# **Short Communication**

# Genetic resources of spine gourd (*Momordica dioica* Roxb. ex Willd.): an underexplored nutritious vegetable from tribal regions of eastern India

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### Received 1 April 2010; Accepted 21 June 2010 - First published online 13 July 2010

### Abstract

Spine gourd (*Momordica dioica* Roxb. ex Willd.) is a highly nutritious and underutilized vegetable containing high amount of protein as compared to other cucurbitaceous vegetables. Twenty-six accessions of spine gourd (*M. dioica* Roxb.), collected from eastern India, were evaluated to study the diversity and other genetic parameters. The estimates of genotypic coefficient of variation and phenotypic coefficient of variation for all the characters indicated that selection can be done on the basis of phenotype alone. On the basis of cluster analysis, spine gourd accessions can be categorized into three distinct groups. The accessions of spine gourd from eastern India showed an immense variation and can be used as a potential source of germplasm for crop improvement. The results presented here on nature and magnitude of genetic variability and diversity existing in this species will provide the foundation for designing an efficient spine gourd breeding programme. As spine gourd is distributed widely almost throughout India, further collections from Peninsular and Western India are suggested for maximum diversity sampling.

Keywords: clustering; genetic variability; germplasm; Momordica dioica; morphological diversity

## Experiment

Spine gourd (*Momordica dioica* Roxb., Cucurbitaceae) is a perennial species propagated by tubers. *M. dioica* (diploid, called spine gourd) and *Momordica subangulata* subsp. *renigera* (tetraploid, called teasle gourd) were found in eastern and north-eastern India, respectively. The latter species has been misidentified as tetraploid *M. dioica* and is cultivated in West Bengal and Orissa states. Cultivation of spine gourd is less demanding and ideal for homestead gardens. The nutritional value of spine gourd fruits is superior with respect to protein, vitamin A and ascorbic acid (Naik *et al.*, 1951; Sastri, 1962). Apart from its nutritive value, spine gourd root contains aliphatic compounds (Ali and Srivastava, 1998) and the plant parts are used in traditional

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ID no.	Source	DF	ΡL	ΓΓ	TL	Π	FP	FW	FL	FD	SL	SF	SW	γP
CH 101	Jharkhand	35.70	2.32	5.10	13.32	4.73	111.13	7.62	4.25	1.85	1.43	8.18	7.56	0.79
CH 103	Orissa	34.73	5.48	7.03	11.31	9.98	83.67	21.88	4.23	2.78	3.63	15.85	9.56	1.64
CH 104	Orissa	31.00	3.04	5.92	14.88	6.47	111.40	16.67	3.55	2.37	2.07	11.05	9.57	1.78
CH 105	Orissa	38.13	4.58	9.10	13.80	6.62	90.73	15.73	5.03	2.35	3.38	19.16	7.67	1.20
CH 108	West Bengal	33.93	2.43	5.98	10.67	5.00	96.27	14.44	2.85	2.38	3.00	24.57	8.43	1.11
CH 109	West Bengal	44.07	3.55	7.20	14.18	8.87	91.80	13.44	3.60	2.32	3.62	18.92	9.64	0.96
CH 110	West Bengal	38.60	2.93	5.70	11.52	6.48	95.33	11.27	3.57	2.33	2.30	20.82	9.34	0.94
CH 111	Orissa	36.73	2.55	5.78	7.73	5.20	114.53	10.84	3.00	2.03	2.40	23.57	9.36	1.12
CH 112	Orissa	33.13	3.32	7.20	10.78	5.90	125.20	15.24	2.82	2.49	2.42	19.65	9.82	1.75
CH 113	Jharkhand	37.53	3.27	6.77	12.56	7.45	133.73	6.44	3.83	1.65	2.37	8.54	12.32	0.76
CH 114	Orissa	38.40	2.32	5.77	12.64	4.80	128.67	15.58	5.15	2.67	2.62	14.00	12.22	1.82
CH 116	Orissa	36.27	6.37	7.02	13.63	6.23	151.33	13.15	3.76	2.78	3.38	28.08	8.55	1.79
CH 117	Orissa	34.53	4.32	7.56	13.06	5.93	143.60	16.11	4.82	2.75	2.32	29.73	8.18	2.08
CH 119	Orissa	39.07	2.15	8.19	9.74	6.58	156.20	13.12	3.48	2.48	1.93	26.22	6.51	1.78
CH 1	Orissa	34.13	2.52	5.72	10.25	5.32	160.87	6.69	2.58	1.83	2.43	8.99	7.61	0.92
CH 8	Orissa	37.73	2.97	8.11	11.73	7.16	127.87	15.20	4.18	2.43	4.23	28.18	9.40	1.52
CH 18	Orissa	37.67	1.70	4.83	12.65	6.79	110.53	14.75	3.45	2.47	2.12	6.75	6.49	1.25
CH 26	Orissa	36.33	4.95	7.89	11.81	7.48	100.07	10.97	2.91	2.41	2.46	11.84	4.74	0.97
CH 27	Orissa	39.67	3.01	6.44	11.30	5.83	116.00	10.77	3.10	2.18	3.02	10.74	7.41	1.16
CH 28	Orissa	33.80	4.53	7.46	11.02	6.62	213.67	11.01	3.01	2.36	2.47	10.46	8.91	1.98
CH 35	Orissa	35.20	2.85	7.39	12.39	4.79	132.13	10.50	3.48	1.80	2.26	16.12	6.83	1.07
CH 52	Orissa	38.00	3.95	9.43	10.15	7.45	116.33	13.37	3.51	2.37	1.83	8.67	9.01	1.24
CH 54	Orissa	41.87	2.89	6.43	12.67	6.28	106.00	10.26	3.76	2.15	2.67	14.44	8.63	0.99
CH 59	Orissa	43.20	5.12	7.66	14.22	8.17	98.93	17.79	3.84	2.07	1.53	10.62	9.89	1.37
CH 60	Orissa	39.93	3.20	8.80	11.17	5.83	107.73	15.25	3.85	2.18	3.17	16.88	10.14	1.31
CH 80	Orissa	40.53	3.78	6.80	9.04	5.08	93.27	20.14	4.39	2.49	2.32	19.21	9.38	1.46
Mean		37.16	3.46	6.97	11.85	6.43	119.89	13.39	3.67	2.31	2.59	16.12	8.74	1.34
CD (5%)		2.07	0.46	0.57	1.58	0.98	8.98	1.41	0.49	0.18	0.34	1.89	0.62	0.12
CV (%)		3.40	8.07	5.01	8.13	9.28	4.57	6.40	8.22	4.76	8.04	7.13	4.32	5.33
GCV (%)		8.4	32.9	16.9	13.6	19.4	23.5	27.8	17.9	12.7	25.5	40.1	18.8	28.6
PCV (%)		9.1	33.8	17.6	15.8	21.5	24.0	28.5	19.1	13.6	26.7	40.7	19.3	29.1
DF, days to	o flowering; PL, p	etiole lengt	, (cm); LL	leaf length	ר (cm); TL,	tendril ler	ngth (cm); IL	., internode	e length (cr	n); FP, nun	nber of fru	its/plant; F\	N, fruit we	ight (g);
FL, fruit le	ngth (cm); FD, tru	it diameter	(cm); SL, ti	uit stalk le	ingth (cm);	SF, numbe	er of seeds/tr	uit; SW, 10	J0-seed we	eight (g); YH	<sup>2</sup> , yield/pla	nt (g); CD,	critical dif	erence;
CV, coeffic	ient of variation; (	<b>GCV</b> , genoty	/pic coeffic	ient of vari	iation; PCV	, phenotyp	oic coefficier	nt of variati	on.					DI

Momordica divica accessions from India used for morphological classification and their quantitative characteristics Table 1.

https://doi.org/10.1017/S1479262110000237 Published online by Cambridge University Press

medicines. Extensive use of its tubers in various ethno-medicinal formulations by different tribal communities of Kerala has been reported (Joseph and Antony, 2008). In addition, the seed kernel oil of spine gourd is used as such or as a source of drying oil in the paint and varnish industry (Chakrabarty *et al.*, 1956).

The objective of the present study is to characterize and evaluate the yield and other horticultural traits of a set of accessions of *Momordica* spp. maintained at the Central Horticultural Experiment Station, Bhubaneswar, Orissa.

All 40 accessions were planted at a spacing of  $2 \times 1$  m in a randomized block design with three replications. The plant population in the experimental field had female and male plants at the ratio of 10:1 to ensure effective pollination. The accessions were evaluated for 12 quantitative, 10 qualitative and a phenological character. Mean, ANOVA, coefficient of variation (CV) and cluster analysis were performed using INDOSTAT statistical package.

#### Discussion

The accessions showed a considerable level of variability (supplementary Fig. S1, available online only at http://journals.cambridge.org) for qualitative traits such as leaf

lobing (entire to deeply lobed), leaf colour (light green to dark green), blossom-end rostration of fruit (faint to appreciable length), fruit shape (globular to elliptic), fruit colour at edible maturity (whitish green to dark green) and ripe fruit colour (yellow to red). The diversity found in fruit colour and shape would be helpful for the selection of genotypes with desired quality. In addition, the accessions displaying contrast characters can be used to study the genetics of trait expression. Among the accessions studied, the earliest flowering was observed in CHSG 104 (31 d), which is a favourable trait to take advantage of early harvest and profitable market avenue owing to its high demand.

A wide range of variation (Table 1) was observed for petiole length (1.7-6.4 cm), leaf length (4.8-9.4 cm), internode length (4.7-9.9 cm), tendril length (7.7-14.2 cm), number of fruits per plant (83.7-213.7) and fruit weight (6.4-21.9 g). Accession CHSG 117 recorded the highest yield (2.1 kg/plant) followed by CHSG 28 (1.9 kg/plant), while other accessions exhibited a yield range of 0.8-1.8 kg/plant. The high CV observed for vegetative characters like internode length (9.3%), tendril length (8.1%), petiole length (8.1%), fruit length (8.2%), fruit stalk length (8%) and number of seeds per fruit (7.1%) could be due to heterogeneity existing among the accessions. The estimates of genotypic coefficient of



Fig. 1. Ward's minimum variance dendrogram of 26 accessions of spine gourd.

variation and phenotypic coefficient of variation for all the characters correspond closely, which indicate that the variability existing in them was mainly due to their genetic makeup, and thus selection on the basis of phenotype alone should be effective. The high level of phenotypic diversity observed among the accessions in this study is in agreement with the earlier findings (Ram *et al.*, 2001; Bharathi *et al.*, 2006; Panchbhai *et al.*, 2006) for different characters in other sets of germplasm.

Following Ward's clustering technique, the accessions were grouped into three main clusters (I, II, III) comprising 6, 8 and 12 accessions, respectively (Fig. 1). The maximum intercluster distance 5.5 was observed between clusters I and III. Cluster I could be characterized with genotypes early in flowering (except CHSG 119), more number of fruits per plant (>100), number of seeds per fruit (>10) and greater yield per plant (>1.8 kg/plant). Cluster II comprised the accessions having high individual fruit weight and 100-seed weight. The accessions grouped in cluster III recorded lower values for most of the traits studied.

This prized vegetable presently restricted to the areas of its natural distribution has the scope for extended cultivation in the context of climatic change, physiographic diversity and adaptation. Its commercial cultivation is meagre due to a lack of availability of improved varieties, seed dormancy, low multiplication rate of tubers, dormancy of tubers and unpredictable sex ratio in seedling progeny (Ali *et al.*, 1991). The results presented here on the nature and magnitude of genetic variability and diversity existing in this species will provide the foundation for designing an efficient spine gourd breeding programme. The present study also highlights the importance of spine gourd germplasm from eastern India for maintaining a considerable range of

diversity for most of the important traits studied. As spine gourd is distributed widely almost throughout India, further collections from Peninsular and Western India are required for maximum diversity sampling.

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