.31 N). There was significant difference noted in plant crop in flavour acceptance among different mutant lines of banana (Table 3). The maximum flavour acceptance was recorded in YB40Gy 03 (7.82), which was on par with YB45Gy - 08 (7.72), YB35Gy - 23 (7.76), YB40Gy - 13 (7.60), RAJ40Gy -23 (7.58), YB45Gy - 03 (7.58), YB45Gy - 07 (7.48). This might be due to genetic characters, environmental conditions prevailed during the crop production, presence of more total soluble solids, superior in taste. There is no report on variations in flavour acceptance in mutant lines in banana. Hence, the results of present study revealed that there is a possibility of variations in flavour acceptance in mutants due to mutations. However, the results of the present study are in line with Sagar *et al.* (2018) who has carried out evaluation of banana genotypes for qualitative traits. These results are in line with Balesh (2020) studied evaluation of elite tissue culture raised banana varieties for growth, yield and quality parameters.

Table 3:	Quality	v parameters	of pl	lant and	l ratoon	crop o	of banana	mutant lines.
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Mutant lines	Fruit firn	nness (N)	Flavour	acceptance	Shelf life (number of days)	
Mutant miles	Plant crop	Ratoon crop	Plant crop	Ratoon crop	Plant crop	Ratoon crop
YB25GY 02	13.54	11.26	6.78	6.20	4.82	6.00
YB25GY 03	12.16	10.57	6.55	5.86	6.25	5.04
YB25Gv 04	16.18	13.50	5.68	6.24	4.75	5.39
YB25Gy 05	9.61	11.78	6.25	4.26	6.00	4.10
YB30Gy 01	14.90	9.15	7.01	5.41	5.43	4.00
YB30Gy 02	18.53	15.62	6.79	6.30	6.00	5.20
YB30Gy 03	11.57	14.33	6.63	6.29	5.33	4.90
YB30Gy 04	7.06	10.96	6.58	6.42	6 50	5.00
YB30Gy 06	13.72	12.24	6.49	5.86	7.00	5.34
YB30Gy 07	13.53	14.40	6.15	5.48	5.72	4.86
VB30Gy 08	17.65	11.56	5 40	4 67	5.11	6.00
VB30Gy 09	10.10	12.39	6.02	6.42	6.00	5.00
VB30Gy 10	16.47	15.64	7.22	6.50	7.00	5.00
VB30Cy 11	9.12	13.04	5.21	5.67	5.22	6.10
VB30Cy 15	12.04	15.00	5.21	4.86	5.56	4.00
VB35Cy 01	0.41	13.30	6.20	4.80	7.00	4.00
VP35Cy 02	12.62	15.32	5.10	5.27	5.60	4.82
VB35Cy 02	15.05	13.56	6.52	5.27	7 20	6.00
VP35Cy 03	12.16	15.30	6.00	5.50	5.50	6.20
1 B35Gy 04 VP35Cy 05	10.20	13.70	6.68	5.66	5.00	0.20
1 B35Gy 05	0.00	12.45	5.61	5.00	5.00	4.10
1 D35Gy 00 VD25C+ 07	9.90	12.43	5.01	4.32	7.20	5.01
1 b35Gy 07	0.37	10.58	5.92	4.80	7.20	0.20
1 B35Gy 09 VD25C+ 10	13.14	11.50	5.83	5.00	0.24	1.38
YB35Gy 10	7.04	13.02	7.30	5.62	4.60	5.04
VB35Gy 11 VB25Cy 12	6.09	11.60	0.03	5.05	5.45	0.00
1 D35Gy 12 VD25Cy 14	0.08	13.02	7.04	5.22	3.30	4.00
1 B35Gy 14	0.21	14.00	5.03	3.55	7.08	3.30
YB35Gy 15	9.31	10.38	5.27	4.36	0.00	7.22
YB35Gy 18	17.00	12.04	5.09	5.21	1.22	5.00
YB35Gy 20	13.14	9.80	0.45	6.02	4.00	6.25
YB35Gy 23	10.00	14.55	7.76	6.62	7.50	6.30
YB35Gy 24	14.90	10.27	1.21	6.97	7.04	5.00
YB35Gy 25	13.20	12.34	6.63	6.30	7.00	6./4
YB40Gy 01	17.55	13.01	6.25	5.74	5.00	7.00
YB40Gy 02	10.47	13.89	6.84	6.31	6.00	5.32
YB40Gy 03	12.45	15.46	/.82	7.34	7.50	6.00
Y B40Gy 04	14.41	10.04	6./3	6.56	6.28	5.00
Y B40Gy 05	10.59	12.68	6.30	6.00	6.00	7.50
Y B40Gy 06	12.74	14.79	7.00	6.58	6.50	8.00
YB40Gy 07	13.53	10.63	/.40	7.29	7.62	6.00
YB40Gy 08	15.88	10.55	6.38	6.41	6.00	5.00
YB40Gy10	15.00	12.30	6.44	5.39	4.50	6.72
YB40Gy 11	14.51	16.50	6.96	4.62	6.20	6.00
YB40Gy 12	18.82	13.24	7.10	5.28	5.00	6.00
YB40Gy 13	16.18	14.78	7.60	6.29	6.41	5.00
YB40Gy 14	10.98	11.21	7.30	5.62	8.10	6.39

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YB40Gy 15	13.04	10.07	6.83	6.29	8.00	6.30
YB45Gy 02	12.55	9.63	6.66	5.14	5.00	7.00
YB45Gy 03	15.69	11.30	7.58	5.30	6.92	6.10
YB45Gy 04	9.51	12.64	6.25	6.01	6.58	6.00
YB45Gy 05	14.31	13.27	6.51	6.22	7.00	5.00
YB45Gy 06	9.31	15.40	6.90	5.75	6.21	7.00
YB45Gy 07	10.39	11.00	7.48	6.38	7.17	6.50
YB45Gy 08	12.74	14.38	7.72	6.42	6.08	9.00
YB45Gy 09	14.61	15.69	7.12	6.31	6.52	7.00
YB45Gy 11	11.76	15.07	6.27	5.87	7.00	5.00
YB45Gy 12	13.43	14.02	6.62	5.93	6.27	6.82
YB45Gy 13	15.29	13.66	6.79	6.65	6.84	5.00
YB45Gy 15	12.55	12.04	6.42	6.30	6.92	6.00
RAJ25Gy 01	9.31	8.75	6.23	5.27	5.92	6.00
RAJ25Gy 02	11.20	9.28	6.67	6.30	6.21	7.00
RAJ30Gy 02	15.88	10.52	5.50	5.51	4.00	6.00
RAJ30Gy 03	7.06	9.96	5.01	4.86	5.33	4.82
RAJ30Gy 05	10.57	11.58	6.04	6.23	6.20	5.00
RAJ30Gy 07	11.37	10.62	6.71	5.40	5.00	6.24
RAJ30Gy 12	9.12	11.40	6.82	5.10	6.21	5.00
RAJ30Gy 13	10.39	8.94	6.30	4.43	6.67	4.00
RAJ35Gy 01	12.35	10.31	6.42	4.82	5.00	5.34
RAJ35Gy 02	8.72	10.42	7.01	5.30	6.35	6.00
RAJ35Gy 03	9.21	13.62	5.40	5.52	5.60	4.10
RAJ35Gy 06	11.24	9.71	6.30	6.00	6.05	7.00
RAJ35Gy 07	14.21	12.09	6.42	0.27	6.60 5.80	5.00
RAJ35Gy 09 DA 135Cy 10	0.92	10.27	5.26	4.20	5.00	7.00
RAJ35Gy 10 RAJ35Gy 12	7.45	9.83	7.07	7.45	6.00	5.03
RA555Gy 12 RA 140Gy 01	9.61	10.21	5 58	6 39	7.05	7.00
RAJ40Gy 02	9.12	8.96	6.21	5.92	7.00	6.00
RAJ40Gy 03	11.76	10.55	7.30	7.28	5.50	5.20
RAJ40Gy 04	10.98	10.82	6.38	5.22	6.00	4.50
RAJ40Gy 05	8.43	13.67	6.92	6.01	6.70	5.50
RAJ40Gy 08	9.82	11.40	6.87	5.37	5.50	4.00
RAJ40Gy 09	8.90	10.02	6.44	5.06	4.00	6.00
RAJ40Gy 11	10.68	13.50	7.11	5.00	5.10	5.00
RAJ40Gy 15	13.04	10.00	6.70	6.26	4.30	6.50
RAJ40Gy 22	11.78	12.80	7.36	7.24	4.00	5.31
RAJ40Gy 23	11.57	14.60	7.58	7.31	5.90	6.00
RAJ40Gy 25	12.45	11.72	6.27	6.80	6.00	8.00
RAJ45Gy 03	10.39	8.63	6.66	7.36	5.20	6.00
RAJ45Gy 04	16.18	11.08	6.12	7.21	5.38	7.00
RAJ45Gy 07	9.81	10.37	6.98	7.48	5.24	5.00
RAJ45Gy 08 PA 145Cy 10	12.60	9.50	5.20	5.00	3.05	5 30
RAJ45Gy 10 RA 145Cy 13	10.46	8.76	6.42	5 36	6.50	6.00
RAJ45Gy 15 RA 145Gy 14	12.05	9.74	6.04	7 42	5.80	6.00
RAJ45Gv 15	10.34	11.50	6.25	7.00	5.00	6.72
RAJ45Gy 19	11.57	14.30	5.42	6.32	7.00	5.00
RAJ45Gy 20	13.33	12.86	5.38	5.20	5.68	6.00
NR30Gy 06	9.12	10.38	4.96	5.03	6.00	7.54
NR30Gy 10	12.75	11.72	5.25	5.12	7.25	6.00
NR35Gy 08	13.57	9.68	6.10	4.50	7.50	6.04
NR35Gy 11	12.50	14.69	5.00	4.89	7.92	7.00
NR35Gy 12	10.57	11.36	5.41	5.65	6.45	4.05
NR35Gy 19	10.03	10.28	5.32	6.31	6.92	5.00
NK40Gy 05	12.16	12.87	6.84	6.27	7.00	6.30
NR40Gy 07	9.81	10.52	7.02	6.90 5.46	/.33	5.34
NR40Gy 09 NR40Gy 12	11.03	11.30	6.20	J.40 1.86	6.00	7.00
NR40Gy 12 NR40Cy 15	10.90	12.70	5.60	5 27	6./5	6.00
NR45Cv 01	12.56	14 50	5.00	4 29	7 20	5.00
NR45Gv 02	10.39	11.00	6.00	5.37	5.35	7.20
NR45Gv 04	9.51	12.04	6.50	5.33	6.28	5.62
NR45Gy 09	10.27	12.35	5.05	6.20	7.12	4.89
YB EMS 0.6% 1	13.72	10.26	7.28	6.80	7.58	6.00
YB EMS 0.6% 4	12.95	16.04	7.04	6.57	6.21	5.34
YB EMS 0.9% 1	10.29	13.38	7.12	6.33	5.00	8.00

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YB EMS 0.9% 2	10.56	11.50	7.34	6.92	7.25	5.76
YB EMS 0.9% 3	11.80	12.62	6.58	5.62	6.89	6.74
YB EMS 0.9% 4	13.05	12.42	6.96	5.60	6.00	5.27
YB SA 0.02% 1	13.57	14.50	5.76	5.21	5.24	4.81
YB SA 0.03% 6	15.60	13.70	7.23	4.80	6.85	5.00
YB BAP15- 1	12.53	10.68	6.39	4.36	5.00	4.27
YB BAP 20-2	10.29	14.32	6.96	6.00	6.67	6.20
RAJ EMS 0.6% 3	8.76	9.56	7.02	6.28	6.00	4.00
RAJ EMS 0.9% 1	10.40	13.66	6.52	6.39	5.64	5.00
RAJ EMS 0.9% 2	10.55	9.84	7.40	6.44	7.00	5.34
RAJ EMS 0.9% 4	11.34	8.31	6.70	7.02	6.00	5.00
RAJ SA 0.02% 1	12.46	10.04	5.22	4.50	5.46	4.73
RAJ SA 0.03% 1	9.28	9.63	4.56	4.62	6.00	5.00
RAJ SA 0.03% 2	9.69	10.00	6.72	6.24	5.62	6.00
RAJ BAP 15- 3	10.38	10.42	5.07	6.78	7.00	5.04
RAJ BAP 20 1	8.94	9.20	6.12	5.60	5.00	7.20
NR EMS 0.6% 3	12.40	11.88	7.00	5.82	6.00	5.30
NR EMS 0.9% 5	11.79	12.60	6.53	6.21	7.00	6.00
NR SA 0.02% 1	13.06	10.53	4.62	5.06	8.00	6.55
NR SA 0.03% 1	12.27	8.61	4.76	5.00	8.00	7.00
NR SA 0.03% 2	10.82	14.57	6.45	6.30	6.34	5.40
NR BAP 15-1	11.51	13.60	6.29	4.52	7.02	6.02
NR BAP20-3	10.91	12.00	6.80	5.21	6.50	7.00
YB	12.72	12.01	6.45	6.07	6.57	6.21
RAJ	12.64	12.77	6.45	6.55	6.21	6.26
NR	11.80	12.00	6.29	5.96	6.45	5.87
S.Em±	2.24	2.10	0.61	0.59	0.87	0.78
CD @ 5%	NS	NS	1.59	1.51	NS	2.02

NS- indicates Non significant; YB - Yelakki bale; RAJ- Rajapuri bale; NR- Nanjangud rasabale

RAJ EMS 0.9% - 2 (7.40), YB EMS 0.9% -2 (7.34), YB35Gy - 10 (7.38), RAJ40Gy - 22 (7.36), RAJ40Gy -03 (7.30), YB40Gy - 14 (7.30), YB EMS 0.6% -1 (7.28), YB30Gy - 10 (7.22), YB35Gy - 24 (7.21), NR40Gy 07 (7.16), YB EMS 0.9% -1 (7.12), RAJ40Gy - 11 (7.11), RAJ35Gy - 12 (7.07), YB EMS 0.6% -4 (7.04), NR40Gy - 09 (7.03), YB30Gy - 01 (7.01),. The minimum flavour acceptance was recorded in RAJ SA 0.03% - 1 (4.56).

In plant crop, among Yelakki bale mutant lines the maximum flavour acceptance compared to check (6.45) was recorded in YB40Gy - 03 (7.82). In Rajapuri bale mutant lines the maximum flavour acceptance compared to check (6.45) was recorded in RAJ40Gy -23 (7.58). In Nanjangud rasabale the maximum flavour acceptance compared to check (6.29) was recorded in NR40Gy - 09 (7.03).

Significant difference in flavour acceptance was noted in ratoon crop among different mutant lines of banana (Table 2). The maximum flavour acceptance was recorded in RAJ45Gy - 07 (7.48) which was on par with RAJ35Gy - 12 (7.45), RAJ45Gy - 14 (7.42), RAJ45Gy - 03 (7.36), YB40Gy - 03 (7.34), YB40Gy -07 (7.29), RAJ40Gy - 03 (7.28), RAJ40Gy - 22 (7.24), RAJ45Gy - 04 (7.21), RAJ EMS 0.9% - 4 (7.02), RAJ45Gy - 15 (7.00), YB35Gy - 24 (6.97), YB EMS 0.9% -2 (6.92), NR40Gy - 07 (6.90), RAJ40Gy - 25 (6.80), YB EMS 0.6%-1 (6.80), YB35Gy - 12 (6.75), YB35Gy - 10 (6.70), YB45Gy - 13 (6.65), YB35Gy -01 (6.63), YB35Gy - 23 (6.62), YB40Gy - 06 (6.58), YB EMS 0.6% - 4 (6.57), YB35Gy - 04 (6.57), YB40Gy - 04 (6.56), YB30Gy - 10 (6.50), YB40Gy - 08 (6.41), RAJ40Gy - 01 (6.39). The minimum flavour acceptance Kirankumar et al., **Biological Forum – An International Journal** 14(1): 215-223(2022)

was recorded in YB25Gy - 05 (4.26). In ratoon crop, in Yelakki bale, the maximum flavour acceptance compared to check (6.07) was recorded in YB40Gy - 03 (7.34). In Rajapuri bale, the maximum flavour acceptance compared to check (6.55) was recorded in RAJ45Gy - 07 (7.48). In Nanjangud Rasabale the maximum flavour acceptance compared to check (5.96) was recorded in NR30Gy - 06 (7.54). There was no significant difference was recorded in plant crop among different mutant lines of banana and data is presented in Table 2. In ratoon crop, the maximum shelf life was recorded in YB45Gy - 08 (9.00 days) and the minimum shelf life was recorded in RAJ40Gy - 08 (4.00 days), RAJ EMS 0.6% -3 (4.00 days), YB35Gy - 12 (4.00 days), YB30Gy - 15 (4.00 days) YB30Gy - 01 (4.00 days). Earlier, the longest green life of banana fruit was observed in irradiated Williams clone w-193/3 while lowest seen in original Williams cultivar. These results are in agreement with Ahmed (2003) who showed that irradiated Grand Naine (clone 1 and clone 2) had longer green life than that of untreated Williams. Similarly, Mohammed (2003) observed highly significant difference between Williams mutants compared to dwarf Cavendish in their fruit green life. Similarly Sales et al. (2013) reported that plants treated with 50Gy in Lakatan cultivar showed mean value for four generation showed longer shelf life (14.5 days) than control (13.15 days). This might be due to mutation induction can lead to enhance heritable variation.

### CONCLUSION

Mutation induction can lead to enhance heritable variation this might be useful for functional genomics 222

or breeding application. This can be exploited for suppressing vegetative character like plant height and improve fruit quality parameters like peel weight, shelf life and flavour acceptance. The application of gamma radiation and chemical mutagen is a promising technique to obtain good quality parameters in banana.

### FUTURE SCOPE

(i) Generation of some more mutant lines *in vitro* using physical mutagens at 40 and 45Gy.

(ii) Evaluation of lines for disease resistance by challenging with artificial inoculation of isolated culture or in the sick plot.

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