### DEVELOPMENT AND EVALUATION OF HYBRIDS IN IVY GOURD (Coccinia grandis L.Voigt.)

#### R. PRABU

## Thesis submitted in partial fulfilment of the requirement for the degree of

### Master of Science in Agriculture

Faculty of Agriculture Kerala Agricultural University, Thrissur

2008



Department of Plant Breeding and Genetics COLLEGE OF AGRICULTURE VELLAYANI, THIRUVANANTHAPURAM-695 522

#### CERTIFICATE

Certified that this thesis entitled 'Development and evaluation of hybrids in ivy gourd (*Coccinia grandis* L. Voigt)' is a record of research work done independently by Mr. R. Prabu (2006-11-113) under my guidance and supervision and that it has not previously formed the basis for the award of any degree, fellowship or associateship to his.

Vellayani, 23-09-2008.

**Dr. D. WILSON** (Chairman, Advisory Committee) Professor, Department of Plant Breeding and Genetics, College of Agriculture, Vellayani, Thiruvananthapuram.

#### DECLARATION

I hereby declare that this thesis entitled 'Development and evaluation of hybrids in ivy gourd (*Coccinia grandis* L. Voigt)' is a bonafide record of research work done by me during the course of research and that the thesis has not previously formed the basis for the award of any degree, diploma, associateship, fellowship or other similar title, of any other university or society.

Vellayani, 23-09-2008.

(2006 - 11 - 113)

#### CERTIFICATE

Certified that this thesis entitled 'Development and evaluation of hybrids in ivy gourd (*Coccinia grandis* L. Voigt)' is a record of research work done independently by Mr. R. Prabu (2006-11-113) under my guidance and supervision and that it has not previously formed the basis for the award of any degree, fellowship or associateship to his.

Vellayani, 23-09-2008.

**Dr. D. WILSON** (Chairman, Advisory Committee) Professor, Department of Plant Breeding and Genetics, College of Agriculture, Vellayani, Thiruvananthapuram.

#### Approved by

#### Chairman

Dr. D. WILSON Professor, Department of Plant Breeding and Genetics, College of Agriculture, Vellayani, Thiruvananthapuram - 695 522.

#### Members

#### Dr. P. MANJU

Professor and Head, Department of Plant Breeding and Genetics, College of Agriculture, Vellayani, Thiruvananthapuram-695522.

**Dr. SUNNY K.OOMMEN** Professor, Department of Plant Breeding and Genetics, College of Agriculture, Vellayani, Thiruvananthapuram-695 522.

#### Dr. VIJAYARAGHAVAKUMAR

Professor, Department of Agricultural Statistics, College of Agriculture, Vellayani Thiruvananthapuram-695522.

12-9-08



External Examiner

23109 S. MURUGAN Professor Dept. of Agri Arnem alan Agni Borton

# **Dedicated** to My dear chairman **Dr.D.Wilson**, **Professor** & his family

#### ACKNOWLEDGEMENT

To Whom be glory for ever and ever. Amen.

It is a great rejoice and privilege for me to express my profound sense of gratitude to Dr. D. Wilson, Professor, Department of Plant Breeding and Genetics and chairman of my advisory committee for his unforgettable guidance, infinity encouragement, care and love rendered during the research period which gave me a feel of comfortness and enabled me to complete the thesis of course the M.Sc, programme successfully.

My sincere thanks to Dr. P. Manju, Professor and Head, Department of Plant Breeding and Genetics and member of my advisory committee for her critical suggestions and timely help during the course of this research.

My profound gratitude to thanks to Dr. Sunny K.Oommen, Professor, Department of Plant Breeding and Genetics and member of my advisory committee for his critical suggestions and valuable advice.

I wish to place on record my sincere thanks to Dr. Vijayaraghavakumar, Professor, Department of Agricultural Statistics and member of my advisory committee for the kind help rendered during the statistical analysis and interpretation, valuable suggestions and thorough scrutiny of the manuscript. I thank Dr. Roy Stephen, Associate Professor and Head and Dr. Manju, Associate Professor, Department of Plant Physiology for their kind help in providing the lab facilities.

I happy to thanks to Dr.K.Arya, Professor and Smt. Seeja, Assistant Professor, Department of Plant Breeding and Genetics for their kind suggestions and technical guidance for thesis work. I also thank all teaching and non-teaching staff of Department of Plant Breeding and Genetics for their co-operation all through the period of the study.

I express my profound gratitude to Dr. N. P. Kumari Sushama, Professor, Department of Agricultural Extension for her timely and sincere help and support provided during the critical stages of my life. My immortal gratitude to madam E. Nirmala Wilson and Nancy Wilson for their loving involvement and care during the entire period of my M.Sc. programme.

I find special pleasure in placing my wholehearted thanks to my classmates Divya Krishnan, Kumanan, Selvaraju, Kumaran, Bavya, Vidya, Saritha, Vimersh, Ponsubrya, Renjani and Rekha and all my seniors Jithesh, Beena Thomas, MadhuKumar and junior Jeyarammegowda for their kind support and affection provided during the various stages of my work. Thanks are due to Rani akka, Sajitha akka, Ligi chechi, Sri ram, Johny, Neelima, Majjusha, Ramlath, Vinitha, jungiah, Princy John, Reddy, Tamil vel, Srikanth, Amstrong Enose A, Saji stephen and Anil chetan for their invaluable help from the very beginning of the PG programme. My special thanks to Ananthi akka and Aravind.J for their technical help for thesis work.

I express my respect and thanks to Dr. Sam, Good Sam Ministries and the brothers for their financial support and care give to me at all times. Once again I thank to all friends who helped me for this M.Sc, programme. I sincerely acknowledge KAU, Thrissur for awarding the KAU Fellowship.

The beginning and end I express my eternal love and affection to most lovable Amma for her patience, prayer and care with me in whole life.

Above all, I thank God Almighty for being with me always.

Prabu.R

#### CONTENTS

.

.

	Page No.
1. INTRODUCTION	1
2. REVIEW OF LITERATURE	3
3. MATERIALS AND METHODS	22
4. RESULTS	31
5. DISCUSSION	66
6. SUMMARY	77
7. REFERENCES	୫୮
ABSTRACT	

### LIST OF TABLES

Table No.	Title	Page No.
1	Female and male parents used for the experiments	23
2	Mean square values of eight characters in 12 families of ivy gourd	33
3	Mean values of eight characters in twelve $F_1$ families of ivy gourd	34
4	Mean values of progenies for different characters	46-50
5	Genetic components of variation	54
6	Phenotypic correlation coefficient	56
7	Genotypic correlation coefficient	57
8	Environmental correlation coefficient	58
9	Direct and indirect effects of path analysis	6 <u>0</u>
10	Standard heterosis (%) for eight characters of 12 families of ivy gourd	65

.

.

### LIST OF FIGURES

SI. No.	Title	Between pages
1	Phenotypic and genotypic coefficient of variation	54-55
2	Heritability and genetic gain	54-55
3	Path diagram	60-61
4	Standard heterosis	65-66

## LIST OF PLATES

Plate No.	Title	Between pages
1	Morphological characters of the female parents	22 - 23
2	Morphological characters of the male parents	22-23
3	Hybridization technique	22 - 23
4	Field view	23-24
5	Variation in fruit characters among the hybrid progenies	35-36
6	Variation in fruit characters among the hybrid progenies continued	35 - 36
7	Variation in fruit characters among the hybrid progenies continued	35-36
8	Variation in fruit characters among the hybrid progenies continued	.35-36
9	Variation in fruit characters among the hybrid progenies continued	35-36
10	Variation in fruit characters among the hybrid progenies continued	35-36

.

## **INTRODUCTION**

#### 1. INTRODUCTION

. :

Cucurbits are a highly evolutionary group and many of them are known to be vegetables for human consumption from the remote age of civilization. The ancient Indian civilization was basically dependent upon and intimately related with the forests and flora. Sanskrit prose, scriptures, epics, poetical works such as "Vedas", "Upanishads", "Ramayana", "Mahabharata", "Brahmanas", "Puranas" and medical treatises dating back to the ages before Christian era mention several kinds of cucurbits.

India is the home of ivy gourd (*Coccinia grandis* L. Voigt.) having primary gene centre of diversification (Seshadri and More, 2004). Ivy gourd otherwise called gentleman's toes (English), Kovakkai (Malayalam), and Kovaikkai (Tamil) is a common vegetable cultivated in South East Asian countries. It is widely distributed throughout the tropics and found in both wild and cultivated state in the plains of India. There are two distinct types sweet and bitter. Several cultivars of the sweet type are extensively cultivated in different parts of Kerala for its edible fruits.

Ivy gourd is dioecious having male and female plants (Veeraragavathatham et al., 1998). Distinct sex forms in *Coccinia* are due to the presence of characteristically distinct sex chromosomes. Its diploid chromosome number is 2n = 24. Gynodioeceious forms consisting of female and hermaphrodite flowers have also been observed. Maleness is expressed by the influence of 'Y' chromosome. The pericarp in *Coccinia grandis* develops from homogeneous ovary wall. It is distinguished into outer, middle and inner regions. There is an increase in the number of cell layers and cell size with some differentiation of the cells during the development of the fruit from the ovary (Ilyas and Muhammad, 1995).

Tender fruits of ivy gourd contain 1-2 per cent protein (Sachan and Chaundawat, 1985). Being rich in beta-carotene it is a good source of dietary vitamin A (Wasantwisut and Viriyapanich, 2003). It has considerable amount of protein, fibre, and modest amount of calcium. The tender fruits and shoots of ivy gourd are eaten raw or cooked while roots, stem and leaves are used as ingredients of medicines for the treatment of skin diseases and bronchitis (Veeraragavathatham et al., 1998). Dhanabal et al. (2004) reported that ivy gourd has been widely used in the traditional treatment of *Diabetes mellitus* disease.

. \*

Ivy gourd is highly cross pollinated in nature. However, due to the non availability androecious plants, viable seeds are not produced. Even though ivy gourd is a potential vegetable crop for the tropics, crop improvement programmes are scanty. However variety the 'Sulabha' released by Kerala Agricultural University is cultivated in many parts of Kerala. Being a clonally propagated crop there is scope for the development of hybrid clones in ivy gourd. Androecious lines having variations in leaf and floral characters have also been identified from different regions but no reports are available on the development of hybrid clones in ivy gourd.

The success of breeding programme depends on the availability of genetic variation in a population. Information on variability, components of phenotypic and genotypic variation, correlation and path analysis are the basic requirements for crop improvement. The present study was undertaken with the following objectives.

- (i) Estimation of genetic variability among the hybrid population.
- (ii) Estimation of genetic variability within the hybrid progenies.
- (iii) Correlation and path analysis to estimate direct and indirect effects of yield contributing characters on yield.
- (iv) Estimation of standard heterosis using the check variety'Sulabha'.
- (v) Identification of superior hybrids over check variety.

## **REVIEW OF LITERATURE**

1

•

.

•

.

•

.

#### 2. REVIEW OF LITERATURE

Estimation of genetic parameters is the primary requirement to formulate efficient breeding strategy in any crop. Genetic studies in ivy gourd are meager. Attempts on hybridization in this crop have also not yet been reported. In this study attempts were made to develop hybrids and estimate their genetic variation between and within families. Hence variability studies reported in ivy gourd and other cucurbitaceous crop are reviewed in a crop wise manner in this chapter.

#### 2.1. Genetic parameters

#### 2.1.1. Variance and coefficient of variation

The effectiveness of selection in crop improvement programmes depends on the extent of genetic variability present in the population. The variability present in the plant population is of three types viz., phenotypic, genotypic and environmental. Of these the genetic variance can be further partitioned into additive, dominance and epistatic components variance. The genetic parameters such as phenotypic coefficient of variation (PCV), genotypic coefficient of variation (GCV), heritability ( $h^2$ ) and genetic advance (GA) serves as a basis for selection (Johnson et al. 1955).

#### Ivy gourd

Raju and Peter (1995) reported considerable variability in fruit length, fruit weight and fruit size. Joseph (1999) observed a wide range of variation for days for first flower opening, fruit length, fruit weight, fruits per plant and yield per plant. The PCV and GCV were high for primary branches and fruit yield. Sarnaik et al. (1999) evaluated 35 genotypes of ivy gourd and reported wide range of variation for fruit diameter, number of fruits and yield per plant. Varghese (2003) reported maximum variability for yield per plant followed by flowers per plant in ivy gourd. Characters viz., days to first flowering, fruit weight, fruit size and vitamin C registered a variability of less than 20 per cent both at genotypic and phenotypic levels.

Suresh (2004) recorded high PCV and GCV for fruits per plant, total number of harvest and yield per plant in ivy gourd. Wilson et al. (2006) recorded high PCV and GCV of 65.58 and 64.17 per cent respectively for yield per plant. The least PCV and GCV were observed for fruit girth.

#### Cucumber

、 '

Rao (1988) reported that in  $F_1$  progenies of cucumber the GCV values ranged from 5.14 to 73.35 per cent while PCV values ranged from 8.52 to 80.13 per cent. Abusaleha and Dutta (1990) studied 75 pure genotypes of cucumber and observed high magnitude of genotypic and phenotypic variance for all the characters studied. Rastogi and Deep (1990) recorded higher PCV and GCV for fruit yield per plant and fruit weight and the lowest for days to fruit maturity in cucumber .

In cucumber, Gayathri (1997) reported highest PCV (95.8 per cent) and GCV (92.9 per cent) for yield per plant and lowest PCV (13.6 per cent) for days to first fruit harvest and lowest GCV (11.2 per cent) for days to first flower opening.

#### Musk melon

Swamy et al. (1985) performed variability studies in muskmelon and they found highest GCV and PCV for marketable yield per plant followed by total yield per plant and average fruit weight. Chacko (1992) observed moderate to high GCV for yield per plant in a study for evaluation of dessert type musk melon. In culinary melon, Kandaswamy (2004) observed highest PCV and GCV for yield per plant followed by average fruit weight and fruits per plant. Parmar and Tarsen Lal (2005) recorded highest genotypic (91.09 per cent) and phenotypic (93.32 per cent) coefficients of variation for number of fruits per vine. Prasad et al. (2002) in their studies with 32 inbred lines of musk melon revealed high genetic variation for node at which first male flower appeared and days to first flower opening. Rakhi and Rajamony (2005) conducted genetic studies in land races of culinary melon and found highest PCV and GCV for fruit yield per plant followed by fruit weight, fruits per plant and keeping quality.

#### Bitter gourd

The highest GCV for yield per plant and the lowest for days to first female flower opening were reported by Mangal et al. (1981) in their studies with 21 varieties of bitter gourd. Suribabu et al. (1986) anlaysed six lines of bitter gourd and found that GCV was moderate to high for all the characters except fruits per plant and percentage of fruit set. Indiresh (1982) studied 24 lines of bitter gourd and observed high GCV for fresh fruit weight, yield per plant and fruit size. While studying the genetic variability among 50 genotypes of bitter gourd, Vahab (1989) observed significant differences for 18 characters. The high PCV were observed for fruit length, yield and fruits per plant, while earliness exhibited low PCV. The GCV was of high magnitude for majority of characters. Jaiswal et al. (1990) found that fruits of various cultivars varied in colour, size, and weight, contents of protein, carbohydrates, vitamin A, vitamin C, acidity and total phenol.

In bitter gourd, Thakur et al. (1994) reported significant variability for vine length, branches per plant, root - shoot ratio and flowers per plant. Study conducted by Rajput et al. (1996) in bitter gourd reported large variations for yield and its components at the phenotypic and genotypic levels. Manju and Wilson (2002) reported the highest phenotypic and genotypic coefficient of variation for fruit yield per plant followed by number of seeds per fruit in their studies using 53 genotypes of diverse origin.

Bhave et al. (2003) reported in bitter gourd that the PCV was higher than the GCV for all the characters studied. The PCV and GCV were higher for total fruit yield per vine. In a study with 46 genotypes of bitter gourd, Raja et al. (2007) observed that genotypic and phenotypic coefficients of variation for different characters ranged from 6.375 to 37.251 and 8.373 to 38.630 per cent respectively. The highest PCV and GCV were observed for number of female flowers per plant followed by fruit weight.

#### Bottle gourd

, **'** 

Tyagi (1972) conducted a study in bottle gourd using 25 inbreds and noted that fruits per plant exhibited the highest GCV (48.26 per cent) followed by fruit length and girth. Kumar et al. (1999) noticed maximum GCV for fruits per plant followed by yield per plant. Mathew et al. (1999) in bottle gourd observed significant differences for days to first female flower opening, sex ratio, fruits per plant, fruit yield per plant and fruit size. Bisognin and Storck (2000) reported significant estimates for genetic variance for fruit diameter and neck diameter in bottle gourd. Mathew et al. (2001) evaluated 28 accessions and recorded high coefficients of variation for fruits per plant (32.50 per cent), fruit length (24.8 per cent) and fruit weight (22.40 per cent).

#### Ridge gourd

High PCV and GCV for fruits per plant, fruit weight and yield per plant were reported by Varalakshmi et al. (1995). In a study in ridge gourd, Anitha (1998) reported high PCV and GCV for number of days to first harvest, fruits per plant, yield per plant, average fruit weight and fruit size. In ridge gourd, Choudhury and Sarma (2002) evaluated 12 cultivars and recorded high phenotypic and genotypic coefficients of variation for fruit weight and yield per hectare. They showed that the phenotypic coefficient of variation was greater than genotypic coefficient of variation for all the characters. Singh et al. (2002) in an experiment with 80 genetically diverse genotypes of ridge gourd observed that PCV and GCV were high for fruits per plant, fruit weight and yield per plant. Karuppaiah and Kavitha (2002) observed GCV values ranging from 13.36 to 33.77 per cent for different characters. High GCV was observed for number of female flowers per plant, yield per plant, and number of fruits per plant. Snake gourd

Varghese (1991) and Varghese and Rajan (1993) observed that, the PCV and GCV were the highest for yield per plant, fruits per plant, fruit length, and average fruit weight while days to female flower opening and fruit picking had low estimates of PCV and GCV. Mathew (1999) reported the highest PCV and GCV for mean fruit weight and the lowest PCV and GCV for flesh thickness. Radhika (1999) found that in snake gourd the PCV values ranged from 5.63 to 21.63 per cent. PCV for yield per plant was the highest followed by fruits per plant. Ashok (2000) found wide variation in seed growth and yield characters in a study of character association of seeds with plant morphology in snake gourd. The GCV was high for fruit yield, number of fruits per plant and fruit length in snake gourd (Rahman et al., 2002).

#### Watermelon

Variability pattern of 48 water melon inbred lines were studied by Prasad et al. (2002) and observed high PCV values of number of fruits, node at which female flower appeared and days to female flower opening,.

#### Ash gourd

George (1981) conducted biochemical studies in ash gourd and reported significant differences for protein content and vitamin C in fruits, fruit yield, fruit weight, fruits per plant and fruit size and the variability was low for days to first female flower anthesis. Lovely and Radhadevi (2007) noted in ash gourd high PCV and GCV for mean fruit weight, fruit yield per plant and fruits per plant which indicates high genetic variability. They also reported comparatively low coefficient of variation for days to first female flower.

#### Pumpkin

Mohanty and Mishra (1999) observed high GCV and PCV for yield and fruits per plant in pumpkin and PCV was greater than GCV for all the traits. Sirohi and Yayasani (2001) revealed variation in the total soluble solids and carotenoid content of eight genetically diverse lines and varieties and their  $F_1$  hybrids of pumpkin. Lakshmi et al. (2002) recorded high PCV GCV for days to first female flowering in pumpkin.

#### 2.1.2. Heritability and genetic advance

Heritability and genetic advance are important selection parameters. The ratio of genotypic variance to the phenotypic variance is known as heritability. The difference between the mean phenotypic value of the progeny of selected plants and the base or parental population is called genetic advance (GA). High heritability means that the character is least influenced by environment. High genetic advance shows that the character is governed by additive genes and low genetic advance shows the existence of non additive gene action.

#### Ivy gourd

In ivy gourd, Joseph (1999) observed high heritability for vine length, primary branches per plant, fruit field per plant and fruits per vine. Varghese (2003) recorded high heritability and genetic advance for all the characters studied, except number of days for first flower opening which exhibited high heritability and low genetic advance in ivy gourd.

Suresh (2004) observed high heritability along with high GA for all the characters in ivy gourd. The range of heritability was 98.83 to 63.02. While studying the genetic variability of 61 clones of ivy gourd, Wilson et al. (2006) observed the highest heritability for days to first harvest (99.62 per cent) followed by days to first flowering (99.19 per cent). Expected genetic advance as percentage of mean was the maximum for days to first harvest (36.23 per cent) and the lowest was recorded for fruit girth (0.92 per cent).

#### Cucumber

Choudhary et al. (1985) reported significant genotypic variance for several yield components in cucumber. Yield per vine and fruits per vine had high GA along with high heritability. High heritability and low GA has been reported for days to first female flower appearance and fruit length. Prasunna and Rao (1988) reported high heritability for fruits per vine and average fruit weight in cucumber.

Mariappan and Pappiah (1990) in their studies on 45 diverse genotypes of cucumber reported high heritability along with high GA for fruit girth. Study with 23 genotypes of cucumber by Prasad and Singh (1992) revealed that the heritability estimates ranged from 0.02 per cent for fruits per plant to 48 per cent for fruit length. High heritability coupled with high GA was observed for fruit length, fruit breadth and fruit weight. High heritability and high GA for more than 12 growth and yield attributes were also observed in cucumber (Prasad ad Singh 1994).

Wehner and Cramer (1996) reported low to moderate heritability for fruit yield, earliness and quality. Paiva and Paiva (1997) derived information on variation and heritability from 36 half sib progenies in cucumber and reported that heritability was the lowest for fruit number and highest for number of fruits. Gayathri (1997) reported that yield per plant, fruits per plant, average fruit weight and node to first female flower had the highest GCV with high heritability and genetic advance. Kanwar et al. (2003) evaluated 26 genotypes of cucumber and observed high heritability estimates associated with high genetic gain for sex ratio, yield per plant and node of first female flower.

#### Bitter gourd

Vahab (1989) observed high heritability coupled with high genetic gain for fruit weight, yield and fruits per plant in bitter gourd. Choudhary et al. (1991) also reported high estimates of heritability and genetic advance for yield per plant and fruits per plant. Rajput et al. (1995) reported high heritability in bitter gourd for almost all the yield and related characters such as fruits per plant, fruit weight and fruit size. Iswaraprasad (2000) recorded high heritability for days to first female flower, days to first fruit harvest, fruits per plant, mean weight of fruit and fruit yield per plant in a study using seven parents and 21 hybrids of bitter gourd.

Manju and Wilson (2002) observed high heritability for 100 seed weight, weight of fruits, days to first male flower anthesis and fruit length. They also found maximum genetic advance for number of seeds per fruit followed by weight of fruit, fruit length and yield. All characters excluding days to first fruit harvest showed high estimates of genetic advance. In bitter gourd, Raja et al. (2007) reported high heritability coupled with greater GA for fruit weight and yield per plot.

#### Ridge gourd

In ridge gourd, Varalakshmi et al. (1995) observed high heritability values for fruit weight, days to first female flower, fruit length and fruits per plant and low heritability for fruit diameter. Anitha (1998) has reported high heritability along with high genetic advance for fruits per plant, yield per plant and days to first female flower opening. Days to first harvest had high heritability but low genetic advance.

High heritability and high GA were recorded for fruit weight and yield per hectare by Choudhury and Sarma (2002) in 12 cultivars of ridge gourd. Singh et al. (2002) in a study with 80 genotypes of ridge gourd found high heritability coupled with high genetic advance for appearance of first male and female flower, fruits per plant, fruit length, fruit weight and yield per plant. Julie Mole et al. (2002) estimated genetic parameters in 53 accessions of ridge gourd and reported high heritability along with high genetic gain for vine length, sex ratio, number of fruits per plant, length of fruit and seeds per fruit. Days to first female flowering and days to first harvest though had high heritability values, showed only low genetic gain. The number fruits per plant had the highest positive and significant correlation with yield. Karuppaiah and Kavitha (2002) observed high heritability for days to first female flowering, number of male flowers per plant, yield per plant, number of female flowers per plant, and flesh thickness. High heritability along with high genetic advance was observed for number of female flowers per plant yield per plant, number of fruits per plant and flesh thickness.

#### Snake gourd

Varghese (1991) in snake gourd observed high heritability coupled with high genetic gains for flowers per plant. Varghese and Rajan (1993) noticed high magnitude of heritability and GA for fruits per plant, while yield per plant, fruit length and days to first harvest showed high heritability coupled with low genetic gain. Mathew and Khader (1999) reported that mean fruit weight, fruit length and fruit yield per plant had high heritability coupled with high GA.

In snake gourd, Radhika (1999) reported the highest and the lowest values of heritability for days to first female flower and vine length respectively. High heritability along with high genetic advance was noticed for days to first female flower opening, fruit yield per plant and fruits per plant. High to moderate heritability as well as genetic advance was observed for fruit yield, number of fruits per plant, fruit length and average fruit weight in snake gourd (Rahman et al., 2002).

#### Bottle gourd

In bottle gourd, Kumar et al. (1999) observed high heritability for all the character studied and high genetic advance was recorded for fruit yield per plant followed by number of fruits per plant and number of branches per plant. Mathew (1999) reported high heritability and genetic advance for fruit length and fruit girth.

#### Pointed gourd

Sarkar et al. (1990) reported high heritability and low GA for fruit diameter in pointed gourd. Singh et al. (1992) reported high heritability for all the characters studied. High heritability with high GA was observed for yield and fruits per plant.

#### Pumpkin

In pumpkin, Mohanty and Mishra (1999) reported moderate heritability with high GA for yield per plant. Days to first anthesis showed moderate to high heritability accompanied by low genetic advance. Bindu et al. (2000) recorded high heritability and high GA for fruit length, fruit weight, fruits per plant and yield per hectare through the evaluation of 24 genotypes of pumpkin. In the study of 19 pumpkin genotypes, Mohanty (2000) observed high heritability and high GA as per cent of mean for the characters fruit weight, fruits per plant and yield per plant and high heritability alone for days to anthesis of first female flower.

#### Melon

Chacko (1992) evaluated desert type of musk melon and reported high heritability with high GA for yield per vine. Deepthy (2000) observed high heritability and genetic advance for yield per plant, mean fruit weight, and fruits per plant in melon. Parmar and Tarsem Lal (2005) found that heritability estimates in the broad sense were very high (>75 per cent) for all the characters except node at which the first female flower opens. Number of fruits per vine showed the highest genetic advance. High heritability along with high GA as percentage of mean was observed for number of fruits per vine. High heritability coupled with high genetic advance was reported by Rakhi and Rajamony (2005) for fruit length and fruit weight in land races of culinary melon. Zalapa et al. (2006) reported that narrow sense heritability was 0.62 for days to anthesis, 0.70 for fruit weight per plant and 0.79 for average weight per fruit in melon.

#### Ash gourd

Lovely and Radhadevi (2007) observed that fruits per plant, mean fruit weight, fruit yield per plant, fruit length and fruit girth had high heritability coupled with high GA indicating additive gene action.

#### Water melon

Prasad et al. (2002) and observed high heritability coupled with high genetic advance for number of fruits, node number, vine length, and yield per plot.

#### 2.2 Correlation Analysis

Selection is the basis for crop improvement in order to increase yield. Several other component characters contributing towards yield make it the subject of a distinct study. Correlation studies would facilitate effective selection for improvement of one or many yield contributing components. Study of relationship of yield with other traits is of immense help in any crop improvement programme.

#### Ivy gourd

In ivy gourd, Joseph (1999) reported that fruits per plant, average fruit weight, fruit girth and fruit length showed significant positive correlation with yield. Highest correlation with yield was recorded for fruits per plant. Yield per plant showed positive and significant correlation with fruits per plant and size of fruits at genotypic and phenotypic levels in ivy gourd (Sarnaik et al. 1999).

Varghese (2003) reported that in ivy gourd yield per plant exhibited positive association with all the characters except number of days for flowering, which had significant negative correlation with yield. Suresh (2004) observed that all characters except days to first flowering are correlated with yield both at phenotype and genotypic levels. Average fruit weight and fruits per plant had positive direct effect on yield.

#### Cucumber

Rastogi and Deep (1990) in a study with 25 genotypes of cucumber revealed positive correlation of yield per plant with fruits per plant, fruit weight and fruit size. Positive correlation of yield with vine length, branches per vine, number of flowers per vine and total fruits per vine and negative correlation with days to flower and percentage of deshaped fruits were recorded by Satyanarayana (1991). Chen et al. (1994) found that there was significant positive genotypic correlation between number of flowers, number of parthenocarpic fruits and yield.

Ma et al. (1995) reported that total yield had a significant positive correlation with total fruit number, fruit size and average fruit weight in cucumber. Damarany et al. (1995) observed a negative relationship between total yield and number of days for first flowering. The total yield had significant positive correlation with total fruit number.

Fruits per plant, average fruit weight, fruit size and fruit diameter were highly correlated with yield in cucumber (Gayathri, 1997). Zhang et al. (1999) reported that the three traits with largest direct positive action on early yield were average fruit weight, harvested fruits per plant and average fruit length in cucumber. Correlations analysis indicated that yield was positively correlated with fruit weight and fruit length both at genotypic and phenotypic levels (Rao et al., 2004).

#### Bitter gourd

In bitter gourd, Lawande and Patil (1989) reported that yield per plant was positively correlated with fruit weight and fruits per plant. In correlation and path analysis of 21 genotypes of bitter gourd they indicated that fruit yield per vine was positively correlated with fruits per vine, average fruit weight and fruit length. Fruit yield per vine was negatively correlated with days to first flower opening both at genotypic and phenotypic levels. Bhave et al. (2003) observed that fruit number per vine was highly correlated with total fruit yield per vine.

#### Snake gourd

Pynadath (1978) reported that yield per plant was highly correlated with days to first female flower, fruit weight and fruit girth. Fruit weight, fruit girth and number of fruits were the characters contributing to yield in snake gourd. Ashok (2000) noted strong correlation between yield and fruit characters and fruits per plant in snake gourd. 14

#### Water melon

In water melon, Shibukumar (1995) observed that yield per plant recorded high positive genotypic and phenotypic correlations with fruits per plant, weight of individual fruit and fruit size. Negative correlation was seen for yield per plant with days to first flower opening. Prasad et al. (2002) revealed that number of fruits per plant had significantly high positive correlation with yield per plot and significant negative association was noticed with days to first male and female flower opening. Saroj Rolania et al. (2003) reported that magnitude of genotypic correlation coefficient was generally higher than that of phenotypic correlation coefficient. Fruit yield was positively correlated with main vine length, number of primary branches per plant, number of nodes per plant, number of female flowers per plant, number of fruit per plant and harvest duration.

#### Pointed gourd

In pointed gourd, Prasad and Singh (1990) reported positive association of yield with fruit number and negative correlation with days to first flowering and picking. Singh et al. (1993) reported that yield was positively correlated with number of fruits per plant in pointed gourd. Fruits per plant, days to first flowering and average fruit weight were responsible for yield increase. Sarkar et al. (1999) conducted correlation studies in pointed gourd and reported that fruit weight and fruit size was positively and significantly correlated with yield per plant at genotypic and phenotypic levels.

#### Ridge gourd

Anitha (1998) reported that days to first female flower, days to first harvest, fruits per plant and average fruit weight had significant positive correlation with yield in ridge gourd. Rao et al. (2000) observed that in ridge gourd yield per vine was significantly and positively associated with fruits girth and fruits per vine. Negative association was observed for days to first female flower. Julie Mole et al. (2002) in their studies using 53 accessions of ridge gourd observed that the number fruits per plant had the highest positive and significant correlation with yield. Karuppaiah and Kavitha (2002) observed significant and positive correlation of yield with single fruit weight, fruit length, fruit girth, number of fruits per plant, fruit size index, flesh thickness, number of male and female flowers per plant, days to first male flowering and days to harvest.

#### 2.3. Path coefficient analysis

Path analysis is an extension of multiple regressions. It goes beyond regression in that it allows for the analysis of more complicated models. In particular, it can examine situations in which there are several final dependent variables and those in which there are "chains" of influence, in that variable A influences variable B, which in turn affects variable C. Despite its previous name of "causal modelling," path analysis cannot be used to establish causality or even to determine whether a specific model is correct; it can only determine whether the data are consistent with the model. However, it is extremely powerful for examining complex models and for comparing different models to determine which one best fits the data.

#### Ivy gourd

In ivy gourd Varghese (2003) reported almost all characters showed positive direct effect on fruit yield per plant except days to first flowering which showed negative direct effect.

#### Cucumber

In cucumber, Prasunna and Rao (1988) reported that fruits per vine and average fruit weight were the important yield contributing characters. Fruits per vine and yield were showed significant positive correlation. Rajput et al. (1991) reported that harvest period also influenced yield but its degree of association was reduced with increasing vine length. Prasad and Singh (1992) in the path analysis of yield in 23 genotypes of cucumber revealed positive direct effect of vine length, days to female flower appearance, fruit weight and fruit length on yield.

Path analysis of eight genotypes of cucumber revealed that fruits per plant had maximum direct positive effect on yield followed by fruit weight (Saika et al. 1995). Gayathri (1997) reported that fruit size exerted maximum direct positive effect on yield followed by average fruit weight and fruits per plant in cucumber. Rao et al. (2004) revealed that characters such as fruit weight and numbers of fruits per plant are highly dependable and reliable for selection to improve yield.

#### Pumpkin

In pumpkin, Kumaran et al. (1998) reported that fruits per plant exhibited the highest direct effect on yield. High positive indirect effect was exerted by fruits per plant and mean fruit weight.

#### Bitter gourd

In bitter gourd, Paranjape and Rajput (1995) found that yield was mainly contributed by fruits per vine and fruit weight. The fruit weight had maximum direct effect on yield and fruits per vine indirectly contributed towards yield. Bhave et al. (2003) reported fruit length and average fruit weight had the highest positive direct effects, followed by harvesting span, fruit length and number of fruits per vine.

#### Ridge gourd

In ridge gourd, Rao et al (2000) reported that fruits per vine and weight of fruit had high direct effect of yield. Karuppaiah and Kavitha (2002) observed that the number of fruits per plant, number of female flowers per plant, flesh thickness, number of male flowers per plant, days to female flower opening exerted maximum positive direct effect on yield in addition to their positive indirect effect through other characters.

#### Ash gourd

Menon (1998) found that in ash gourd average fruit weight exhibited highest positive direct effect on fruit yield followed by number of fruits.

#### Watermelon

. 1

Shibukumar (1995) reported that in watermelon, fruit yield was directly affected by days to flower followed by fruits per vine and fruit weight.

#### Musk melon

Dhaliwal et al. (1996) reported that in musk melon, fruit yield was directly related by days to first flowering followed by fruit per vine and fruit weight. Lal and Singh (1997) observed that fruits per vine and fruit weight had direct effect on yield.

#### 2.4. Standard heterosis

The term heterosis was coined by Shull in 1914. It refers to the superiority of  $F_1$  hybrids over its parents. In other words, heterosis refers to increase of  $F_1$  in fitness and vigour over the parental values. While heterosis refers to the phenomenon (cause), hybrid vigour is the phenotypic expression (effect) of the genetical phenomenon. In current usuage, heterosis and hybrid vigour are used as synonyms and interchangeable.

Existence of significant amount of dominance variance is essential for undertaking heterosis breeding programme. Dominance effects are associated with heterozygosity. Therefore, in plant populations, dominance effects are expected to the maximum in cross pollinated crops (Frey, 1966). For this reason, occurrence of heterosis has been reported for most of the traits. The available literature on heterosis in cucurbits is presented in a crop wise manner.

#### Cucumber

Imam et al. (1977) reported that heterosis ranged from 15.34 per cent for fruit diameter to 59.22 per cent for fruit shape index in cucumber. Pyzhenkov and Kosareva (1979) made hybrids between four male and four female parents in cucumber and reported that heterosis for yield was reflected as increased number of fruits per plant. The mean fruit weight of the hybrids was not more than parents.

Kasem and Somsak (1991) evaluated the hybrid performance of crosses among 21 mini- cucumber lines and reported significant heterosis for characters like flowering habits yield and fruit characters like fruit length, fruit width and average fruit weight.

Fang et al. (1994) developed a hybrid 'Zhongnong 8' from a cross between 90271 and line 90211 which was heterotic over standard variety for total yield per vine and average fruit weight in cucumber. Significant positive heterosis for total yield, fruit number and average weight were reported in cucumber by Li et al. (1995). Gayathri (1997) reported all the three types of heterosis for days to first female flower opening in cucumber. Wilson and Deepthy (2006) studied heterosis for fruit characters in melon and found significant standard heterosis for number of fruits in seven hybrids and six hybrids exhibited significant standard heterosis for yield per plant.

Susheel Sharma et al. (2007) reported maximum heterobeltiosis of 39.57 per cent in the hybrid AAUC  $-1 \times K$ . Paprola and the minimum heterobeltiosis was observed in the hybrid Sel. 75-1-10 x K. Paprola for marketable yield per plant when Pusa Sanyog was used as a standard check.

#### Bitter gourd

Lawande and Patil (1990) reported that heterosis for yield per vine was 86.1 per cent in bitter gourd. Ranpise et al. (1992) derived information on heterosis from diallel analysis of eight lines and 28  $F_1$ hybrids in bitter gourd and reported 64 per cent heterosis for yield in the most promising hybrid.

Mishra et al. (1994) indicated a high level of heterosis for fruits per plant, fruit length, weight and yield in bitter gourd after performing a diallel analysis using nine varieties. Kenndy et al. (1995) reported that a heterosis of 65.7 per cent over standard parent. Ram et al. (1997) observed negative heterosis for days to female flower anthesis. Fruits per plant and yield per plant were the most heterotic characters.

#### Bottle gourd

Janakiram and Sirohi (1992) in their studies on heterosis in round fruited bottle gourd revealed that the best performing hybrid S 46 x S 54 gave 148.97 per cent higher yield over the commercial cultivar Pusa Summer Prolific Round and 84.5 per cent over best parental line. Sharma et al. (1995) gathered information on heterosis in bottle gourd from a line x tester cross and observerd that, the cross Summer Long Green Selection 2 x Faizabadi Long had the largest heterosis over control cultivar Pusa Summer Prolific Long for number of fruits (106.63 per cent) and total yield per plant (110.33 per cent). Another cross showed 22.93 per cent heterosis for fruit length. Dubey and Maurya (2002) reported heterosis over the three best parents ranging from 89.76 to 121.35 per cent over the mid parent, 68.44 to 91.02 percent over the better parent and 80.94 to 89.47 per cent over the check variety for fruit yield per plant. Significant heterobeltiosis and standard heterosis for fruit yield per plant, total soluble sugars, fruits per plant and fruit weight were recorded by Bhalala et al. (2002) evaluated 45 crosses in bottle gourd and found 74.4 per cent heterobeltiosis and 74.5 per cent standard heterosis for fruit yield per plant. High magnitude heterosis for total soluble sugars, fruits per plant and fruit weight also reported in their study.

#### Pumpkin and Squashes

Sirohi (1993) noticed appreciable heterosis for important quantitative characters including yield in pumpkin. The  $F_1$  'Pusa Hybrid 1' showed significant heterosis for yield over the commercial check Pusa Vishwas. According to Ghai et al. (1998)  $F_1$  hybrids in summer squash showed significant heterosis for yield. Pandey et al. (2002) evaluated 12  $F_1$  hybrids of pumpkin and reported 82 per cent heterosis for number of fruits per plant, 44.84 per cent for average fruit weight and 27.33 per cent for length of fruits.

#### Snake gourd

Varghese and Rajan (1993 b) studied heterosis for growth characters in snake gourd and found standard heterosis for days to first fruit picking maturity. Radhika (1999) reported that manifestation of heterosis for all the characters studied. Among the hybrids Thrikkannapuram Local x Kaumudi had maximum standard heterosis (73.28 per cent) for yield and yield related characters.

#### Ash gourd

Mandal and Sirohi (2002) reported heterosis for early maturity, longer vine, more number of fruits per plant, bigger fruit size and higher fruit weight in ash gourd.

#### Water melon

Reddy et al. (1987) evaluated six watermelon cultivars with their 15  $F_1$  hybrids in a diallel analysis and reported that hybrids showed significant heterosis for yield per plant, number of fruits per plant and average fruit weight.

#### Musk melon

Choudhary et al. (2002) reported heterobeltiosis for fruit yield per plant in three crosses viz., MS<sub>1</sub> x Hara Madhu, Jobner Local x Durgapura Madhu and Jobner Local x Hara Madhu.

## **MATERIALS AND METHODS**

#### 3. MATERIALS AND METHODS

The present study was carried out at the Department of Plant Breeding and Genetics, College of Agriculture, Vellayani during 2007-2008 as two experiments with a view to develop and evaluate hybrids in ivy gourd. The details of materials used and methods adopted for the study are presented below.

Ivy gourd is propagated by vegetative method. Clonal variation exists among the clones grown in different regions. Being a dioecious crop, conventional biometric analysis is not possible in this crop. However hybrid seeds can be produced through heterosis breeding. The estimation of variation between and with in the hybrid progenies facilitates selection of superior hybrids.

#### 3.1 Materials

The experimental material comprised of four gynoecious lines of ivy gourd (*Coccinia grandis* L Voigt.) namely Royappanpatty 1 (F.P-1), NBPGR-9 (F.P-2), Kannur-6 (F.P-3), and Karuvalur (F.P-4) and three androecious lines namely Royappanpatty local (M.P-1), Vellayani local (M.P-2) and Karyavattom (M.P-3). The fruit characters of the female parents and the floral and leaf characters are presented in Plate 1. The leaf and flower characters of the male parents are presented in Plate 2. The selected parents were crossed with each other to produce twelve hybrid combinations.

# Morphological Characters of the Female Parents

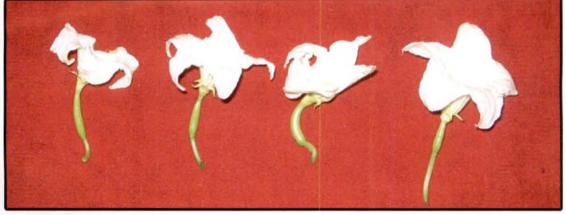


FP 1

FP 2

FP 3

FP 4

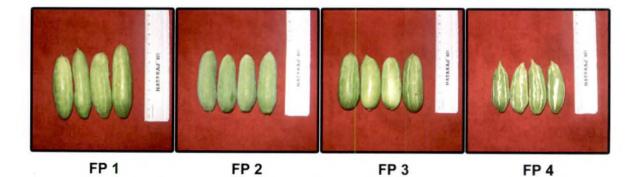


FP 1

FP 2

FP 3

FP 4



# Morphological Characters of the Male Parents



MP 1



MP 2







MP 1



MP 2 Plate 2



MP 3

# Hybridization Technique



Crosses	Combinations	Parentage
F <sub>1</sub>	FP1 x MP1	Royappanpatty 1 x Royappanpatty local
F <sub>2</sub>	FP1 x MP2	Royappanpatty 1 x Vellayani local
F <sub>3</sub>	FP1 x MP3	Royappanpatty 1 x Karyavattam
F <sub>4</sub>	FP2 x MP1	NBPGR-9 x Royappanpatty local
F <sub>5</sub>	FP2 x MP2	NBPGR-9 x Vellayani local
F <sub>6</sub>	FP2 x MP3	NBPGR-9 x x Karyavattam
F <sub>7</sub>	FP3 x MP1	Kannur-6 x Royappanpatty local
F <sub>8</sub>	FP3 x MP2	Kannur-6 x Vellayani local
F9	FP3 x MP3	Kannur-6 x Karyavattam
F <sub>10</sub>	FP4 x MP1	Karuvalur x Royappanpatty local
F <sub>11</sub>	FP4 x MP2	Karuvalur x Vellayani local
F <sub>12</sub>	FP4 x MP3	Karuvalur x x Karyavattam

Table 1 Female and male parents used for the experiments

#### 3.1.1 Experiment I: Production of F1 seeds

The parents for hybridization were raised in the field. Staggered planting was done to facilitate synchronous flowering to ensure successful production of hybrids in all combinations.

Four female parents and three male parents were crossed in all possible combinations to get hybrid seeds. Mature male and female flower buds in the parental lines were kept covered with a cover on the previous day of anthesis. On the following day, hand pollination was done between 8.30 am to 10 am (Plate 3) and pollinated female flowers were kept covered and labeled. The covers kept over female flowers were removed after two or three days. The ripe fruits were harvested for collecting seeds. The seeds were dried in shade and stored for raising  $F_1$  progeny.

**Field View** 



Plate 4

### 3.1.2. Experiment II: Evaluation of F<sub>1</sub> hybrids

The hybrid seeds obtained from 12 cross combinations were raised in the field for evaluation (Plate 4)

#### 3.2 Design and Layout

The experiment was laid out in Compact Family Block Design (CFBD) with three replications at spacing of  $3 \times 3 \text{ m}$ . Each hybrid progeny consists of seven plants. The male progenies were removed from the experimental field.

#### 3.2.1 Cultural practices

The seeds of twelve crosses were sown in pots separately to raise seedlings for planting in the main experimental field.

The experimental field was prepared and pits were taken at a spacing of 3 x 3 m. FYM @ 25 kg/pit was given in two split doses. Twenty five days old hybrid seedlings were used for planting. Three seedlings were planted per each pit but after the establishment of seedlings only two seedlings were retained. Pandals were raised for trailing. Basal application of fertilizers at the rate of NPK 28:10:10 g per pit was given followed by top dressing at three weeks interval. Regular irrigation was given during dry periods. Need based application of insecticides was done to protect the crop from insect pests.

#### 3.2.2. Biometric observations

Biometric observations were taken from seven plants in each treatment adopting standard procedures and average was worked out for each replication.

#### 3.2.2.1 Days to first flowering

Number of days taken from sowing to anthesis of first female flower in each plant was recorded.

#### 3.2.2.2 Days to first harvest

Number of days taken from sowing to first harvest in each plant was recorded.

### 3.2.2.3 Fruit length (cm)

The length of five fruits taken at random from each plant was measured and the average worked out.

### 3.2.2.4 Fruit girth (cm)

The girth of five fruits taken at random from each plant was measured and the average worked out.

# 3.2.2.5 Fruit weight (g)

The weight of five randomly selected fruits from each plant was taken and the average worked out to obtain the mean fruit weight in grams.

### 3.2.2.6 Number of fruits per plant

The number of fruits obtained from each plant in fifteen harvests was recorded separately and the total was worked out to record the number of fruits per plant.

### 3.2.2.7 Vitamin C

Vitamin C content in tender fruits was estimated using volumetric method and expressed as mg/100 g (Sadasivam and Manickam, 1996).

### 3.2.2.8 Fruit yield per plant

The weight of fruits obtained from each plant in fifteen harvests was recorded separately in grams and the total was worked out to record the fruit yield per plant.

### 3.2.2.9 Morphological appearance

Colour and pigmentation of fruits in each progeny were recorded.

### 3.2.2.10 Incidence of pests and diseases

Incidence of pests or diseases were observed and recorded.

# 3.3 Statistical Analysis

The data collected were subjected to statistical analysis.

### 3.3.1 Analysis of variance

Analysis of variance was carried out for all the traits to find out whether there is any difference among the families.

Source	df	SS	MS	F
Replication	r-1	SSR	MSR	
Families	f-1	SSF	MSF	
Error	(r-1) (f-1)	SSE	MSE	

# Analysis of variance for families

# Analysis of variance for progenies within the family

SSR	MSR		
SSP	MSP	i	
1) (p-1) SSE	MSE		
	SSP	SSP MSP	SSP MSP

# Pooled analysis of variance

Source	df	SS	MS	F
Replication	r-1	SSR	MSR	
Families	f-1	SSF	MSF	
Error	(r-1)(f-1)	SSE	MSE	
Progeny in F <sub>1</sub>	(p-1)	SSP	MSP	
F <sub>2</sub>	(p-1)	SSP	MSP	
F <sub>3</sub>	(p-1)	SSP	MSP	
F_4	(p-1)	SSP	MSP	
F5	(p-1)	SSP	MSP	
F_6	(p-1)	SSP	MSP	
F <sub>7</sub>	(p-1)	SSP	MSP	
F8	(p-1)	SSP	MSP	-
F <sub>9</sub>	(p-1)	SSP	MSP	
F <sub>10</sub>	(p-1)	SSP	MSP	
F <sub>11</sub>	(p-1)	SSP	MSP	
F <sub>12</sub>	(p-1)	SSP	MSP	
Pooled error	F(r-1)(p-1)	SSE	MSE	

Where,  $\mathbf{r} =$  Number of replication,

- P = Number of progenies,
- MSR = Replication mean square
  - MSF = Family mean square

f = Number of treatments

SSR = Replication sum of square

SSF = Family sum of square

When the treatments differed significantly by the F test, the pair wise comparison of the treatment means are made by using critical difference as

Critical difference (CD) = 
$$t_{(\infty)} \sqrt{\frac{2MSE}{r}}$$

Where, t  $\infty$  is the students't' table value for  $\infty$  (5 per cent or 1 per cent) level of significance corresponding to the error degree of freedom.

#### 3.3.2 Components of variance

Based on the initial analysis homogeneous group of progenies within each family were identified. These groups of progenies within the family were used for the estimation of genetic components. The mean squares between treatments consisted of variances attributable to genotype, environment and phenotype (Singh and Chaudhary, 1985).

For each character the phenotypic and genotypic components of variance were estimated by equating the expected value of mean squares (MS) to the respective variance components (Jain, 1982). Based on this the following variance components were estimated.

- i) Genotypic variance,  $\sigma^2 g = \frac{MST MSE}{r}$
- ii) Environmental variance,  $\sigma^2 e = MSE$
- iii) Phenotypic variance,  $\sigma^2 p = \sigma^2 g + \sigma^2 e$

#### 3.3.3 Coefficient of variation

It is a unit of measurement used for comparison of variation of different characters measured in different units. Genotypic and phenotypic coefficients of variation were worked out using the estimate of  $\sigma^2 g$  and  $\sigma^2 p$  and expressed in percentage (Burton, 1952) for each trait.

i) Phenotypic coefficient of variation (PCV)

$$=\frac{\sigma p}{Mean}$$
 x 100

ii) Genotypic coefficient of variation (GCV)

$$= \frac{\sigma g}{Mean} \times 100$$

#### 3.3.4 Heritability

For each trait heritability (broad sense) was estimated as the ratio of genotypic variance to phenotypic variance and expressed as percentage (Jain, 1982).

Heritability (H<sup>2</sup>) = 
$$\frac{\sigma^2 g}{\sigma^2 p} \times 100$$

Heritability per cent was categorized as suggested by Johnson et al. (1955) viz., low (0 - 30), moderate (30 - 60) and high (above 60).

#### 3.3.5 Genetic Advance (Johnson et al. 1955 and Allard 1960)

Genetic advance which measures the change in mean genotypic level of the population brought about by selection depends upon standardised selection differential, heritability and phenotypic standard deviation (Allard, 1960).

Genetic advance as percentage of mean was estimated as per the method suggested by Lush (1940) and Johnson et al. (1955) for each trait as

Genetic advance, GA = 
$$\frac{\text{k H}^2 \sigma \text{p}}{\overline{X}}$$
 100

Where, k is the standardised selection differential (k = 2.06) at five per . cent selection intensity and  $\overline{X}$  is the mean of the character over all accessions. Genetic advance was categorized into low (below 10 per cent), moderate (10-20 per cent) and high (above 20 per cent) as suggested by Johnson et al. (1955).

#### 3.3.6 Correlation Analysis

Phenotypic, genotypic and environmental correlation coefficients were worked out for two characters  $X_i$  and  $X_j$  as

Genotypic correlation  $(r_{g_{ij}}) = -\frac{\sigma_{g_{ij}}}{\sigma_{g_i} \times \sigma_{g_i}}$ 

Phenotypic correlation  $(r_{p_{ij}}) = \frac{\sigma_{p_{ij}}}{\sigma_{p_i} \times \sigma_{p_j}}$ 

Environmental correlation  $(r_{e_{ij}}) = \frac{\sigma_{e_{ij}}}{\sigma_{e_i} \times \sigma_{e_i}}$ 

where,  $\sigma_{g_{ij}}$ ,  $\sigma_{P_{ij}}$  and  $\sigma_{e_{ij}}$  denote the genotypic, phenotypic and error covariances between two traits  $X_i$  and  $X_j$  respectively.

#### 3.3.7 Path coefficient analysis

Path analysis is applied to identify relatively important component characters (which are the independent 'v' variables) as a dependent variable on the basis of their direct and indirect effects and it helps the plant breeder to lay emphasis on component characters during selection. The solution of the matrix equation

 $A \underline{B} = \underline{C}$ 

Where A is the genotypic inter-correlation matrix with respect to independent variables, B is the column vector of genotypic correlation co efficient between the dependent and independent variables. Vector <u>B</u> provides estimates of path coefficient which means the direct effect of the independent variable on dependent variable, and also the indirect effect of each independent variable on dependent variable through other variables. Residual variation which could arise from unknown and uncontrollable factor was also estimated using vector <u>B</u> (Dabholkar, 1992).

30

The direct and indirect effects were calculated and classified into very high (> 1), high (0.30 - 0.99), moderate (0.20 - 0.29), low (0.10 - 0.19) and negligible (0.00 - 0.09) (Lenka and Mishra, 1973)

#### 3.3.8 Standard heterosis

Heterosis can be estimated in three different ways.

- As the percentage deviation of the mean performance of F<sub>1</sub>'s from its mid parent which is referred as relative / average heterosis (RH)
- As the percentage deviation of the mean performance of F<sub>1</sub>'s from better parent which is referred as heterobeltiosis (HB)
- iii) As the percentage deviation of mean performance of F<sub>1</sub>'s from a standard parent which is referred as standard heterosis (SH).

Since ivy gourd is a dioecious crop average heterosis and heterobeltiosis cannot be estimated. The standard heterosis was estimated using "Sulabha" as the standard parent.

$$SH = \frac{\overline{F_1} - \overline{SP}}{\overline{SP}} \times 100$$

To test the significance of  $\overline{F_1} - \overline{SP}$  observed in standard heterosis Critical Difference is worked out as

$$CD (0.05) = t_{\alpha} \sqrt{\frac{2MSE}{r}}$$

Where,  $t_{\alpha} = t$  value for error degrees of freedom at 5 per cent level of significance

MSE = Error mean square

r = Number of replications.

# **RESULTS**

.

#### 4. RESULTS

The results of the study entitled "Development and evaluation of hybrids in ivy gourd (*Coccinia grandis* L.Voigt.)" are presented below.

#### 4.1 Production of hybrids in ivy gourd

The hybrid seeds were produced successfully between four gynoecious and three androeicous lines of ivy gourd. The best time of hand pollination for maximum seed set was found to be between 8.30 and 10 am. There was no dormancy and hybrid seeds recorded more than eighty per cent germination in all the crosses.

#### 4.1.1 Evaluation of hybrids

The twelve  $F_1$  families and each family consisting of seven progenies were evaluated in the field for 10 biometric and quality characters namely days to first flowering, days to first harvest, average fruit length (cm), average fruit girth (cm), average fruit weight (g), number of fruits per plant, vitamin C, fruit yield per plant (g), morphological appearance and incidence of pests and diseases.

#### 4.1.1.1 Variability among the families

The analysis of variance conducted for different characters are presented in Table 2. The results showed significant difference for all the characters among the different families.

#### 4.1.1.2 Mean Performance of the families

The mean performances of 12  $F_1$  families for eight biometric characters studied are given in Table 3.

#### 4.1.1.2.1 Days to first flowering

Days to first flowering showed significant differences among the families. The mean values for this character ranged from 49.62 to 87.99 days. The  $F_1$  (49.62 days) took the minimum number of days to first flowering, which was on par with  $F_3$  (50.58 days). Other families were

significantly different from these two families for days to flowering. The  $F_{12}$  (87.99 days) took the maximum days to first flowering.

#### 4.1.1.2.2 Days to first harvest

There was no significant difference among the families for days to first harvest. The mean values for this character ranged from 66.16 to 101.28 days. The  $F_1$  took the minimum days to first harvest (66.16 days) followed by  $F_9$  (74.86 days). The  $F_{12}$  took maximum days (101.28 days) to first harvest.

# 4.1.1.2.3 Average fruit length (cm)

Average fruit length showed significant differences among the families. Fruit length ranged from 4.87 to 8.34 cm. The maximum average fruit length was recorded in the  $F_1$  (8.34 cm) followed by  $F_3$  (8.14 cm) and the minimum fruit length was recorded in the family  $F_4$  (4.87 cm) followed by  $F_8$  (5.58 cm).

#### 4.1.1.2.4 Average fruit girth (cm)

Fruit girth showed significant difference among the families. The mean values for this character ranged from 5.74 to 7.20 cm. The family  $F_1$  showed maximum fruit girth (7.20 cm). The lowest fruit girth was observed in  $F_4$  (5.74 cm).

### 4.1.1.2.5 Average fruit weight (g)

Fruit weight showed significant differences among the families. Fruit weight ranged from 7.44 to 20.93 g. The family  $F_1$  showed maximum average fruit weight (20.93g) whereas the  $F_4$  showed the least average fruit weight (7.44g).

#### 4.1.1.2.6 Number of fruits per plant

Number of fruits per plant showed significant differences among the families and ranged from 219.95 to 501.52. The maximum number of fruits per plant was observed in the family  $F_1$  (501.52) followed by  $F_2$ (467.27). The minimum number of fruits per plant was observed in the family  $F_{12}$  (219.95) which was on par with the families  $F_6$  (240.27),  $F_5$ (258.09) and  $F_7$  (263.79).

Sources	df	Days to	Days to	Average	Average	Average	Number of	Vitamin C	Fruit yield per
			fruit length	fruit girth	fruit weight	fruits per	mg/100g	plant (g)	
	L	flowering		(cm)	(cm)	(g)	plant	88	P (8)
Replication	2	994.938	736.545	1.776	0.219	6.957	5470.506	5.823	1467737,000
Families	11	2777.048	1951.730	17.908	4.644	271.489	149892.200	57.856	72131390.000
Error	_ 22_	192.549	50.195	0.891	0.302	8.990	7587.567	9.009	1605516.000
Progeny in F <sub>1</sub>	6	7.603	1.714	0.084	0.043	3.710	6216.000	2.039	871537.800
F_2	6	6.635	9.857	0.153	0.149	3.124	2844.389	1.749	585048.900
F <sub>3</sub>	6	10.968	9.539	4.250	0.109	21.102	7358.180	9.177	4012089.000
<u>F4</u>	6	4.873	16.158	0.744	0.418	3.660	2590.222	15.577	398612.000
<u> </u>	6	29.539	57.650	0.054	0.291	6.850	675.181	15.704	1054546.000
F <sub>6</sub>	6	18.762	4.190	0.391	0.506	12.788	1498.215	7.203	443637.300
F7_	6	19.762	19.858	0.297	0.167	4.595	622.708	12.705	419166.200
<u>F</u> 8	6	3.159	9.190	0.099	0.462	1.893	800.660	6.341	447713.800
<u>F9</u>	6	1.080	7.270	0.597	0.368	5.888	1106.181	4.408	361464.900
F <sub>10</sub>	6	2.873	2.096	0.153	0.153	4.734	808.319	0.772	601484.400
<u>F11</u>	6	18.158	7.207	0.586	0.497	16.827	8247.111	3.297	214000.000
F <sub>12</sub>	6	13.763	1.430	0.355	0.265	1.123	221.934	1.779	48050.670
Pooled Error	144	9.320	9.741	0.581	0.268	7.730	3166.442	5.663	1215664.000

-

# Table 2 Mean square values of eight characters in 12 families of ivy gourd

.

Families	Days to	Days to	Average	Average	Average	Number	Vitamin C	Fruit
	first	first	fruit	fruit girth	fruit	of fruits	Content	yield per
	flowering	harvest	length	(cm)	weight	per plant	(mg/100g)_	plant
			(cm)		(g)			(g)
<u> </u>	49.62	66.16	8.34	7.20	20.93	501.52	20.95	8309.33
2	<u> </u>	76.69	7.08	6.96	16.74	467.27	16.32	7137.96
3	50.58	77.40	8.14	6.86	20.07	422.54	19.43	8058.61
4	<u> </u>	80.70	4.87	5.74	7.44	293.06	20.66	2175.44
5	68.85	<u> </u>	6.70	6.99	14.75	258.09	18.45	4253.63
6	64.87	78.04	6.81	6.76	14.94	240.27	21.12	3476.52
7	69.82	84.43	6.15	6.46	13.02	263.79	20.88	3623.49
8	73.17	83.56	5.58	6.61	12.29	301.51	20.84	3920.12
9	67.76	74.86	6.13	6.10	13.10	298.98	17.73	3764.50
10	<u> </u>	90.42	6.90	5.85	12.65	382.83	18.02	4714.59
11	84.00	96.64	6.78	6.59	16.61	316.93	18.93	3705.52
12	87.99	101.28	6.30	6.03	10.55	219.95	16.82	3148.57
Sulabha	48.40	64.80	7.86	6.43	17.82	356.75	17.93	6031.90
CD @ 5 %	8.80	NS	0.68	0.33	2.44	65.34	2.01	1015.22

.

-

# Table 3 Mean values of eight characters in twelve $F_1$ families of ivy gourd.

# 4.1.1.2.7 Vitamin C (mg/100g)

The vitamin C content of fruits varied significantly among the families. The mean values for this character ranged from 16.32 to 21.12 mg/100g of sample. The maximum vitamin C content was observed in the family  $F_6$  (21.12 mg/100g). The minimum vitamin C content was recorded in the family  $F_2$  (16.32 mg/100g) which was on par with the families  $F_{12}$  (16.82 mg/100g),  $F_9$  (17.73 mg/100g) and  $F_{10}$  (18.02 mg/100g).

#### 4.1.1.2.8 Fruit yield per plant (g)

Fruit yield per plant showed significant differences among the families. The mean values for this character ranged from 2175.44 to 8309.33 g. The family  $F_1$  showed maximum fruit yield per plant (8309.33 g) followed by  $F_3$  (8058.61 g). The family  $F_4$  showed the least fruit yield per plant (2175.44 g) which was on par with the family  $F_{12}$  (3148.57 g).

#### 4.1.1.2.9 Morphological appearance of fruits

There was a lot of variation for the morphological appearance of fruits. The variations were evident in the colour and shape of fruits and presence or absence of prominent streaks and patches. (Plate 5-10)

#### 4.1.1.2.10 Incidence of pests and diseases.

There was no incidence of major pests and diseases in the experimental plots during the course of study. However irrespective of genotypes minor incidence of fruit fly attack was noticed.

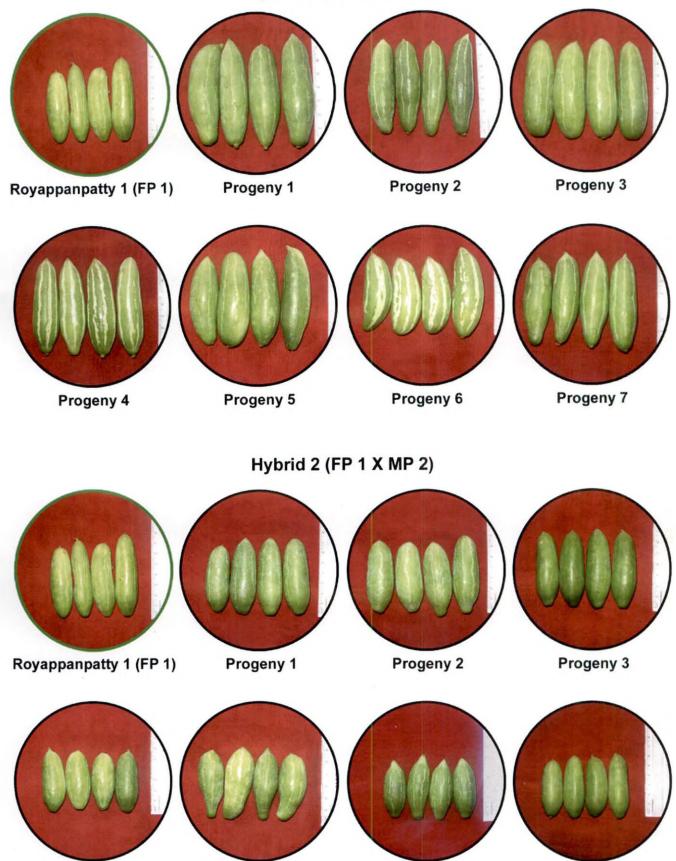
#### 4.1.2 Mean performance of the progenies within families

The mean performance of the progenies in 12 families for different characters is presented in Table 4 and Plate 5 - 10.

#### Family 1

There were no significant differences among the progenies for different characters studied. The mean values for days to first flowering ranged from 48.00 to 52.33 days, days to first harvest from 65.00 to 67.33 days, average fruit length from 7.92 to 8.42 cm, average fruit girth from 7.00 to 7.30 cm, average fruit weight from 18.83 to 21.83 g, number of

Hybrid 1 (FP 1 X MP 1)



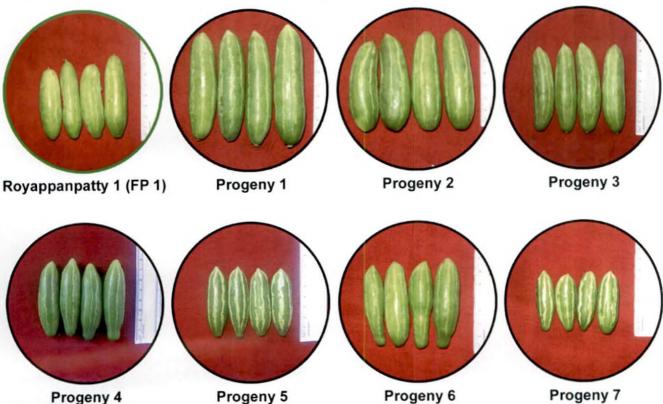
Progeny 4

Progeny 5

Progeny 6

Progeny 7

Hybrid 3 (FP 1 X MP 3)



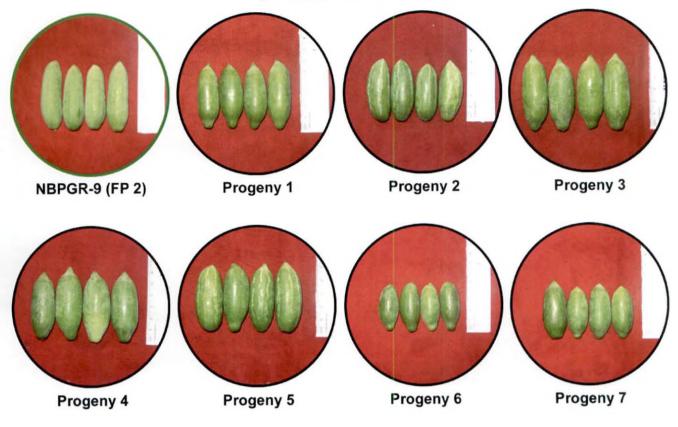
Progeny 4

Progeny 5

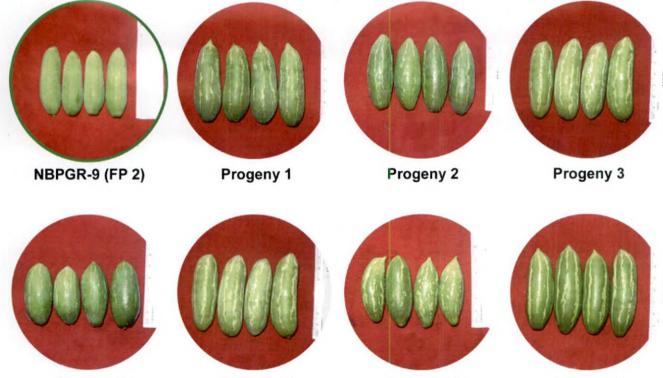
Hybrid 4 (FP 2 X MP 1)



Hybrid 5 (FP 2 X MP 2)



Hybrid 6 (FP 2 X MP 3)



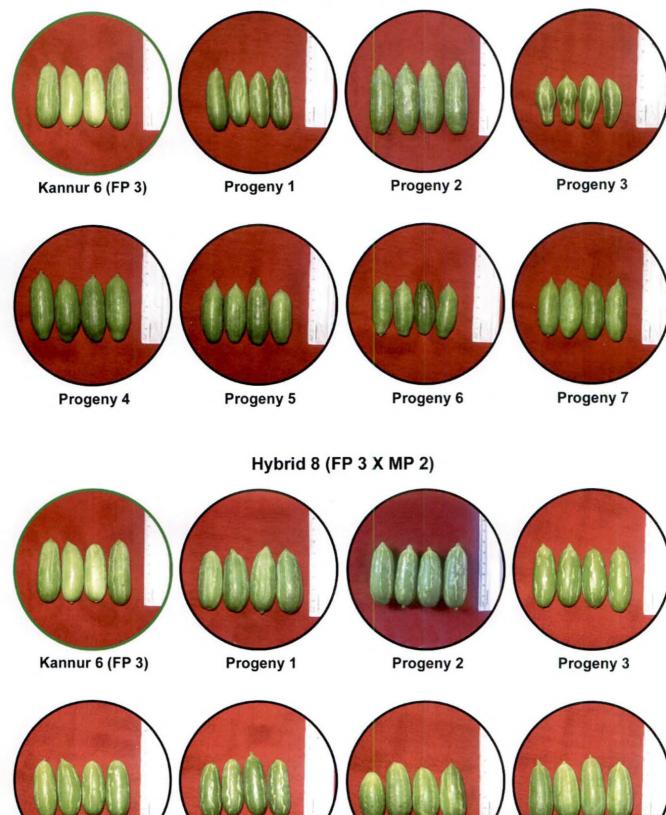
Progeny 4

Progeny 5

Progeny 6

Progeny 7

Hybrid 7 (FP 3 X MP 1)



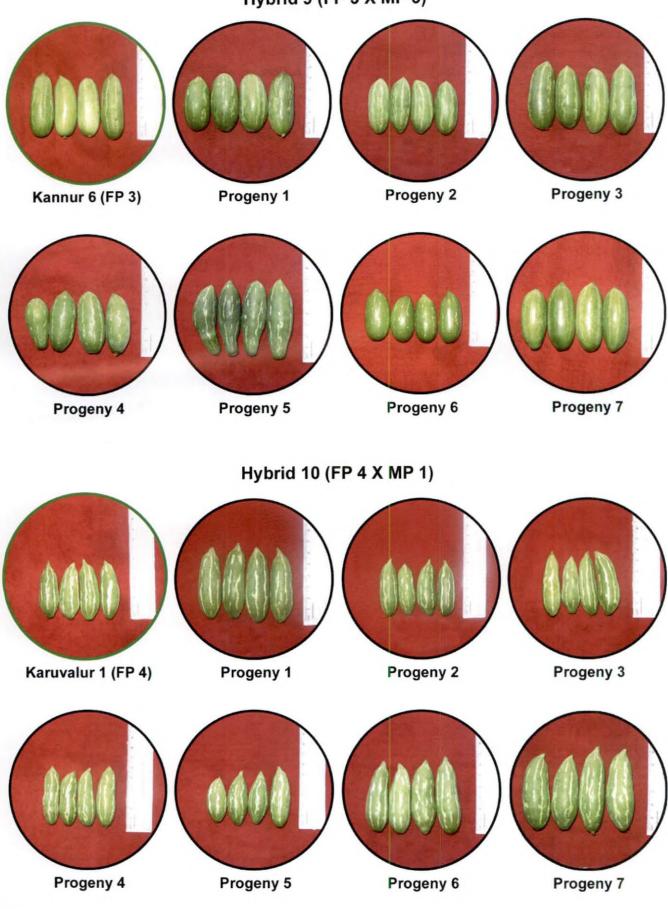
Progeny 4

Progeny 5

Progeny 6

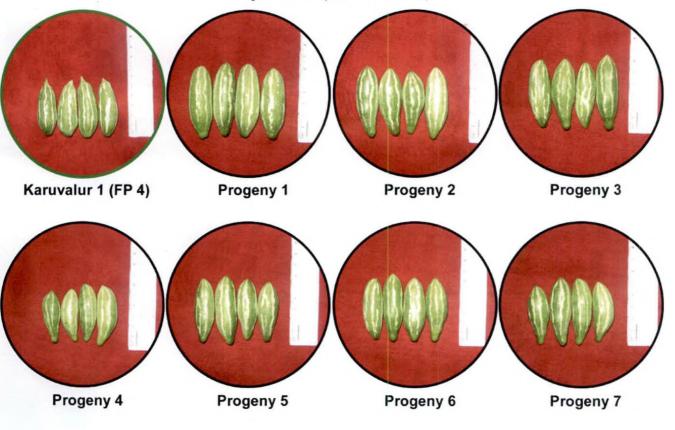
Progeny 7

Hybrid 9 (FP 3 X MP 3)



Hybrid 11 (FP 4 X MP 2) Weight of the formula of t

Hybrid 12 (FP 4 X MP 3)



fruits per plant from 447.66 to 577.00, vitamin C content from 19.04 to 21.42 mg/100 g and fruit yield per plant from 7222.33 to 8686.33 g. Family 2

Days to first flowering significant differences among the progenies of family 2 and ranged from 57.33 to 61.66 days. The progeny 5 (57.33 days) took minimum days to flowering which was on par with the progenies 7 (58.66 days), 6 (59.66 days), 4 (60.33 days) and 1 (60.66 days). The progeny 2 (61.66 days) took the maximum days to first flowering. There was no statistically significant difference in days to first harvest among the progeny of family 2 but exhibited a narrow range of 73.00 to 78.33 days.

Fruit length showed significant difference among the progenies. The mean values ranged from 6.69 to 7.27 cm. The maximum average fruit length was observed in the progeny 1 (7.27 cm) followed by 2 (7.19 cm). The minimum average fruit length was observed in 5 (6.69 cm) which was on par with the progenies 7 (6.72 cm), 4 (6.93 cm), 3 (7.07 cm), 6 (7.11 cm).

Average fruit girth showed significant difference among the progenies. The mean values ranged from 6.64 to 7.20 cm. The maximum average fruit girth was observed in the progeny 1 (7.20 cm). The minimum average fruit girth was observed in 3 (6.64 cm) which was on par with the progenies 7 (6.68 cm), 4 (6.87 cm), 6 (6.86 cm), and 2 (6.83 cm). The average fruit weight, number of fruits per plant, vitamin C and fruit yield per plant content showed no significant difference among the progenies. The mean values for average fruit weight ranged from 14.83 to 17.66g. The number of fruits per plant among the progenies of family 2 ranged from 415.00 to 501.00. The vitamin C mean values ranged from 15.26 to 17.19 mg/100g. The range of mean values for fruit yield per plant was 6399.66 g to 7662.00g.

#### Family 3

Significant differences were observed among the progenies of family 3 for days to first flowering. The mean values ranged from 48.66 to 54.33 days. The minimum number of days to first flowering was observed in the progenyl (48.66 days) which was on par with the progenies 5 (49.33 days), 7 (49.66 days), 3 (50.66 days), 6 (50.66 days) and 2 (52.00 days). The progeny 4 (54.33 days) took the maximum number of days to first flowering. The number of days to first harvest showed no significant difference among the progenies. The mean values ranged from 75.00 to 79.66 days.

The average fruit length showed significant differences among the progenies. The mean values for this character ranged from 5.30 to 8.92 cm. The progeny 3 (8.92 cm) showed maximum fruit length. The minimum average fruit length was observed in the progeny 6 (5.30 cm) which was on par with the progenies 7 (7.51 cm), 4 (7.73 cm), 5 (7.82 cm), 2 (8.23 cm) and 1 (8.66 cm). The characters average fruit girth, average fruit weight, number of fruit per plant, and fruit yield per plant did not show any significant difference among the progenies. The mean values for these characters varied from 6.46 to 7.04 cm, 14.65 to 21.91 g, 333.66 to 458.00 and 5585.66 to 8487.00 g respectively. Vitamin C content showed significance differences in which the maximum vitamin C content was recorded in the progeny 3 (22.07 mg/100g) followed by 5 (20.88 mg/100g). The minimum vitamin C content was recorded in the progeny 4 (17.45 mg/100g) which was on par with he progenies 6 (17.98 mg/100g), 7 (17.98 mg/100g), 1 (18.25 mg/100g) and 2 (19.84 mg/100g). The mean values for this character ranged from 17.45 to 22.07 mg/100g.

#### **Family 4**

The mean values of days to first flowering ranged from 64.33 to 67.66 days. The minimum number of days to first flowering was observed in the progeny 2 (64.33 days) which was on par with 3 (65.00 days), 5 (65.66 days), 1 (66.00 days) and 7 (66.66 days) progenies. The progeny 4

(67.66 days) took maximum number of days to first flowering. Days to first harvest ranged from 78.66 to 84.66 days. The minimum number of days to first harvest was recorded in the progeny 7 (78.66 days) which was on par with the progenies 5 (79.00 days), 6 (79.66 days), 2 (79.66 days), and 3 (80.00 days). The maximum number of days to first harvest was observed in the progeny 4 (84.66 days) followed by the progeny 1 (83.33 days).

The average fruit length ranged from 4.08 to 5.48 cm. The mean values for average fruit girth ranged from 5.17 to 6.30 cm. The maximum average fruit girth was observed in the progeny 1 (6.30 cm) followed by 6 (5.85 cm). The minimum average fruit girth was observed in the progeny 4 (5.17 cm) which was on par with the progenies 5 (5.28 cm), 3 (5.66 cm), 7 (5.69 cm) and 2 (5.72 cm). Fruit weight ranged from 5.59 to 9.00 g. The maximum average fruit weight was observed in the progeny 1 (9.00g). The minimum average fruit weight was observed in the progeny 5 (5.59g) which was on par with all other the progenies 4 (6.25g), 2 (6.70g), 3 (6.75g), 7 (7.25g) and 6 (7.75g).

The maximum number fruits per plant were recorded in the progeny 1 (333.33). The mean values for number of fruits per plant ranged from 246.33 to 333.33. The progeny 5 (246.33) showed minimum number of fruits per plant which was on par with all other progenies, 4 (263.33), 2 (272.66), 7 (284.66), 6 (302.33), and 3 (306.33). The mean values for vitamin C ranged from 17.98 to 24.45 mg/100g. The maximum vitamin C content was observed in the progeny 1 (24.45 mg/100g). The progenies 3 (17.98 mg/100g) and 7 (17.98 mg/100g) recorded the minimum vitamin C content which was on par with 5 (19.44 mg/100g), 2 (19.57 mg/100g), and 6 (19.84 mg/100g). Fruit yield per plant ranged from 1489.00 to 2397.00g. The maximum fruit yield per plant was recorded in the progeny 1 (2397.00g) and the lowest fruit yield was observed in the progeny 5 (1459.00g) which was on par with 4 (1559.66g), 2 (1700.00g), 3 (2123.00g) and 6 (2263.33g) progenies.

#### Family 5

The mean values for days to first flowering ranged from 65.33 to 73.33 days. The progeny 1 took the minimum number of days to first flowering (65.33 days) which was on par with the progenies 2 (66.00 days), 3 (68.33 days) and 5 (69.33 days). The maximum number of days was taken by the progeny 7 (73.33 days) followed by 6 (72.66 days) and 4 (71.33 days). Days to first harvest also showed significant differences among the progenies and ranged from 71.66 to 84.33 days. The progeny 5 (71.66 days) took the minimum number of days to the first harvest which was on par with all other progenies 7 (74.33 days), 6 (76.66 days), 1 (78.33 days), 4 (80.00 days), 2 (84.33 days) except the progeny 2 (84.33 days). Average fruit length and average fruit girth did not show any significant difference among the progenies. The mean values for average fruit length ranged from 6.42 to 5.48 cm and for average fruit girth from 6.57 to 7.35 cm.

The average fruit weight showed significance difference among the progenies. The mean values for this character ranged from 12.50 to 16.25 g. The maximum fruit weight was recorded by the progeny 1 (16.25 g) followed by 6 (16.16 g). The minimum fruit weight was observed in the progeny 2 (12.50 g) which was on par with the progenies 3 (12.70 g), 4 (13.67 g), 7 (13.75 g) and 5 (13.96 g).

The number of fruits per plant showed no significant difference among the progenies. The mean values for this character ranged from 236.66 to 279.66. Vitamin C content varied significantly among the progenies. The mean values ranged from 13.75 to 21.16 mg/100g. The maximum vitamin C content was observed in the progeny 7 (21.16 mg/100g) followed by 3 (19.16 mg/100g) and the minimum vitamin C content was observed in the progeny 1 (13.75 mg/100g) which was on par with the progenies 5 (16.66 mg/100g), 2 (16.93 mg/100g), 6 (17.64 mg/100g), and 4 (18.02 mg/100g). The fruit yield per plant showed significant difference among the progenies and ranged from 3445.66 to 4953.66 g. The maximum fruit yield per plant was observed in the progeny 6 (4953.66 g). The minimum fruit yield per plant was observed in the progeny 3 (3445.66 g) which was on par with the progenies 2 (3471.00 g), 5 (3981.00 g), 7 (4083.00 g), 4 (4123.00 g) and1 (4839.66 g).

#### Family 6

Days to first flowering showed significant differences among the progenies. The mean values ranged from 62.33 to 69.00 days. The progeny 4 (62.33 days) took the minimum number of days to first flowering which was on par with the progenies 3 (62.66 days), 5 (63.66 days), 6 (64.66 days), 7 (66.33 days) and 1 (67.33 days). The maximum number of days was taken by the progeny 2 (69.00 days). Days to first harvest showed no significant differences among the progenies. The mean values ranged from 77.33 to 80.66 days. Average fruit length showed significant differences among the progenies. The mean values for this character ranged from 6.20 to 7.38 cm. The maximum average fruit length was observed in the progeny 4 (7.38 cm). The minimum average fruit length was observed in the progeny 6 (6.20 cm) which was on par with other progenies 2 (6.52 cm), 5 (6.64 cm), 3 (6.69 cm), 1 (6.82 cm), and 7 (6.84 cm). Average fruit girth showed significant differences among the progenies. The mean values for this character ranged from 5.87 to 7.04 cm. The maximum average fruit girth was observed in the progeny 1 (7.04 cm) followed by 4 (6.97 cm). The minimum average fruit girth was observed in the progeny 6 (5.87 cm) which was on par with other progenies 5 (6.30 cm), 2 (6.55 cm), 7 (6.67 cm) and 3 (6.81 cm).

The average fruit weight showed no significance difference among the progenies. The mean values for this character ranged from 11.16 to 17.89 g. The number of fruits per plant showed significant difference among the progenies. The mean values for this character ranged from 208.66 to 269.66. The maximum number of fruits per pant was observed in progeny 1 (269.66) followed by progeny 4 (259.00). The minimum number of fruits per pant was observed in progeny 2 (208.66) which was on par with 6 (215.00), 3 (225.66), 5 (229.33), and 7 (231.66). Vitamin C content showed no significant differences among the progenies. The mean values ranged from 19.16 to 23.80 mg/100g. The fruit yield per plant showed significant difference among the progenies. The mean values ranged from 2923.33 g to 4113.00 g. The maximum fruit yield per plant was observed in the progeny 4 (4113.00 g). The minimum Fruit yield per plant was observed in the progeny 6 (2923.33 g) which was on par with the progenies 1 (3050.66 g), 2 (3203.33 g), 7 (3213.00 g), 3 (3230.66 g) and 5 (3250.66 g).

#### Family 7

In this family the characters showed non significant differences among the progenies except days to first harvest and vitamin C. The mean values for days to first flowering ranged from 67.00 to 73.66 days. Days to first harvest showed significant difference among the progenies and the mean values ranged from 80.66 to 87.33 days. The progeny 6 (80.66 days) took minimum number of days to first harvest which was on par with the progenies 3 (82.00 days), 5 (82.00 days), 7 (83.66 days), and 4 (84.66 days). The progeny 1 (87.33 days) took maximum number of days to first harvest followed by the progeny 2 (87.00 days).

Average fruit length showed non significant differences among the progenies. The mean values for this character ranged from 5.60 to 6.44 cm. Average fruit girth showed non significant differences among the progenies. The mean values for this character ranged from 5.90 to 6.59 cm. Average fruit weight showed non significant differences among the progenies. The mean values for this character ranged from 9.75 to 13.58 g. Number of fruits per plant showed non significant differences among the progenies. The mean values for this character ranged from 233.00 to 279.33. Vitamin C content showed significant differences among the progenies. The mean values for this character ranged from 233.00 to 279.33. The mean values for this character ranged from 17.72 to 23.80

mg/100g. The maximum vitamin C content was observed in the progeny 2 (23.80 mg/100g). The minimum vitamin C content was observed in the progeny 4 (17.72 mg/100g) which was on par with the progenies 1 (18.25 mg/100g), 7 (18.78 mg/100g), 3 (19.57 mg/100g), 6 (20.48 mg/100g), and 5 (20.89 mg/100g). Fruit yield per plant showed no significant difference among the progenies. The mean values ranged from 2526.66 to 3654.33 g. Family 8

Days to first flowering showed no significant differences among the progenies. The mean values ranged from 72.66 to 75.66 days. Days to first harvest showed no significant differences among the progenies. The mean values varied from 81.33 to 85.66 days. Average fruit length showed no significant differences among the progenies. The mean values for this character ranged from 5.20 to 5.74 cm. Average fruit girth showed significant differences among the progenies. The mean values for this character ranged from 5.20 to 5.74 cm. Average fruit girth showed significant differences among the progenies. The mean values for this character ranged from 5.81 to 6.98 cm. The maximum average fruit girth was observed in the progeny 7 (6.98 cm) which was on par with the progenies 6 (6.91 cm), 2 (6.80 cm), and 3 (6.60 cm). The minimum average fruit girth was observed in the progeny 5 (5.81 cm) which was on par with other progenies 4 (6.47 cm) and 1 (6.54 cm).

The average fruit weight showed significance difference among the progenies. The mean values for this character ranged from 10.50 to 12.74 g. The maximum average fruit weight was observed in the progeny 7 (12.74 g). The minimum average fruit weight was observed in the progeny 5 (10.50 g) which was on par with the progenies 1 (11.00 g), 4 (11.41 g), 2 (12.08 g), 3 (12.09 g) and 6 (12.33 g). The number of fruits per plant showed non significant difference among the progenies. The mean values for this character ranged from 274.33 to 321.66. Vitamin C content showed significant differences among the progenies. The mean values ranged from 19.31 to 23.54 mg/100g. The maximum vitamin C content was observed in the progeny 1 (23.54 mg/100g). The minimum vitamin C content was observed in the progeny 4 (19.31 mg/100g) which was on par

with the progenies 7 (19.42 mg/100g), 5 (1984 mg/100g), 3 (20.46 mg/100g), 2 (20.89 mg/100g) and 6 (21.16 mg/100g). Significant differences in fruit yield per plant among the progenies were not noticed. The mean values ranged from 3153.66 to 4124.33 g.

#### Family 9

Days to first flowering did not show any significant differences among the progenies. The mean values ranged from 67.66 to 69.33 days. Days to first harvest also showed non significant differences among the progenies. The mean values ranged from 73.33 to 77.66 days. Average fruit length showed significant differences among the progenies. The mean values for this character ranged from 5.41 to 6.79 cm. The maximum average fruit length was observed in the progeny 3 (6.79 cm) which was on par with progenies 5 (6.51 cm) and 1 (6.30 cm). The minimum average fruit length was observed in the progeny 4 (5.41 cm) which was on par with the progenies 6 (5.90 cm), 7 (5.93 cm) and 2 (5.97 cm).

Fruit girth showed no significant difference among the progenies. The values for this character ranged from 5.53 to 6.51 cm. The average fruit weight also showed non significance difference among the progenies. The mean values for this character ranged from 10.91 to 14.58 g.

The number of fruits per plant showed no significant difference among the progenies. The mean values for this character ranged from 263.00 to 322.66. Vitamin C content showed significant difference among the progenies. The mean values ranged from 15.87 to 19.31 mg/100g. The maximum vitamin C content was observed in the progeny 3 (19.31 mg/100g) which was on par with the progenies 6 (18.51 mg/100g) and 1 (17.98 mg/100g). The minimum vitamin C content was observed in the progeny 2 (15.87 mg/100g) which was on par with the progenies 7 (16.40 mg/100g), 5 (16.93 mg/100g) and 4 (17.19 mg/100g). The fruit yield per plant showed no significant difference among the progenies. The mean values ranged from 3159.00 to 3998.00 g. Days to first flowering showed no significant differences among the progenies. The mean values ranged from 73.00 to 75.66 days. Days to first harvest also showed no significant differences among the progenies. The mean values ranged from 89.33 to 91.66 days. Average fruit length showed significant differences among the progenies. The mean values for this character ranged from 6.35 to 6.99 cm. The maximum average fruit length was observed in the progeny 6 (6.99 cm) followed by 2 (6.98 cm). The minimum average fruit girth was observed in the progeny 5 (6.35 cm) which was on par with the progenies 3 (6.66 cm), 4 (6.85 cm) and 7 (6.91 cm). Average fruit girth showed no significant differences among the progenies. The mean values for this character ranged from 5.40 to 6.04 cm.

The average fruit weight showed significant difference among the progenies. The mean values for this character ranged from 10.16 to 13.66 g. The maximum average fruit weight was observed in the progeny 4 (13.66 g) followed by the progeny 2 (13.50 g). The minimum average fruit weight was observed in the progeny 5 (10.16 g) which was on par with the progenies 3 (11.20 g), 1 (11.62 g), 7 (12.41 g) and 6 (12.57 g). The number of fruits per plant showed non significant difference among the progenies. The mean values for this character ranged from 352.00 to 400.66. Vitamin C content showed non significant differences among the progenies. The mean values ranged from 16.66 to 18.25 mg/100g. The fruit yield per plant also showed non significant difference among the progenies. The mean values ranged from 3665.33 to 5023.33 g.

#### Family 11

Days to first flowering showed significant difference among the progenies. The mean values ranged from 80.00 to 87.00 days. The minimum number of days to first flowering was observed in the progeny 1 (80.00 days) which was on par with the progenies 4 (83.00 days), 5 (85.33 days), 6 (84.00 days) and 7 (85.33 days). The maximum number of days to first flowering

was taken by the progenies 2 (87.00 days) and 3 (87.00 days). Days to first harvest showed no significant difference among the progenies. The mean values ranged from 95.00 to 99.00 days. Average fruit length showed significant differences among the progenies. The mean values for this character ranged from 5.75 to 7.20 cm. The maximum average fruit length was observed in the progeny 2 (7.20 cm). The minimum average fruit length was observed in the progeny 7 (5.75 cm) which was on par with the progenies 4 (6.39 cm), 1 (6.48 cm), 5 (68.53 cm), 3 (6.74 cm) and 6 (6.75 cm). Average fruit girth showed no significant differences among the progenies. The mean values for this character ranged from 5.75 to 7.20 cm.

The average fruit weight showed non significant difference among the progenies. The mean values for this character ranged from 11.09 to 18.25 g. The number of fruits per plant showed significant difference among the progenies. The mean values for this character ranged from 232.00 to 383.66. The maximum number of fruits per plant was observed in the progeny 3 (383.66). The minimum number of fruits per plant was observed in the progeny 7 (232.00) which was on par with 4 (238.66), 6 (273.00), 5 (307.33), 1 (310.66) and 2 (319.66). Vitamin C content showed significant differences among the progenies. The mean values ranged from 17.19 to 20.21 mg/100g. The maximum vitamin C content was observed in the progeny 2 (20.21 mg/100g) followed by 3 (19.42 mg/100g). The minimum vitamin C was observed in the progeny 1 (17.19 mg/100g) which was on par with 5(17.98 mg/100g), 7 (17.98 mg/100g), 6 (19.04 mg/100g) and 4 (19.31 mg/100g). The fruit yield per plant showed non significant difference among the progenies. The mean values ranged from 3323.33 to 4035.66 g. Family 12

Days to first flowering showed significant difference among the progenies. The mean values ranged from 83.66 to 90.33 days. The minimum number of days to first flowering was observed in the progeny 1 (83.66 days) which was on par with the progenies 4 (87.66 days), 7 (88.33 days), 6 (88.66 days) and 3 (89.00 days). The maximum number of days to first flowering

٠

							Cha	racters		
	Progenies	Days to first flowerin	Days to first harvest	Average fruit length	Average fruit girth	Average fruit weight	Number of fruits per plant	Vitamin C	Fruit yield per plant	Morphological characters
		g		<u>(g)</u>	_(g)	(g)		g)		
	1	48.00	65.00	8.42	7.30				7837.66	Light green with irregular white patches
	2	52.33	66.66	8.36	7.25	21.01	447.66	20.63	7222.33	Green with narrow white patches
1	3	50.66	67.00	- 8.17	7.29	21.83	468.66	20.21	8433.00	Light green with narrow white streaks
Family	4	50.00	66.66	8.10	7.10	20.35	577.00	21.16	8686.33	Light green with broad prominent white streaks
Far	5	48.00	66.66	8.11	. 7.07	20.16	455.33	19.67	7311.33	Light green with irregular white patches
	6	50.33	67.00	7.92	7.09	19.41	464.66	21.42	7882.66	Light green with broad prominent white streaks
	7	48.66	67.33	8.22	7.00	18.83	448.33	19.04	7686.66	Light green with narrow white patches
	CD @ 5 %	NS	NŠ	NS	NS	NS	NS	NS	NS	· · · · · · · · · · · · · · · · · · ·
	1	60.66	77.66	7.27	7.20	17.25	501.00	17.19	7662.00	Green with irregular white patches
2	2	61.66	77.33	7.19	- 6.83	16.08	450.33	15.26	6702.00	Light green with irregular white patches
	3	61.00	77.33	7.07	6.64	14.83	415.00	16.15	6399.66	Light green with irregular white patches
Family	4	60.33	75.66	6.93	6.87	16.51	439.00	15.32	6694.00	Green with irregular white patches
	5	57.33	75.66	6.69	7.19	16.66	468.33	15.26	7131.66	Light green with irregular white patches
	6	59.66	73.00	7.11	6.86	17.66	464.33	15.71	7261.66	Light green with irregular white patches
	7	58.66	78.33	6.72	6.68	15.25	415.33	16.66	6622.00	Green with irregular white patches
	CD @ 5 %	3.59	NS	0.55	0.40	NS	NS	NS	NS	

•

.

# Table 4 continued...

	1	48.66	75.00	8.66	6.87	20.66	457.00	18.25	8487.00	Light green with narrow white patches
	2	52.00	76.33	8.23	6.74	20.75	431.00	19.84	7834.00	Light green with narrow white streaks
13	3	50.66	79.33	8.92	7.04	21.91	458.00	22.07	8465.00	Light green with narrow white patches
Family	4	54.33	75.66	7.73	6.46	14.65	364.66	17.45	6078.00	Green with narrow white patches
Fai	5	49.33	76.66	7.82	6.73	18.32	333.66	20.88	7106.66	Green with broad prominent white patches
	6	50.66	79.66	5.30	6.57	17.08	382.66	17.98	6485.66	Light green with irregular white patches
	7	49.66	77.66	7.51	6.66	16.50	365.33	17.98	5585.66	Green with broad regular white patches
	CD @ 5 %	5.52	NS	3.46	NS	NS	NS	3.03	NS	
	1	66.00	83.33	5.48	6.30	9.00	333.33	24.45	2397.00	Dark green with no white patches
4	2	64.33	79.66	4.92	5.72	6.70	272.66	19.57	1700.33	Dark green with scattered white patches
ily	3	65.00	80.00	4.96	5.66	6.75	306.33	17.98	2123.00	Dark green with irregular white patches
Family	4	67.66	84.66	4.08	5.17	6.25	263.33	21.69	1559.66	
	5	65.66	79.00	4.12	5.28	5.59	246.33	19.44	1489.00	
	6	67.66	79.66	4.91	5.85	7.75	302.33	19.84	2263.33	Dark green with irregular white patches
	7	66.66	78.66	4.86	5.69	7.25	284.66	17.98		Dark green with irregular white patches
	CD @ 5 %	3.08	3.47	NS	0.65	2.95	64.26	5.16	773.98	
	1	65.33	78.33	6.83	7.35	16.25	279.66	13.75	4839.66	patches
<u>ہ</u>	2	66.00	84.33	6.52	6.65	12.50		16.93	3471.00	Green, ashy coat with broad white patches
	3	68.33	82.00		6.68	12.70		19.16		
Family	4	71.33	80.00	6.55	6.80	13.67	254.00	18.02	4123.00	Green with irregular white patches
L L	5	69.33	71.66	6.63	6.68	13.96	248.00	16.66	3981.00	Green, ashy coat with irregular white patches
	6	72.66	76.66	6.68	7.25	16.16	274.66	17.64	4953.66	
	7	73.33	74.33	6.42	6.57	13.75	260.66	21.16	_	
	CD @ 5 %	4.22	10.66	NS	NS	3.36	NS	4.97	1478.18	

.

.

.

.

.

.

•

# Table 4 continued...

(

	·									
	<u> </u>	67.33	77.66	6.82	7.04	13.66	269.66	19.84	3050.66	Green with irregular white patches
	2	69.00	78.00	6.52	6.55	13.49	208.66	21.16	3203.33	Dark green with irregular white patches
y 6	3	62.66	77.33	6.69	6.81	14.82	225.66	23.80	3230.66	Green with irregular white patches
Family	4	62.33	77.66	7.38	6.97	17.89	259.00	21.42	4113.00	Dark green with irregular white patches
Fai	5	63.66	78.33	6.64	6.30	13.16	229.33	19.16	3250.66	Green with irregular white patches
	6	64.66	80.66	6.20	5.87	11.16	215.00	20.89	2923.33	Green with irregular white patches
	7	66.33	79.33	6.84	6.67	14.91	231.66	19.69	3213.00	Green with irregular white patches
	CD @ 5 %	6.38	4.15	NS	1.04	NS	38.63	NS	1040.02	
	1	69.00	87.33	6.36	6.59	13.00	279.33	18.25	3654.33	Green with irregular white patches
ļi	2	73.66	87.00	6.20	6.45	12.75	266.00	23.80	3415.33	Dark green with irregular white patches
6 2	3	69.00	82.00	5.60	5.90	9.75	233.00	19.57	2526.66	Light green with narrow white patches
Family	4	70.00	84.66	6.44	6.48	13.58	260.33	17.72	3565.33	Dark green with irregular white patches
Fai	5	67.00	82.00	5.94	6.24	11.75	256.00	20.89	3338.33	Dark green with irregular white patches
	6	68.66	80.66	6.06	6.36	12.58	269.33	20.48	3467.33	Dark green with irregular white patches
	7	73.66	83.66	5.73	6.13	12.16	258.00	18.78	3242.66	Light green with irregular white patches
	CD @ 5 %	NS	5.70	NS	NS	NS	NS	5.71	NS	
	1	74.00	85.66	5.20	6.54	<u>11.00</u>	291.33	23.54	3343.00	Green with broad regular white patches
	2	75.66	85.33	5.42	6.80	12.08	300.66	20.89	3791.66	Green with scattered white patches
80	3	73.66	82.33	5.62	6.60	12.09	291.66	20.46	3960.66	Light green with scattered white patches
Family	4	72.66	84.66	5.31	6.47	11.41	305.33	19.31	3441.33	Light green with scattered white patches
Fai	5	72.66	82.00	5.41	5.81	10.50	274.33	19.84	3153.66	Green with scattered white patches
	6	73.33	81.33	5.42	6.91	12.33	321.66	21.16	4124.33	Green with irregular white patches
	7	73.33	82.66	5.74	6.98	12.74	317.33	19.42	4082.33	Green with broad prominent white streaks
	CD @ 5 %	NS	NS	NS	0.76	2.11	NS	2.89	NS	

.

.

# Table 4 continued...

Γ.	1	68.00	77.00	6.04	6.30	13.25	295.00	17.98	3725.33	Dark green with irregular white patches
	2	69.33	77.66	5.97	6.10	14.58	298.66	15.87	3755.00	Green with irregular white patches
6	3	68.33	75.00	6.79	6.27	13.50	301.00	19.31	3998.00	Light green with broad irregular white patches
Family	4	67.66	73.33	5.41	5.53	10.91	275.33	17.19	3159.00	Green with irregular white patches
Fai	5	67.66	74.00	6.51	6.51	13.75	322.66	16.93	3980.00	Dark green with irregular white patches
	6	67.66	75.33	5.90	5.67	11.25	263.00	18.51	3174.00	Green with scattered white patches
	7	68.00	74.66	5.93	6.01	11.75	287:66	16.40	3521.00	Light green with irregular white patches
	CD @ 5 %	NS	NS	0.61	NS	NS	NS	2.82	NS	
	1	75.00	90.00	6.81	5.65	11.62	376.33	17.72	4434.00	Light green with narrow white patches
	2	75.66	89.66	6.98	6.04	13.50	396.33	16.66	4917.66	Green with narrow white patches
2	3	73.00	90.66	6.66	5.89	11.20	373.66	17.98	4424.33	Light green with irregular white patches
	4	7300	90.33	6.85	5.89	13.66	400.66	17.72	5023.33	Light green with irregular white patches
Family	5	74.00	89.33	6.35	5.40	10.16	352.00	17.72	3665.33	Light green with irregular white patches
L CL	6	74.00	89.33	6.99	5.81	12.57	370.00	18.25	4297.00	Light green with irregular white patches
	7	74.33	91.66	6.91	6.03	12.41	375.66	17.98	4323.00	Light green with irregular white patches
	CD @ 5 %	NS	NS	0.62	NS	2.91	NS	NS	NS	
	1	80.00	97.33	6.48	7.03	14.97	310.66	17.19	3893.00	Green with irregular white patches
	2	87.00	98.66	7.20	6.21	16.33	319.66	20.21	3792.33	Green with irregular white patches
1	3	87.00	96.33	6.74	6.94	18.25	383.66	19.42	4035.66	Green with narrow white streaks
Family	4	83.00	95.00	6.39	6.17	12.67	238.66	19.31	3421.00	Green with narrow white streaks
Far	5	83.66	99.00	6.53	6.18	13.46	307.33	17.98	3562.66	Green, with narrow white streaks
	6	84.00	95.33	6.75	6.64	14.50	273.00	19.04	3466.66	Light green with irregular white patches
	7	85.33	96.33	5.75	6.04	11.09	232.00	17.98	3323.33	Light green with irregular white patches
	CD @ 5 %	6.78	NS	1.08	NS	NS	159.85	1.90	NS	

Table 4 continued...

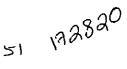
	nies in ent family			_						
CD @		9.63	6.36	1.27	0.84	4.54	99.63	4.00	1822.86	
	CD @ 5 %	5.55	NS	NS	NS	NS	NS	I.77	NS	
	7	88.33	101.33	6.44	6.09	10.83	232.00	16.66	3115.66	Light green, regular white patches with white spots
	6	88.66	100.33	6.12	5.85	10.54	211.00	15.34	3263.33	Light green, irregular white patches with white spots
Fami	5	90.33	100.33	6.64	6.19	10.75	225.66	16.40	2941.66	Light green, regular white patches with white spots
hily	4	87.66	102.00	5.97	5.63	10.00	213.33	17.19	2916.00	Light green, irregular white patches with white spots
17	3	89.00	101.00	5.60	5.43	9.25	212.66	17.19	3198.33	Light green, irregular white patches
	2	89.33	102.00	6.33	6.11	10.00	216.33	17.72	3121.66	Light green, regular white patches with white spots
	1	83.66	101.00	6.35	6.18	10.95	228.66	16.40	3115.00	Light green, regular white patches with white spots

.

-

.

.



was taken by the progeny 5 (90.33 days) followed by 2 (89.33 days) progenies. Days to first harvest showed no significant difference among the progenies. The mean values ranged from 100.33 to 102.00 days. Average fruit length showed no significant differences among the progenies. The mean values for this character ranged from 5.60 to 6.64 cm. Average fruit girth also showed no significant differences among the progenies. The mean values for this character ranged from 5.43 to 6.19 cm.

The average fruit weight showed non significance difference among the progenies. The mean values for this character ranged from 9.25 to 10.95 g. The number of fruits per plant also showed non significant difference among the progenies. The mean values for this character ranged from 211.00 to 232.00. Vitamin C content showed significant differences among the progenies. The mean values ranged from 15.34 to 17.72 mg/100g. The maximum vitamin C content was observed in the progeny 2 (17.72 mg/100g) followed by the progenies 3 (17.19 mg/100g) and 4 (17.19 mg/100g). The minimum vitamin C content was observed in the progeny 6 (15.34 mg/100g) which was on par with 1 (16.40 mg/100g), 5 (16.40 mg/100g) and 7 (16.66 mg/100g). The fruit yield per plant showed non significant difference among the progenies. The mean values ranged from 2916.00 to 3263.33 g.

# 4.1.3 Variability among the progenies of 12 different families (pooled analysis)

The pooled analysis of the data showed significant difference among the progenies for all the characters. The variations for days to first flowering ranged from 48.00 to 90.33 days. The minimum days to flowering was recorded in progeny 1 and 5 of family 1 followed by progeny 7 of the same family and progeny 1 of family 3 (48.33). The maximum time for flowering (90.33) was taken by progeny 5 of family 12. All progenies of family 12 took more than 83 days. All progenies of family 1 generally took less than 53 days for flowering.

The same trend was observed for days to first harvest of fruits. The progeny 1 of family 1 was found to be earliest to harvest within 65 days. It



was followed by three other progenies viz., progeny 2, 4 and 5 which took 66.66 days. The progenies 3 and 6 of the same family 3 and 6 took 67 days and progeny 7 took 67.33 days. The progenies of family 12 recorded more than 100 days for harvest. The maximum duration (102.00 days) was taken by progenies 2 and 4.

Significant variation for average fruit length was observed among the progenies. Progeny 1 of family 1 recorded the longest fruits having the length of 8.42 cm. All other progenies of the same family except progeny 6 produced fruits having more than 8 cm. The shortest fruits having an average length of 4.08 cm were produced by progeny 4 of family 4. The progenies of different families recorded a wide range of variation from 4.08 to 8.42 cm. The magnitude of variation was high for fruit girth. The progeny 4 of family 4 recorded the lowest fruit girth (5.17 cm). The highest fruit girth was observed in progeny 1 of family 5 (7.35 cm).

Average fruit weight was found to be highly variable from 5.59 to 21.91 cm among the progenies. The progeny 3 of family 3 produced fruits having the maximum weight followed by progeny 3 of family 1 (21.83).

Analysis of the pooled data revealed significant differences among the progenies for number of fruits. The mean values ranged from 208.66 to 577.00. The maximum number of fruits was produced by progeny 4 of family 1 and the minimum was recorded by progeny 2 of family 6.

Significant variation ranging from 13.75 to 24.45 mg/100g was recorded among the progenies for vitamin C content of fruits. Progeny 1 of family 4 recorded the highest vitamin C content and the lowest being in progeny 1 of family 5.

Fruit yield per plant showed significant variation ranging from 1489.00 to 8686.33g among the progenies of 12 families. The highest yield was recorded by progeny 4 of family 1 followed by progeny 1 of family 3 (8487.00) and progeny 3 (8465.00) of the same family.

#### 4.1.3 Coefficients of variation

The estimates of the components of variance viz., phenotypic and genotypic coefficients of variation are given in Table 5 fig.1.

Phenotypic coefficient of variation (PCV) was found to be slightly higher than genotypic coefficient of variation (GCV) for all the biometric characters studied.

The phenotypic coefficient of variation ranged from 7.78 to 44.06. The maximum phenotypic coefficient of variation was recorded for fruit yield per plant (44.06) followed by number of fruits per plant (29.35), average fruit weight (27.66), days to first flowering (17.97), average fruit weight (15.39), days to first harvest (12.17) and vitamin C content (10.29). The minimum PCV was observed for average fruit girth (7.78).

The genotypic coefficient of variation ranged from 7.15 to 42.17. The highest genotypic coefficient of variation was recorded for fruit yield per plant (42.17) followed by number of fruits per plant (26.93), average fruit weight (25.79), days to first flowering (16.26), average fruit length (14.14), days to first fruit harvest (11.71) and vitamin C content (8.23).The lowest GCV of variation was observed for average fruit girth (7.15).

#### 4.1.4 Heritability and Genetic advance

Heritability estimates ranged from 63.87 to 92.48 per cent (Table 5 fig.2). The highest heritability estimates was recorded for days to first harvest (92.48 per cent) followed by fruit yield per plant (91.59 per cent), average fruit weight (86.92 per cent), average fruit length (84.43 per cent), average fruit girth (84.36 per cent), number of fruits per plant (84.18 per cent) and days to first flowering (81.91 per cent ). The lowest heritability estimates was recorded for vitamin C content (63.87 per cent).

Genetic advance as per cent of mean ranged from 13.53 to 83.13. The highest genetic advance was recorded for fruit yield per plant (83.13 per cent) followed by number of fruits per plant (50.90 per cent), average fruit weight (49.53 per cent), days to first flowering (30.32 per cent), average fruit length (26.74 per cent), days to first harvest (23.20 per cent) and vitamin C content (13.55 per cent). The least genetic advance expressed for average fruit girth (13.53 per cent).

Table 5 Genetic components of variation

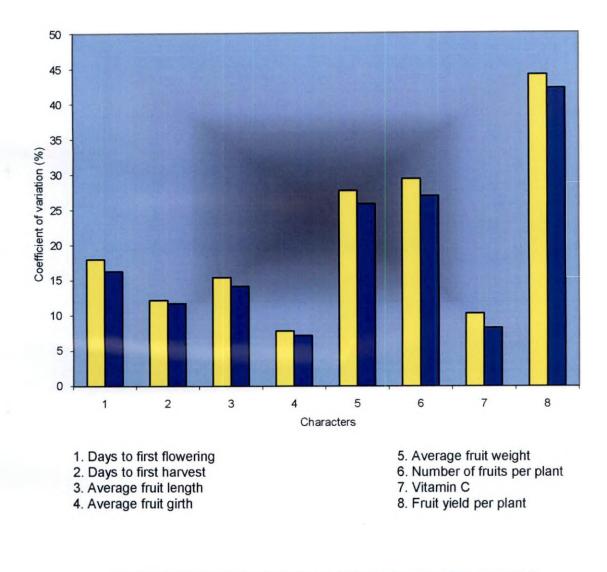
SI. no	Characters	PCV	GCV	Heritability (%)	Genetic advance (% of <sup>.</sup> mean)
1	Days to first flowering	17.97	16.26	81.91	30.32
2	Days to first harvest	12.17	11.71	92.48	23.20
3	Average fruit length (cm)	15.39	14.14	84.43	26.74
4	Average fruit girth (cm)	7.78	7.15	84.36	13.53
5	Average fruit weight (g)	27.66	25.79	86.92	49.53
6	No. of fruits per plant	29.35	26.93	84.18	50.90
7	Vitamin C	10.29	8.23	63.87	13.55
8	Fruit yield per plant (g)	44.06	42.17	91.59	83.13

#### 4.1.5 Correlation analysis

#### 4.1.5.1 Phenotypic correlation

Phenotypic correlation coefficients estimated for eight characters are presented in Table 6

Days to first flowering showed positive association (0.852) with days to first harvest and showed significant negative correlation with fruit yield per plant (-0.688), number of fruits per plant (-0.605), average fruit weight (-0.542) and average fruit length (-0.484). Negative association was noticed for days to first harvest with all other characters. Average fruit length showed positive correlation with average fruit weight (0.876),



Phenotypic coefficient of variation

## Fig. 1 Phenotypic and geotypic coeffficient of variation of eight characters of ivy gourd

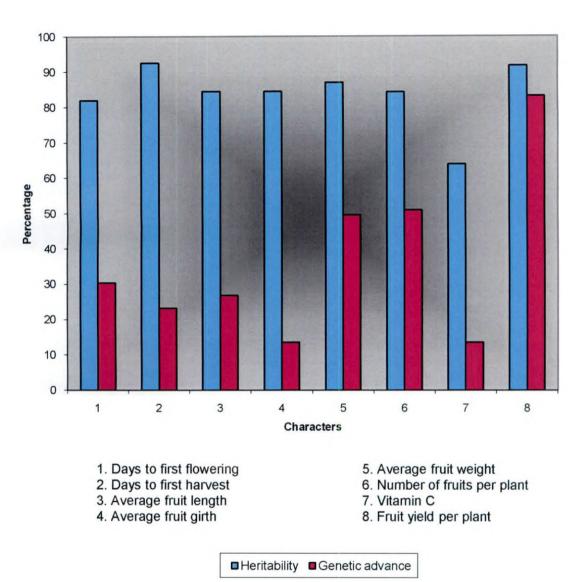


Fig. 2 Heriatability and genetic advance for eight characters in ivy gourd

fruit yield per plant (0.831), number of fruits per plants (0.612) and average fruit girth (0.592). Average fruit length showed negative phenotypic correlation with days to first flower (-0.484) and days to first fruit harvest (-0.307).

55

Average fruit girth showed significant positive correlation with average fruit weight (0.772), fruit yield per plant (0.638), and significant negative correlation was recorded with days to first fruit harvest (-0.487).

Average fruit weight recorded significant positive association with average fruit length (0.876), fruit yield per plant (0.836), average fruit girth (0.772) and number of fruits per plant (0.663) and significant negative correlation with days to first flowering (-0.542).

Fruit yield per plant recorded significant positive phenotypic correlation with average fruit weight (0.836), average fruit length (0.831), number of fruits per plant (0.663) and average fruit girth (0.638). Significant negative association was noticed with days to first flowering (-0.688) and days to first harvest (-0.497).

The number of fruits per plant recorded significant positive correlation with fruit yield per plant (0.872), average fruit weight (0.663) and average fruit length (0.612). Significant negative association was recorded with days to first flower (-0.605). The vitamin C content of the fruits had no significant correlation with any other characters.

#### 4.1.5.2 Genotypic correlation

The genotypic correlation for different characters is presented in Table 7.

The number of days to flower had significant positive genotypic correlation with days to first fruit harvest (0.929) and negative association with all other characters. Days to first harvest showed significant negative correlation with all other characters except days to first flower. Average fruit length showed significant positive genotype correlation with average fruit weight (0.963), fruit yield per plant (0.898), average fruit girth (0.726) and number of fruits per plant (0.717) and significant negative

### Table 6 Phenotypic correlation coefficient

Character	X1	X2	X3	X4	X5	X6	X7	X8
X1	1.000							
X2	0.852**	1.000	+					
X3	-0.484*.	-0.307	1.000					
X4	-0.430	-0.487*	0.592*	1.000				
X5	-0.542	-0.407	0.876**	0.772**	1.000			
X6	-0.688**	-0.497	0.831	0.638**	0.836**	1.000		
X7	-0.605*	-0.437	0.612*	0.414	0.663**	0.872**	1.000	
X8	-0.251	-0.286	-0.090	0.109	0.060	-0.090	-0.097	1.000

X 1 = Days to first flowering, X 2 = Days to first harvest, X 3 = Average fruit length, (cm) X 4 = Average fruit girth (cm)

X 5 = Average fruit weight (g) X 6 = Fruit yield per plant (g) X 7 = Number of fruits per plant X 8 = Vitamin C

.

### Table 7 Genotypic correlation coefficient

Character	<b>X</b> 1	X2	X3	X4	X5	X6	X7	X8
X1	1.000	+	+		+			
X2	0.929**	1.000						
-X3	-0.588*	-0.372	1.000	<u> </u>				
X4	-0.608	-0.559*	0.726**	1.000				
X5	-0.639**	-0.496	0.963**	0.866	1.000			
X6	-0.780**	-0.566	0.898**	0.699	0.887**	1.000		
X7	-0.738	-0.549	0.717**	0.453	0.700**	0.902**	1.000	
X8	-0.364	-0.372	-0.081	0.216	0.038	-0.088	-0.084	1.000

X 1 = Days to first flowering, X 2 = Days to first harvest, X 3 = Average fruit length (cm), X 4 = Average fruit girth (cm),

X 5 = Average fruit weight (g), X 6 = Fruit yield per plant (g), X 7 = Number of fruits per plant, X 8 = Vitamin C (mg/100g)

			X4	X5	X6	X7	X8
1.000							
0.371	1.000						<u> </u>
0.028	0.193	1.000					
0.449	0.058	-0.127	- 1.000				
-0.019	0.376	0.358	0.214	1.000			<u> </u>
-0.097	0.303	0.360	0.207	0.425	1.000		
0.045	0.430	0.050	0.206	0.443	0.689**	1.000	
0.047	-0.002	-0.128	-0.208	0.144	-0.126	-0.148	1.000
	0.371 0.028 0.449 -0.019 -0.097 0.045	0.371       1.000         0.028       0.193         0.449       0.058         -0.019       0.376         -0.097       0.303         0.045       0.430	0.371         1.000           0.028         0.193         1.000           0.449         0.058         -0.127           -0.019         0.376         0.358           -0.097         0.303         0.360           0.045         0.430         0.050	0.371       1.000         0.028       0.193       1.000         0.449       0.058       -0.127       1.000         -0.019       0.376       0.358       0.214         -0.097       0.303       0.360       0.207         0.045       0.430       0.050       0.206	0.371       1.000         0.028       0.193       1.000         0.449       0.058       -0.127       1.000         -0.019       0.376       0.358       0.214       1.000         -0.097       0.303       0.360       0.207       0.425         0.045       0.430       0.050       0.206       0.443	0.371       1.000	0.371       1.000

 Table 8 Environmental correlation coefficient

X 1 = Days to first flowering, 2 = Days to first harvest, X 3 = Average fruit length (cm), X 4 = Average fruit girth (cm),

X 5 = Average fruit weight (g), X 6 = Fruit yield per plant (g), X 7 = Number of fruits per plant, X 8 = Vitamin C (mg/100g)

correlation was observed with days to first flower (-0.588) and non significant negative correlation with days to first harvest (-0.372).

Positive genotypic correlation was recorded for average fruit girth with average fruit weight (0.866), average fruit length (0.726) and fruit yield per plant (0.699) and significant negative correlation was recorded with days to first flower (-0.608) and days to first harvest (-0.559).

Average fruit weight recorded significant positive genotypic correlation with average fruit length (0.963), fruit yield per plant (0.0887), average fruit girth (0.866) and number of fruits per plant (0.700). Significant negative correlation was recorded with days to first flower (-0.639) and days to first harvest (-0.496).

Fruit yield pert plant showed significant positive genotypic correlation with number of fruits per plant (0.902), average fruit length (0.898), average fruit weight (0.887) and average fruit girth (0.699). Significant negative genotypic correlation was showed with days to first flower (0.780) and days to first fruit harvest (-0.566).

Significant positive genotypic correlation was recorded for number of fruits per plant with fruit yield per plant (0.902), average fruit length (0.717) and average fruit weight (0.700). Significant negative correlation was observed with days to first flower (-0.738) and days to first harvest (-0.549). There was no significant correlation for vitamin C content with any other characters.

The estimates of the environmental correlation coefficient of variation presented in Table 8 showed no significant correlation except for fruit yield per plant.

#### 4.1.6 Path analysis

The direct and indirect effect of different characters on yield is presented in Table 9 and Fig. 3.

Fruit yield per plant had positive correlation with length of fruit (0.898), girth of fruit (0.699), average fruit weight (0.887) and number of fruits per plant (0.902). Vitamin C content of fruit had negative

### Table 9 Direct and indirect effects

.

	X1	X2	X3	X4	X5	X6	X7	Correlation
X 1	0.088	-0.169	-0.783	-0.378	0.884	-0.438	0.017	-0.780
X 2	0.082	-0.182	-0.495	-0.348	0.686	-0.326	0.018	-0.566
X 3	-0.052	0.067	1.333	0.452	-1.333	0.426	0.003	0.898
X 4	-0.054	0.101	0.967	0.623	-1.198	0.269	-0.010	0.699
X 5	-0.056	0.090	1.283	0.539	-1.384	0.416	-0.001	0.887
X 6	-0.065	0.100	0.955	0.282	-0.968	0.594	0.003	0.902
X 7		0.067	-0.107	0.134	-0.052	-0.049	-0.047	-0.088

Residual effect = 0.1448

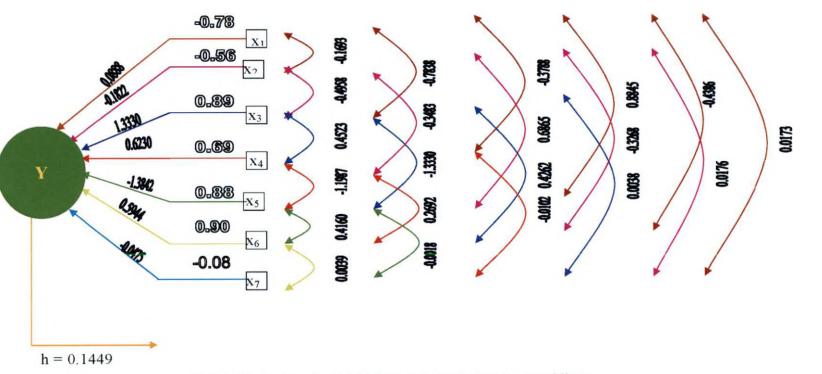
X 1 = Days to first flowering

X 2 = Days to first harvest

- X 3 = Average fruit length (cm)
- X 5 = Average fruit weight (g)

- X 4 = Average fruit girth (cm)
- X 6 = Number of fruits per plant

X 7 = Vitamin C (mg/100g)



Direct effects given in straight lines and correlations in curved lines

- X1Days to first floweringX4Average fruit girthX2Days to first harvestX5Average fruit weightX3Average fruit lengthX6Number of fruits per plant
  - X7 Vitamin C

#### Fig. 3 Path diagram showing direct and indirect effects of components on yield

correlation with yield. The average length of fruit has the maximum direct effect on yield (1.333) followed by girth of fruits (0.623), number of fruits per plant (0.594). However days to first fruit harvest and vitamin C content recorded negative direct effect with fruit yield.

Number of days to flower  $(x_1)$  had positive direct effect (0.088) and negative correlation (-0.780). Number of days to first harvest  $(x_2)$  and vitamin C content of fruits  $(x_7)$  recorded negative genotypic correlation (-0.169) and negative direct effect (-0.047) respectively. Average weight of fruit  $(x_5)$  recorded negative direct effect (-1.384) and positive genotypic correlation (0.884) and it had high indirect effect through average length of fruits  $(x_3)$ , average girth of fruits  $(x_4)$  and number of fruits per plant  $(x_6)$ . The direct effect of number of fruits per plant  $(x_6)$  is positive (0.594) which is indirectly contributed by length of fruits  $(x_3)$  and girth of fruits  $(x_4)$ .

The average length of fruits recorded the highest direct effect (1.333) and it had positive correlation (0.898). The residual effect (r = 0.1448) indicated that the selected characters explain the total correlation well and the remaining characters have minor contributions on the variability of yield.

#### 4.1.7 Standard heterosis

The standard heterosis was estimated in comparison with the standard check variety "Sulabha" for different characters are presented in Table 10 and Fig. 4

#### Days to first flowering

Days to first flowering showed positive heterosis among the families ranging from 2.52 to 81.79 per cent. The family  $F_1$  recorded the lowest percentage (2.52) followed by the families  $F_3$  (4.50 per cent) and  $F_2$  (23.30 per cent). The maximum percentage was expressed by the family  $F_{12}$  (81.80 per cent).

#### Days to first harvest

Positive heterosis for days to first fruit harvest ranged from 2.10 to 56.30 per cent. The minimum percentage was recorded by the family  $F_1$  (2.09 per cent) followed by the families  $F_9$  (15.52 per cent),  $F_2$  (18.35 per cent),  $F_3$  (19.44 per cent),  $F_5$  (20.05 per cent) and  $F_6$  (20.43 per cent). The maximum percentage was expressed by the family  $F_{12}$  (56.30 per cent).

# Average fruit length

Two families viz,  $F_1$  (6.11 per cent) and  $F_3$  (3.56 per cent) recorded positive heterosis for average fruit length. The negative heterosis ranged from -9.92 to -38.04 per cent. The maximum negative heterosis was observed in the family  $F_4$  (-38.04 per cent) followed by the families  $F_8$  (-29.01 per cent),  $F_9$  (-22.01 per cent),  $F_7$  (-21.76 per cent),  $F_{12}$  (-19.85 per cent) and  $F_5$  (-14.75 per cent). The minimum negative heterosis was recorded in the family  $F_2$  (-9.92 per cent) followed by the families  $F_{10}$  (-12.21 per cent),  $F_6$  (-13.36 per cent) and  $F_{11}$  (-13.74).

#### Average fruit girth

The positive heterosis ranged from 0.45 to 11.96 per cent for average fruit girth. The maximum percentage was observed in the family  $F_1$  (11.96 per cent) followed by the families  $F_5$  (8.69 per cent),  $F_2$  (8.23 per cent),  $F_3$  (6.67 per cent) and  $F_6$  (5.12 per cent). The minimum positive heterosis was observed in the family  $F_7$  (0.45 per cent) followed by the families  $F_{11}$  (2.47 per cent) and  $F_8$  (2.78 per cent). The negative heterosis for this character ranged from to -5.15 to -10.74 per cent. The maximum negative heterosis was observed in the family  $F_4$  (-10.74 per cent) followed by  $F_{10}$  (-9.03 per cent) and  $F_{12}$  (-6.23 per cent). The minimum negative heterosis was observed in the family  $F_9$  (-5.15 per cent).

#### Average fruit weight

The positive heterosis for average fruit weight was observed in the families  $F_1$  (17.42 per cent) and  $F_3$  (12.60 per cent). The negative heterosis for this character ranged from -6.08 to -58.26 per cent. The maximum negative heterosis was recorded in the family  $F_4$  (-58.26 per cent) followed by the families  $F_{12}$  (-40.81 per cent),  $F_8$  (-31.04 per cent),  $F_{10}$  (-29.03 per cent),  $F_7$  (-26.95 per cent),  $F_9$  (-26.50 per cent),  $F_5$  (-17.25 per cent) and  $F_6$  (-16.18 per cent). The minimum negative heterosis was observed in the family  $F_2$  (-6.08 per cent) followed by the family  $F_{11}$  (-6.81 per cent)

#### Number of fruits per plant

Four families recorded positive heterosis for number of fruits per plant which ranged from 7.31 to 40.58 per cent. The maximum positive heterosis was observed in the family  $F_1$  (40.58 per cent) followed by  $F_2$ (30.98 per cent),  $F_3$  (18.44 per cent) and  $F_{10}$  (7.31 per cent). The negative heterosis for this character ranged from -11.16 to -38.35 per cent. The maximum negative heterosis was observed in the family  $F_{12}$  (-38.35 per cent) followed by  $F_6$  (-32.65 per cent),  $F_5$  (-27.66 per cent),  $F_7$  (-26.06 per cent),  $F_4$  (-17.85) and  $F_9$  (-16.19 per cent). The minimum negative heterosis was observed in the family  $F_{11}$  (-11.16 per cent) followed by the family  $F_8$  (-15.48 per cent).

#### Vitamin C

The positive heterosis for vitamin C content was ranged from 0.45 to 17.73 per cent. The maximum positive heterosis was observed in the family  $F_6$  (17.73 per cent) followed by the families  $F_1$  (16.78 per cent),  $F_7$  (16.39 per cent),  $F_8$  (16.17 per cent),  $F_4$  (15.17 per cent) and  $F_{11}$  (5.52 per cent). The minimum positive heterosis was observed in the family  $F_{10}$  (0.45 per cent) followed by the families  $F_5$  (2.85 per cent). The negative heterosis was recorded by three families which ranched from -1.17 to -9.03 per cent. The maximum negative heterosis was recorded in the family  $F_2$  (-9.32 per cent) followed by the families  $F_{12}$  (-6.24 per cent) and  $F_9$  (-1.17 per cent).

#### Fruit yield per plant

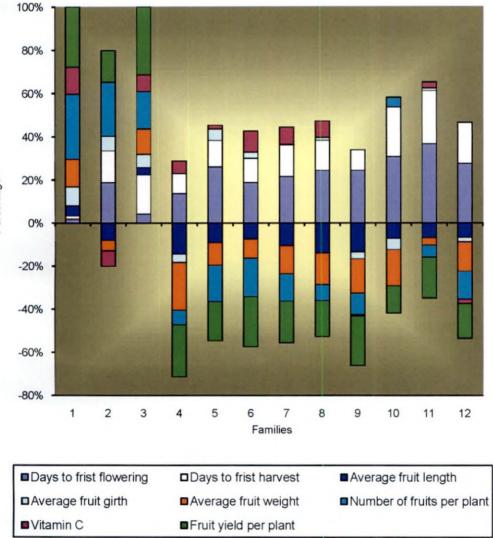
The positive heterosis for fruit yield per plant was observed in the families  $F_1$  (37.76 per cent),  $F_3$  (33.60 per cent) and  $F_2$  (18.34 per cent).

The negative heterosis for this character ranged from -21.84 to -63.93 per cent. The maximum negative heterosis was observed in the family  $F_4$  (-63.93 per cent) followed by the families  $F_{12}$  (-47.80 per cent),  $F_6$  (-42.36 per cent),  $F_7$  (-39.93 per cent),  $F_{11}$  (-38.57 per cent) and  $F_9$  (-35.01 per cent). The minimum negative heterosis was observed in the families  $F_{10}$  (-21.84 per cent) and  $F_5$  (-29.48 per cent).

Families	Days to	Days to	Average	Average	Average	Number of	Vitamin C	Fruit
	first	first	fruit length	fruit girth	fruit	fruits per		yield per
	flowering	harvest			weight	plant		plant
F <sub>1</sub>	2.52*	2.10*	6.11	11.96*	17.43	40.58	16.78	37.76*
F <sub>2</sub>	23.31	18.35	-9.92	8.23	-6.08	30.98	-9.03	18.34*
F <sub>3</sub>	4.50	19.44	3.56	6.67	12.60	18.44	8.31	33.60
F <sub>4</sub>	36.18	24.54	-38.04	-10.74	-58.26	-17.85	15.17	-63.93
F5	42.25	20.05	-14.36	8.69*	-17.25	-27.66	2.85	-29.48
F <sub>6</sub>	34.03	20.43	-13.36	5.12*	-16.18	-32.65	17.73	-42.36
F <sub>7</sub>	44.26	30.29	-21.76	0.45	-26.95	-26.06	16.39	-39.93
F <sub>8</sub>	51.18	28.95	-29.01	2.78*	-31.05	-15.48	16.17	-35.01
F9	40.00	15.52	-22.01	-5.15	-26.50	-16.19	-1.17	37.59
F10	52.69	39.54	-12.21	-9.03	-29.03	7.31	0.45	-21.84
F11	73.55	49.14	-13.74	2.47*	-6.81	-11.16	5.52*	-38.57
F <sub>12</sub>	81.80*	56.30	-19.85	-6.24	-40.81	-38.35	-6.24	-47.80

Table.10 Standard heterosis (%) for eight characters of 12 families of ivy gourd

.



# Fig. 4 Standard heterosis among the 12 families of ivy gourd

Percentage

# **DISCUSSION**

•

.

#### 5. DISCUSSION

The results of the experiment conducted on "Development and evaluation of hybrids in ivy gourd (*Coccinia grandis* L. Voigt.) are discussed in this chapter.

# **Evaluation of hybrids**

Ivy gourd is a typical dioecious crop. The tender parthenocarpic fruits are used as vegetable. Being a clonally propagated crop conventional biometric approach to study heterosis combining ability and gene action is not possible especially for the yield and yield attributes. However viable seeds can be produced with help of male parents.

Morphologically distinct male plants have been identified in different ivy gourd growing tracts. In this investigation attempts were made to study the variation induced by hybridization between different female parents and morphologically distinct male parents. Ivy gourd is highly heterozygous because it is cross pollinated in nature. A desirable attribute of clonally propagated plants amenable to seed production is fixation of heterosis in  $F_1$  generation.

#### Variability among the F<sub>1</sub> families

The analysis of variance conducted for 12  $F_1$  families of ivy gourd showed significant differences among the progenies for the different characters studied. This clearly showed that the families are different from each other. The compact family block design used for the conduct of experiment provides an opportunity to assess the variability among and within the families. Since fixation of heterosis is possible in  $F_1$ generation, identification of superior hybrid progeny is useful for developing a hybrid clone.

There was significant variation among the families for days to first flowering which is evident from the range of variation showed for this character. Early flowering is a desirable attribute. Two families  $F_1$  and  $F_2$  produced using the same female parent was significantly different from the rest for days to flowering. However the other two crosses produced using the same female parent did not show any significant difference in the days taken for first flower opening. The family 12 took the maximum number of days for flowering. Joseph (1999) and Varghese (2003) also reported a wide range of variation for days to flowering in ivy gourd.

The family 1 which took the minimum days for flowering also took minimum days for first harvest. The family 12 took maximum days for flowering and days for first harvest of fruit.

There were significant differences among the families for average fruit length and girth of fruits. The family 1 recorded the maximum length and girth of fruits.

The number of fruits per plant showed significant differences among the families. The family 1 (Royappanpatty 1 x Royappanatty local)produced the maximum number of fruits and was on par with family 1 (Royappanpatty 1 x Vellayani local). The family 1 was found to be the best to give the highest fruit yield. The family 12 recorded the lowest fruit yield per plant. The vitamin C content of the fruits also showed significant difference among the families. Among the twelve families family 1 was found to be best for seven out of eight characters. Hence it is possible to select plants from this family for desirable traits.

#### Variability among the progenies

The mean values of progenies for different characters showed wide variation among the progenies of the same parentage. The compact family block design facilitates the analysis of progeny of different families.

The variation for different characters among the progenies in family 1 was not significant. It indicates the homogeneity of the progenies. The progenies of family 1 had desirable attributes like early flowering, early fruit harvest, longest fruits, more number of fruits and fruit yield per plant. Hence all progenies of family 1 can be selected for further evaluation. But significant variation was observed for days to first flowering and vitamin C content in Family 2. The  $F_1$  population is highly heterozygous and heterogeneous in nature. The progeny 7 of the family 2 took the minimum days for the first harvest of fruits. Variation within the family was observed for other progenies 1 and 2 were significantly different from other progenies for fruit length. The progeny 1 was found to have the maximum fruit girth. Fruit length and fruit girth are desirable traits.

The progenies having desirable traits can be selected from this family.

The variations showed for the number of fruits and fruit yield per plant was found to be not significant. Hence all the progenies can be selected for getting more number of fruits and better fruit yield. The variations among the progenies were significant for days to first flowering and vitamin C content. Among the progenies of family 3 (Royappanpatty 1 x Karyavattam) significant variations was observed for days to flowering. The progeny 1 was the earliest to flower. For fruit length progeny 3 was significantly superior to the rest. The progenies in general had desirable traits viz., fruit girth, weight, number of fruits and fruit yield per plant for which no significant difference was noticed among the progenies. Hence all the progenies can be utilized for further improvement programmes. Selection of progeny 3 is desirable of early flowering.

Maximum heterogeneity among the progenies was observed in family 4. Significant variations were recorded for all the characters except for average fruit length. Progenies for various desirable characters can be selected from this family. In family 4, progeny 2 took the minimum days for first flowering. However early harvest was taken from progeny 7 of the same family. Among the different progenies in family 4, progeny 1 had the maximum girth of fruit. The same progeny also recorded the maximum weight and number of fruits and yield per plant. Progeny 1 can be selected from family 4 because of the desirable yield attributes for further evaluation.

Among the different progenies of family 5 significant variations was recorded for days to first flowering, days to first fruit harvest, average fruit weight, vitamin C content and fruit yield per plant. The progeny 1 took minimum days for flowering followed by progeny 2. Regarding the early harvest of fruits progeny 5 came first followed by progeny 7. The variations among the progenies for fruit length and fruit girth were not significant. Average fruit weight showed significant differences among the progenies. Progeny 1 of family 5 recorded the maximum fruit weight. Significant variation was observed for vitamin C content of fruits. Progeny 7 of family 5 had the maximum content of vitamin C. Significant variations among the progenies of family 5 were recorded for fruit yield per plant. The progeny 6 recorded the maximum fruit yield and can be selected.

The progenies of family 6 showed significant variations for days to first flower but the variation was not significant for days first fruit harvest. The progenies showed significant variation for average fruit length. Progeny 4 recorded the highest value for average fruit length. Average fruit girth also showed significant variation among the progenies. Progeny 1 recorded the highest fruit girth.

There was significant variation among the progenies in family 6 for fruit yield per plant. Progeny 4 recorded the maximum fruit yield per plant. The same progeny also had the maximum girth of fruits. Progeny 4 can be selected from family 6 for further evaluation.

Heterogeneity for different characters was less in family 7. Out of the eight characters significant differences among the progenies were observed only for days to first fruit harvest and vitamin C content. Progeny 6 took the minimum days for fruit harvest. For vitamin C content progeny 2 was found to be the best.

The variations shown among the progenies in family 8 were not significant for day to first flowering, day to fruit harvest, average length of fruit, number of fruits per plant and fruit yield per plant. Average girth of fruit showed significant differences among the progenies. Progeny 7 had the maximum average fruit girth which was on par with progeny 6. Average fruit weight had significant differences among the progenies. The maximum average fruit weight was recorded in the progeny 7. Progeny 7 can be selected for maximum length and girth of fruits.

Significant variation for average fruit length and vitamin C content was observed among the progenies in family 9. The average fruit length was the maximum in progeny 3 which was on par with the progeny 5 and 1. The progeny 3 had the highest content of vitamin C in family 9. Within family variation for other characters were not significant.

Family 10 showed significant variation among progenies for fruit length and fruit weight. The progeny 6 produced fruits with the maximum average length whereas progeny 4 produced fruits having the maximum average weight. The variation shown for other characters were not found to be significant.

The family 11 showed significant differences among the progenies for four characters viz., days to first flowering, average fruit length, number of fruits per plant and vitamin C content. The progeny 1 took the minimum number of days for flowering. The progeny 2 produced fruits having the maximum average length. The progeny 3 produced the maximum number of fruits. There was no significant difference among the progenies for other characters.

Significant variation among the progenies for days for first flowering and vitamin C content was noticed in family 12.

The magnitude of variation shown among the progenies of 12 families revealed the heterogeneous nature of the  $F_1$  families. Significant variations among the progenies for vitamin C content was expressed in eight families viz., family 3, 4, 5, 7, 9, 11 and 12. Seven families viz., family 2, 3, 4, 5, 6, 11 and 12 showed significant

variations among progenies for days to first flowering. Variations among progenies for fruit length were expressed by the families 2, 3, 4, 6, 9, 10, and 11. The families 4, 5, 8 and 10 showed significant variations for average fruit length among the progenies. The families 4, 5 and 7 showed variations among progenies for days to first fruit harvest. The families 2, 6 and 8 showed variations among the progenies for girth of fruits. Families 4, 6 and 11 recorded significant variations among the progenies for number of fruits. Only two families viz., families 4 and 6 showed significant variations among progenies for fruit yield per plant.

#### Variability among the progenies of 12 different families (pooled analysis)

The pooled analysis of the data showed significant difference among the progenies for all the characters. The minimum days to flowering was recorded in progeny 1 and 5 of family 1 followed by progeny 7 of the same family and progeny 1 of family 3. The maximum time for flowering was taken by progeny 5 of family 12. The same trend was observed for days to first harvest of fruits. The progeny 1 of family 1 was found to be earliest to take harvest. It was followed by three other progenies viz., progeny 2, 5 and 6.

Significant variation for average fruit length was observed among the progenies. Progeny 1 of family 1 recorded the longest fruits. The shortest fruits were produced by progeny 4 of family 4. The progeny 4 of family 4 recorded the lowest value for fruit girth. The highest fruit girth was observed in progeny 1 of family 4. The progeny 3 of family 3 produced fruits having the maximum weight followed by progeny 3 of family 1 Analysis of the pooled data revealed significant differences among the progenies for number of fruits. The maximum number of fruits was produced by progeny 4 of family 1 and the minimum was recorded by progeny 2 of family 6.

Significant variation was recorded among the progenies. Progeny 1 of family 4 recorded the highest vitamin C content and the lowest being in progeny 2 and 4 of family 4.

The highest fruit yield was recorded by progeny 3 of family 1 followed by progeny 1 of family 3 and progeny 3 of the same family. Most of the families showed highly heterogeneous nature for different characters. Therefore superior progenies identified for different characters from the hybrid population can be utilized for further improvement. The advantage of clonal propagation and fixation of heterosis in the  $F_1$  generation promotes the development hybrid

clones in ivy gourd. In the present investigation seedling progenies were analysed. Cuttings from the selected plants can be used for growing the next generation for the evaluation of hybrid clones.

#### Variability and Genetic parameters

The results of the analysis of variance showed significant differences among the genotypes for all the characters studied viz., days to first flowering, days to first fruit harvest, average length of fruits, average girth of fruits, average weight of fruits, number of fruits per plant, vitamin C content and fruit yield per plant.

The genetic parameters of different characters in 12 families of ivy gourd were estimated in this study. In ivy gourd heterosis can be fixed in  $F_1$  generation because of the clonal propagation. Selection of plants in  $F_1$  population helps to identify desirable clones. Selection of desirable plants from a genetically variable population is the basic step in crop improvement. The success of selection and extent of progress in a selection programme depends upon the magnitude of variability existing in a crop. In crops like ivy gourd hybridization is one of the methods to induce variability.

A wide range of variation observed among the hybrid population in this study indicated the scope for selection. The variability available in a population could be partitioned into heritable and non-heritable components with the aid of genetic parameters like genotypic coefficient of variation (GCV), heritability ( $h^2$ ) and genetic advance which serve as useful guidelines for selection.

The phenotypic coefficient of variation (PCV) was found to be higher than genotype coefficient of variation for all the characters studied. Bhave et al. (2002) in bitter gourd and Choudhary and Sharma (2002) in ridge gourd also reported that phenotypic coefficient of variation was higher than GCV for all the characters they studied. High estimates of PCV and GCV were observed for fruit yield per plant followed by number of fruits per plant. Suresh (2004) also reported the highest PCV and GCV for fruit yield per plant followed by number of fruits per plant in ivy gourd. Similar reports were also reported by Kandaswamy (2004) in muskmelon. Joseph (1999), Varghese (2003) and Wilson et al. (2006) also reported high PCV and GCV for fruit yield per plant. High PCV and GCV for fruit yield per plant was also reported by Gayathri (1997), Rastogi and Deep (1990) in cucumber, Swamy and Dutta (1985) in musk melon, Vahab (1989) in bitter gourd, Varghese (1991), Varghese and Rajan (1993) in snake gourd. Parmer and Tarsen Lal (2005) reported the highest PCV and GCV for number of fruits per plant in muskmelon. The phenotypic and genotypic coefficients of variation were found to be low for average fruit girth whereas GCV for average weight of fruit (27.66) was high. Mathew et al. (2001) reported high GCV values (22.40) for average fruit weight in bottle gourd. The GCV for fruit length estimated in the study was 14.14. In bottle gourd Mathew et al. 2001 reported a higher estimate (24.80) of GCV for fruit length. The GCV for days to first flowering was 16.26. However Gayathri (1997) in cucumber, Varghese and Rajan (1993) in snake gourd and George (1981) in ash gourd reported the low estimates of GCV for days to first flowering.

The GCV was comparatively low for days to first harvest. Similar results were also reported by Varghese and Rajan (1993) in snake gourd.

The vitamin C content of fruit also recorded low estimates PCV and GCV values. Varghese (2003) reported that the variability for vitamin C content of fruit is less than 20 per cent.

The lowest PCV and GCV values were estimated for average fruit girth. Tyagi (1972) reported a high estimate of GCV for girth of fruits in bitter gourd.

The GCV provides a valid basis for comparing and assessing the range of genetic variability and PCV measures the extent of total variation. High values of PCV with correspondingly high values of GCV observed for fruit yield per plant, number of fruits per plant and average fruit weight indicated the presence of high degree of genetic component of variation and better scope for improvement through selection. Low estimates of GCV for average fruit girth of fruits, vitamin C content of fruits and days to first harvest indicated limited scope for improvement of these traits.

For all the characters PCV and GCV values were found to be closer indicating the predominant influence of genetic component over the environmental effect for these characters.

#### Heritability and Genetic advance

Heritability estimates provide exact and precise information of the influences of environment on various characters. Robinson et al. (1966) classified heritability into low (< 30 per cent), medium (30-60 per cent) and high (> 60 per cent). Burton (1952) suggested that GCV along with heritability would provide a picture of the account of advance to be expected by genotype selection.

High heritability was recorded for all the characters in this study. The highest heritability of 92.48 per cent was recorded for days to first harvest followed by fruit yield per plant, average fruit weight, average fruit length, average fruit girth, number of fruits per plant, days to first flowering and vitamin C content. The high heritability of more than 60 per cent recorded for all the characters may be due to the fact that these characters are least influenced by the environment. This is in line with the findings of Suresh (2004) in ivy gourd. Johnson et al. (1955) pointed out that high heritability estimates along with high genetic advance were more useful than the heritability estimates alone in predicting the resultant effect for selecting the best plant.

High heritability estimates were obtained for fruit yield per plant followed by number of fruits per plant, average fruit weight and days to first flowering. The lowest heritability estimates were recorded for vitamin C content followed by average girth of fruits, days to first flowering, days to first harvest and average fruit length.

High heritability and high genetic advance estimated for fruit yield per plant, number of fruits per plant and average fruit weight in this study is in line with the findings of Suresh (2004) in ivy gourd. The lowest genetic advance recorded for average fruit girth in the study is in conformity with the reports of Wilson et al. (2006) in the same crop and Varalakshmi et al. (1955) in ridge gourd and Sarkar et al. (1990) in pointed gourd. However Lovely and Radhadevi (2007) reported high heritability coupled with high genetic advance for fruit girth in ash gourd.

Gayathri (1997) in cucumber and Anitha (1998) in ridge gourd has also reported high genetic advance for yield per plant, average fruit weight and number of fruits per plant.

Gayathri (1997) in cucumber, Anitha (1998) in ridge gourd, Mathew and Khader (1999) in snake gourd and Bindu et al. (2000) in pumpkin also reported high heritability and genetic advance for fruit yield per plant, average fruit weight and number of fruits per plant.

The differences between PCV and GCV were very low for all the characters and indicated the major contribution genetic component of variation. High heritability estimates coupled with high genetic advance for important characters like fruit yield per plant, average fruit weight and number of fruits per plant indicated the effectiveness selection of for the improvement of these characters.

#### Correlation analysis

Yield is a complex character influenced by a number of other component characters. For successful improvement through selection it is essential to ascertain the importance and extent of inter association with yield. The extent of relationship between yield and its component traits as well as among the component traits is revealed through correlation analysis.

The genotypic correlation between the characters provides a reliable measure of genetic association between the characters and helps to differentiate the vital association useful in breeding from the non-vital ones (Falconer, 1981)

The genotypic correlations were higher than the phenotypic correlation for most of the characters indicating the reduced influence of environment for variation of characters. Saroj Rohana et al. 2003 also obtained similar results in water melon. Fruit yield per plant showed significant positive genotypic correlation with number of fruits per plant, average fruit length, average fruit weight and average fruit girth. This is in line with the findings of Joseph (1999) and Sarnaik et al. (1999) in ivy gourd. Similar results were also reported by Rastogi and Deep (1999), Ma et al. (1995) and Rao et al. (2004) in cucumber, Lawande and Patil (1989) in bitter gourd, Shibukumar (1995) in water melon, Singh et al. (1993) and Sarkar et al. (1999) in pointed gourd. Positive correlation indicates the importance of these traits in improving fruit yield.

Fruit yield per plant was found to be negatively correlated with days to first flowering and days to first harvest. Varghese (2003) and Suresh (2004) also reported the negative correlation of days to first flowering with fruit yield. Similar results were also reported by Rastogi and Deep (1990) and Damarany et al. (1995) in cucumber, Lawande and Patil (1990) in bitter gourd, Shibukumar (1995) in water melon and Prasad and Singh (1990) in pointed gourd. However, positive correlation of yield with days to flower was also reported by Pynadath (1978) in snake gourd and Anitha (1998) in ridge gourd. Negative correlation of yield with days to first fruit harvest was also reported by Anitha (1988) in ridge gourd.

Number of fruits per plant showed significant positive correlation with fruit yield per plant, average fruit length and average fruit weight. From these positive associations it can be inferred that simultaneous improvement of these characters is possible. Average fruit weight also had significant positive correlation with average fruit length, average fruit girth and fruit yield per plant. Vitamin C content of fruit did not show any significant correlation with yield or any other characters.

#### Path analysis

As the correlation coefficients are insufficient to explain the cause and effect relationship among the traits for an effective manipulation of the characters, the path analysis was carried out. The path analysis furnishes a method of partitioning the correlation coefficients into direct and indirect effects and provides the actual contribution of an attribute and its influence through other traits.

Average length of fruits, girth of fruits, average weight of fruits and number of fruits per plant exerted significant positive direct effect on fruit yield per plant. Therefore improvement of anyone of these characters can bring out simultaneous improvement of yield. This is in conformity with the findings of Varghese (2003) in ivy gourd. Positive direct effect of fruit weight and fruit length on yield had been reported by Prasad and Singh (1992) in cucumber. Positive direct effect of number of fruit and weight of fruits has also been reported by Saika et al. (1995) and Gayathri (1997) and Rao et al. (2004) in the same crop and Bhave et al. (2003) in bitter gourd. Days to flowering had negative correlation and positive direct effect in this study. Shibukumar (1995) in water melon and Dhaliwal et al. (1996) in musk melon reported positive direct effect of days to flowering on yield.

#### Standard heterosis

The families showed standard heterosis at varying levels for different characters. In this study mean values of the families were taken for comparison with the standard parent. The progenies were found to be highly heterogeneous. Since heterosis can be fixed in  $F_1$  generation due to clonal propagation superior plants having desirable traits can be selected for further evaluation.

Early flowering is a desirable trait. Compared to the standard check "Sulapba" the hybrid population took more days to flower. However among the hybrid progenies family 1 took the minimum days to flower. Family 12 recorded 81.8 per cent heterosis which is not a desirable attribute. Gayathri (1997) in cucumber also reported heterosis for days to flower, whereas Ram et al. (1997) observed negative heterosis for days to flower.

The family which took minimum days to flower was also found to be the earliest to harvest. All the hybrid progenies showed positive heterosis for days to first fruit harvest. Similar result was also reported by Varghese and Rajan (1993b). But, Mandal and Sirohi (2002) observed heterosis for early maturity in ash gourd.

The magnitude of heterosis for length and girth of fruits were low. Only two families manifested marginal positive heterosis for fruit length. All other families expressed negative heterosis. Significant heterosis for fruit length was reported by Kasem and Somsak (1991) in cucumber, Mishra et al. (1994) in snake guard, and Pandey et al. (2002) in pumpkin.

Heterosis for fruit girth was exhibited by eight families and four families expressed heterosis in the negative direction. Negative heterosis for fruit girth was also noticed by Pyzhenkov and Kosareva (1979). Imam et al. (1977) and Kasem and Somsak (1991) also reported positive heterosis for fruit girth in cucumber.

Only two hybrids were found to have positive heterosis and all other hybrids had negative heterosis for fruit weight. Heterosis for fruit weight was also reported by Pyzhenkov and Kosareva (1979), Fang et al. (1994), Kasem and Somsak (1991) in cucumber and Pandey et al. (2002) in pumpkin.

Regarding the fruit number, four hybrid progenies expressed positive heterosis, other families were inferior to the check variety. Pyzhenkov and Kosareva (1979), Fang et al. (1994), Kasem and Somsak (1991) in cucumber and Pandey et al. (2002) in pumpkin also reported heterois for number of fruits.

High magnitude of heterosis for fruit yield was exhibited by the families 1, 2 and 3 in which the same female parents were used. Pyzhenkov and Kosareva (1979) in cucumber, Lavende and Patel (1990), Ram et al. (1997), Ranpise et al. (1992), Mishra et al. (1994), Ram et al. (1997), and Radhika, (1999) in snake gourd, Janakiram and Sirohi (2002), Sharma et al. (1995), Bhalala et al. (2002) and Dubey and Mourya (2002) in bottle gourd, Choudhary et al. (2002) in musk melon, Mandal and Sirohi (1993) in pumpkin, Sirohi (1993) and Pandey et al. (2002) in pumpkin, Ghai et al. (1998) in summersquash also reported heterosis for fruit yield. Superior hybrids obtained from this study can be multiplied vegetatively for the fixation of heterosis.

# **SUMMARY**

.

. .

#### 6. SUMMARY

The present investigation on "Development and evaluation of hybrids in ivy gourd (*Coccinia grandis* L.Viogt.)" was carried out at the College of Agriculture, Vellayani during 2007-2008. The objectives of the study were to develop hybrids in ivy gourd, to estimate genetic parameters, variability among and within families and to identify superior hybrids over the check variety "Sulabha".

Hybridization was carried out between four genetically diverse gynoecious and three morphologically distinct androecious lines. Based on the mean values of the twelve families and their progenies, variability among the families and progenies were studied. Genetic parameters and standard heterosis were also estimated in this study.

The analysis of variance conducted for 12  $F_1$  families showed significant differences among the progenies for different characters. The mean values recorded for eight characters showed wide variation among the families. The family 1(Royappanpatty 1 x Royappanatty local) revealed superiority over the other families for days to first flower, first harvest, length, girth, number of fruits and fruit yield per plant.

The magnitude of variation among the progenies of 12 families revealed the heterogeneous nature of the  $F_1$  families. The heterogeneous nature of the  $F_1$  population along with the advantage of clonal propagation facilitates fixation of heterosis in the  $F_1$ . Significant variations among the progenies for vitamin C content was expressed in eight families viz., family 3, 4, 5, 7, 9, 11 and 12. Seven families viz., family 2, 3, 4, 5, 6, 11 and 12 showed significant variations among progenies for days to first flowering. Variations among progenies for fruit length were expressed by the families 2, 3, 4, 6, 9, 10, and 11. The families 4, 5, 8 and 10 showed significant variations for average fruit girth among the progenies. The families 4, 5 and 7 showed variations among progenies for days to first fruit harvest. The families 2, 6 and 8 showed variations among the progenies for girth of fruits. Families 4, 6 and 11 recorded significant variations among the progenies for number of fruits. Families 4 and 6 showed significant variations among progenies for fruit yield per plant. Superior plants can be selected from the families showing significant variation for different characters.

Pooled analysis of the data showed significant differences among the progenies for all the characters. The minimum days to flowering was recorded in progeny 1 and 5 of family 1 (Royappanpatty 1 x Royappanatty local) followed by progeny 7 of the same family and progeny 1 of family 3. (Royappanpatty 1 x Karyavattam). The maximum time for flowering and fruit harvest was taken by progeny 5 of family 12 (Karuvalur x Karyavattam). The progeny 1 of family 1 was found to be earliest to harvest followed by progenies 2, 5 and 6 of the same family.

Significant variation for average fruit length was observed among the progenies. Progeny 1 of family 1 recorded the longest fruits. The shortest fruits were produced by progeny 4 of family 4 (NBPGR x Royappanpatty local) also recorded the lowest value for fruit girth. The highest fruit girth was observed in progeny 1 of family 4. The progeny 3 of family 3 produced fruits having the maximum weight followed by progeny 3 of family 1. The maximum number of fruits were produced by progeny 2 of family 6 (NBPGR x Karyavattam). Progeny 1 of family 4 recorded the highest vitamin C content and the lowest being in progeny 1 of family 5.

The highest fruit yield was recorded by progeny 3 of family 1 followed by progeny 1 of family 3 and progeny 3 of the same family. Most of the families were heterogeneous for different characters.

The analysis of variance revealed significant differences among the genotypes for all the characters studied viz., days to first flowering, days

to first fruit harvest, average length of fruits, average girth of fruits, average weight of fruits, number of fruits per plant, vitamin C content and fruit yield per plant.

The genetic parameters of different characters in 12 families of ivy gourd were estimated in this study. The differences between phenotypic and genotypic coefficient coefficients of variation for different characters were found to be low in this study indicating the major contribution of genetic component.

The highest heritability was estimated for days to first harvest followed by fruit yield per plant, fruit weight, fruit length, fruit girth, number of fruits per plant, days to first flower and vitamin c content. Heritability was generally high for all the characters in this study. High heritability and high genetic advance for important traits like fruit yield per plant, number of fruits per plant, and fruit weight indicated the effectiveness of selection.

The correlation analysis revealed significant positive correlation of number of fruits per plant with yield per plant, fruit length and fruit weight. Positive association of these characters indicated the possibility of simultaneous improvement by selection.

Number of fruits per plant as well as length, girth and weight of fruits exerted significant positive direct effect on fruit yield per plant. Therefore improvement of anyone of these characters would bring out simultaneous improvement of yield.

The families showed standard heterosis at varying levels for different characters. In this study mean values of the families were taken for comparison with the standard parent. Compared to the standard check variety "Sulabha" the hybrid population took more days to flower. However, among the hybrid progenies family 1 took the minimum days to flower and fruit harvest. The families 1 (Royappanpatty 1 x Royappanatty local) and 3 (Royappanpatty 1 x Karyavattam) manifested marginal positive heterosis for fruit length and families 1 (Royappanpatty 1 x Royappanatty local), 2 (Royappanpatty 1 x Vellayani local), 3 (Royappanpatty 1 x Karyavattam) 5 (NBPGR 1 x Vellayani local and) and 6 (NBPGR x Karyavattam) expressed heterosis for fruit girth. The positive heterosis for average fruit weight was observed in the families 1 and 3. The maximum positive heterosis for number of fruits per plant was observed in the family 1 followed by 2 and 3. Nine hybrids expressed positive heterosis for vitamin C content. The heterosis for fruit yield per plant was the maximum in family 1 followed by 3 and 2. The superior progenies identified for different characters from the hybrid population can be utilized for further improvement.

# **REFERENCES**

#### 7. REFERENCES

- Allard, R.W. 1960. Principles of Plant Breeding, (John Wiley and Sons, 485p.
- Anitha, C.A. 1998. Variability in ridge gourd (*Luffa acutangula* (Roxb.(L)). M.Sc. (Ag) thesis, Kerala Agricultural University, Thrissur, p.112.
- Abusaleha, A. and Dutta, O.P. 1990. Studies on variability, heritability and scope of improvement in cucumber. Haryana J. Hort. Sci., 19: 349-352.
- Ashok, P. 2000. Character association of seeds on plant morphology in snake gourd (*Trichosanthes anguina* L.). M.Sc.(Ag) thesis, Kerala Agricultural University, Thrissur, p.121.
- Bhalala, M.K., Nag, M., Kathiria, K.B., Patel, A.D and Acharya, R.R.
  2002. Studies on heterosis in bottle gourd, International Conference on Vegetables, Abstracts of Invited and Contributed Papers, p.97.
- Bhave, S.G., Mehta, J.L., Bendale, V.W., Mhatre P.P., and Pethe, U.B. 2003. Character association and path coefficient analysis of bitter gourd, Orissa J. Hort., 31: 44-46.
- Bindhu, S., Mahakal, K.G., Kale, P.B. and Sakahre, S.B. 2000. Genetic variability in pumpkin (*Cucurbita moschata Duch ex. Poir*). A. Pl. Physiol., 14: 66-68.
- \*Burton, G.W. 1952.Quantitative inheritance in grasses. Proc. 6<sup>th</sup> Int. Grassland Congress, 277-283.
- Chacko, E. 1992. Evaluation of desert type of muskmelon (*Cucumis melo*L) for southern region of Kerala. M.Sc (Ag) thesis, KeralaAgricultural University, Thrissur, p.109.

- Chen, X.H., Cao, P.S., Xu, Q., Dong G. and Meng, L.Y. 1994. Genetic correlation and co-efficient analysis of perthenocarpic yield components in cucumber. Advances in Horticulture, 249-251
- Choudhary, B.R., Dhaka, R.S. and Fageria, 2002. Heterosis for yield and yield attributes in musk melon (*Cucumis melo* L.) International Conference on Vegetables, Abstracts of Invited and Contributed Papers, p.123.
- Choudhary, M.L., Joshi, S. and Amar Singh. 1985. Genetic studies in cucumber (*Cucumis sativus* L.). Prog. Hort., 17: 236-240.
- Choudhary, S.M., Kale, P.N. and Desai, U.T. 1991. Variability studies and scope of improvement for fruit yield in bitter gourd. J. Maharashtra agric. Univ., 16: 15-17.
- Choudhury, D. and Sarma, K.C. 2002. Studies on variability, heritability, genetic advance and correlations in ridge gourd (*Luffa acutangula* Roxb.). Hort. J., 15: 53-58.
- Dabholkar, A.R. 1992. *Elements of Biometrical Genetics*, Concept Publishing Company, New York, p431.
- Damarany, A.M., Adoul Nasr, M.H. and Abdulla, M.M.A. 1995. Yield and yield components of some cucurbita spp. Cultivar and hybrids under Assiut conditions L Summer squash (*Cucurbita pepo L.*). Assiut J. Agric. Sci., 26: 51-58.
- Deepthy, R. 2000. Heterosis and combining ability in melon (*Cucumis melo L.* var. conomon) M.Sc. (Ag) thesis, Kerala Agricultural University, Thrissur, p.116.
- Dhaliwal, M.S., Dhiman, J.S and Lal, T. 1996. Character association and causation in musk melon. Indian J. agric. Res., 30: 80-84.

- Dhanabal, S. P., Koate, C. K., Ramanathan, M., Elango, K., and Suresh, B. 2004. The hypoglycemic activity of *Coccinia indica* Wight & Arn. and its influence on certain biochemical parameters. Indian J. Phamacol., 36: 244-250.
- Dubey, S.K and Maurya, I.B. 2002. Heterosis and combining ability in bottle gourd (*Lagenaria siceraria* (Molina) Standl.), International Conference on Vegetables, Abstracts of Invited and Contributed Papers, p.102.
- \*Fang, X.J., Gu, X.F. and Han, X. 1994. New cucumber cultivar. 'Zhongnong 8' for outdoor cultivation. Chinese Veg., 3:2.
- Frey, K. J. 1966. Plant Breeding. Iowa State University Press, Ames, USA
- George, T.E. 1981. Biometrical studies in ash gourd (*Benincasa hispida* (Thumb.) Cogn.). M.Sc (Ag) thesis, Kerala Agricultural University, Thrissur, p.78
- Ghai, T. R., Singh, J., Arora, S.K. and Singh, J. 1998. Heterosis studies for earliness and yield in summersquash (*Cucurbita pepo* L.) Punjab, Vegetable Grower, 33: 35-40.
- Ilyas and Muhammad, M.H. 1955. Acta Botanica India, 23: 143-144.
- Imam, M.K., Abobaker, M.A. and Yacoub, H.M. 1997. Inheritace of some characters in cucumber, some quantitative characters. Libyan J. Agric., 6: 115-125.
- Indiresh, B.T. 1982. Studies on genotypic and phenotypic variability in bitter gourd (*Momordica charantia* L). Thesis abstracts, 8: 52.
- Iswaraprasad, C.M. 2000. Combining ability and heterosis in bitter gourd (Momordica charantia L). M.Sc (Ag) thesis, Kerala Agricultural University, Thrissur, p.107.
- Jain, J.P.1982. Statistical Techniques in Quantitative Genetics, Tata Mc Graw Hill Co. New Delhi, p.281.

- Jaiswal, R.C., Kumar, S., Raghav, M. and Singh, D.K. 1990. Variation in quality traits of bitter gourd (Momordica charantia L) cultivars. Veg. Sci., 17:186-190.
- Janakiram, T. and Sirohi, P.S. 1992. Studies on heterosis for quantitative characters in bottlegourd. J. Maharashtra agric. Univ., 17: 204 206.
- Johnson, H.W., Robinson, H.F. and Comstock. 1955. Genotypic and phenotypic correlations in soybeans and their implications in selection. Agron. J., 47: 477 – 483.
- Joseph, S. 1999. Evaluation of diploids and polyploids of ivy gourd (*Coccinia grandis* (L) Voigt.). M.Sc (Ag) thesis, Kerala Agricultural University, Thrissur, p.98
- Kandaswamy, R. 2004. Morphological, biochemical and molecular characterization of melon (*Cucumis melo* L). Ph.D thesis, Kerala Agricultural University, Thrissur, p.103.
- Kanwar, M.S., Korla, B.N. and Sanjeev Kumar. 2003. Evaluation of cucumber genotypes for yield and quantitative traits. Himachal J. Agric. Res., 29: 43-47.
- Karuppaiah, P and Kavitha, R. 2002. Genetic variability, correlation, path and D<sup>2</sup> studies in ridge gourd (*Luffa acutangula* L.) International Conference on Vegetables, Abstracts of Invited and Contributed Papers, p.60.
- \*Kasem, P. and Somak, R. 1991. Hybrid performance of mini cucumber (*Cucumis sativus*). Kasetsart – Journal, 25: 54 – 57.
- Kennedy, R.R., Arumugam, R., Kandasamy, G and Suresh, S. 1995. Heterosis studies in bitter gourd (Memordica charantia L). Madras Agrci. J., 82: 121-123.

- Kumar, R., Singh, D.K. and Ram, H.H. 1999. Manifestation of heterosis in bottle gourd (*Lagenaria siceraria* (Mol.) Standl.). A. agric. Res., 20: 177-179.
- Kumaran, S.S., Natarajan, S., and Thamburaj, S. 1998. Genetic variability in pumpkin (*Cucubita moschata* Poir). S. Indian Hort., 46: 138-142.
- Lakshmi, L. M., Haribabu, K, and Reddy, G.L.K. 2002 Genetic variability in pumpkin (*Cucubita moschata* Duch ex. Poir). S. Indian Hort., 45: 1-2.
- Lal, T. and Singh, S. 1997. Genetic variability and selection indices in muskmelon. (Cucumis melo L). Veg.Sci., 24: 111-117
- Lawande, K.E and Patil, A.V. 1989. Correlation studies in bitter gourd. J. Maharashtra agric. Univ., 14: 77-79.
- Lawande, K.E. and Patil, A.V. 1990. Heterosis in bitter gourd. Haryana J. Hort. Sci., 19: 388 – 391.
- Lenka, D and Mishra, R. 1973. Path coefficient analysis of yield in rice varieties, Indian J. Agric. Sci., 43: 376-379.
- \*Li Jianwu, Li-J.W. and Zhu DeWei. 1995. Genetic analysis of major agronomic characters in *Cucumis sativus* L. International Symposium on cultivar improvement of horticultural Part I: Vegetable Crops. Acta Horticulturae, 402: 388 – 391.
- Lovely, B. and Radhadevi, D.S. 2007. Estimates of genetic parameters in ashgourd (Benincasa hispida Cogn.) 19<sup>th</sup> Kerala Science Congress, Extended Abstracts, 481-483.
- \*Lush, J.L. 1949. Animal Breeding Plans. Iowa State Univ, Press, p.473.
- \*Ma, D.H., Lu, S.Z., Shen, W.Y., Huo, Z.R., S.J and Zhang, Q.D. 1995. Phenotypic correlation and path analysis on some characters in cucumber. Acta Agriculturae Boreali Sinica, 10: 34-37

- Mandal, J and Sirohi, P.S.2002. Heterosis and gene effects in ash gourd (*Benincasa hispida* (Thunbi) Cogn.), International Conference on Vegetables, Abstracts of Invited and Contributed Papers, p.63.
- Mangal, J.L., Pandita, M.L. and Sidhu, A.A. 1981. Variability and correlation studies in pumpkin (*Cucurbita moschata* Poir.). Haryana J. hort.Sci., 8: 82-86
- Manju, P and Wilson, D. 2002. Variability and genetic divergence in bottle gourd, International Conference on Vegetables, Abstracts of Invited and Contributed Papers, p.83.
- Mariappan, S. and Pappiah, C.M. 1990. Genetic studies in cucumber (Cucumis sativus L.).S. Indian Hort., 38: 70-74.
- Mathew, A. 1999. Genetic variability in bottle gourd (*Lagenaria siceraria* (Mol.) Standl.) in relation to yield and yield attributes. M.Sc (Ag) thesis, Kerala Agricultural University, Thrissur, p.106.
- Mathew, A., Markose, B.L., Rajan, S. and Nirmaladevi, S. 2001. Genetic divergence in bottle gourd (*Lagenaria siceraria* (Mol.) Standl.).
  Veg. Sci., 28: 121-123
- Mathew, S.S. and Khader, A.K.M. 1999. Genetic studies in snake gourd (*Trichosanthes anguina* L.). J. trop. Agric., 37: 71-72.
- Menon, M.P. 1998. Cataloguing and identification of promising ash gourd ecotypes in relation to season and maturity. M.Sc (Hort.) thesis, Kerala Agricultural University, Thrissur. p.105.
- Mishra, H.N., Mishra, R.S., Mishra, S,N. and Parhi, G. 1994. Heterosis and comining ability in bitter gourd. Indian J. Agric., 64: 310-313
- Mohanty, B.K. 2000. Selection on variability and selection parameters in pumpkin (*Cucurbita moschata* Duch ex. Poir). S. Indian Hort., 48: 111-113.

- Mohanty, B.K. and Mishra, R.S. 1999. Variability and genetic parameters of yield and its components in pumpkin. Indian J. Hort., 56: 337-342.
- \*Paiva, Wode, De Paiva, Wo. 1997. Genetic evaluation and correlation study in cucumber. Pesquisa Agropecuaria Brasileii, 32: 719-723.
- Pandey, S. Jagadish Singh and Banerjee, M.K. 2002. Evaluation of F, hybrids of pumpkin, International Conference on Vegetables, Abstracts of Invited and Contributed Papers, p.63.
- Paranjape, S.P., and Rajput, J.C. 1995. Association of various characters in bitter gourd and direct and indirect effects on yield. J. Maharashtra agric., Univ., 20: 192-193.
- Parmar, A.M., and Tarsem Lal, 2005. Variability studies in melon (Cucumis melo L.) Research on crops, 6: 314-327.
- Prasad, V.S.R.K and Singh, D.P. 1994b. Genetic association and interrelationship between yield components in cucumber. J. Maharashtra agric. Univ., 19: 147-148
- Prasad, V.S.R.K. and Singh, D.P. 1990. Studies on morphological and agronomical components of pointed gourd (*Trichsanthes dioica* Roxb.). Indian J. Hort., 47: 337-340
- Prasad, V.S.R.K. and Singh, D.P. 1992. Estimates of heritability, genetic advance and association between yield and its components in cucumber (*Cucumber sativus* L). Indian J. Hort., 49: 62-69
- Prasad, V.S.R.K., Pitchaimuthu and Dutta, O.P. 2002. Variation and component association in watermelon (*Citrullus lanatus* L. Thunb) inbreds grown under semi aridlateritic alfisols, International Conference on Vegetables, Abstracts of Invited and Contributed Papers, p.90.

- Prasad, V.S.R.K., Pitchaimuthu and Dutta, O.P. 2002. Genetic variation and utilization of inbred lines possessing multiple disease resistance in muskmelon (Cucumis meloL.) improvement. International Conference on Vegetables, Abstracts of Invited and Contributed Papers, p.119.
- Prasunna, M.N and Rao, M.R. 1988. variability studies in cucumber (Cucumis spp) SouthIndian Hort., 36: 237-241
- Pynadath, J.S 1978. Genetic variability and correlation studies in sanke gourd (*Trichosanthes anguina* L). M.Sc. (Ag) thesis, Kerala Agricultural University, Thrissur, p.124
- \*Pyzhenkov, V. I. and Kosareva., G.A. 1979. Effect of heterosis on yield structure in cucumber. Trudy po Príkladnoi Botanika, genetika [ Selekstii, 65: 112-118
- Radhika, V.S. 1999. Estimation of comining ability and heterosis in snake gourd (*Trichosanthes anguina* L). M.Sc. (Ag) thesis, Kerala Agricultural University, Thrissur, p.117
- Rahman M.A., Hossain M.D., Islam M.S., Biswas, D.K., and M. Ahiduzzaman. 2002, Genetic variability, Heritability and Path Analysis in snake gourd, Pakistan. J. Biol., 5: 284-286
- Raja, S., Bagle, B.G., and Dhandar D.G. 2007. Genetic variability studies in bitter gourd for zero irrigated condition of semi-arid ecosystem, Indian J. Hort. Vol. 64, Issue 4.
- Rajput, J.C., Paranjape, S.P. and Jamadadagni, B.M. 1995. Correlation and path analysis studies in bitter gourd. J. Maharashtra agric. Univ., 20:377-379
- Rajput, J.C., Paranjape, S.P. and Jamadagni, B.M. 1996. Variability, heritability and scope of improvement for yield components in bitter gourd (*Momodica charantia* L.). Annals Agric. Res. 17: 90-93

- Rajput, P.C., Palve, S.B. and Jamadagni, B.M. 1991. Correlation and path analysis studies in cucumber (*Cucumis sativus* L). Maharashtra J. Hort., 5:52-55
- Raju, V.K. and Peter, K.V. 1995. Final Report of the ICAR Adhoc scheme on srvey, collection, evaluation, conservation and cataloguing of germplasm of certain under-exploited perennial vegetables. Kerala Agricultural University, Thrissur, p.18-60
- Rakhi,S and Rajamony,L.2005. Variability, heritability and genetic advance in land races of culinary melon ((Cucumis meloL.) J. Trop. Agric., 43: 79-82.
- Ram, D., Kalloo, G. and Singh, M. 1997. Heterosis in bitter gourd. (Momodica charantia L.). Veg. Sci., 24: 99-102
- Ranpise, S.A., Kale, P.N., Desale, G.Y. and desai, U.T. 1992. Heterosis in bitter gourd (*Memordica charantia* L.). S. Indian Hort., 40 6: 313-315
- Rao, C.R 1952. Advanced Statistical Methods in Biometric Research. John Wiley & Sons, New York, p-390
- Rao, E.S., Munshi, A.D., and Verma, V.K. 2004. Genetic correlation and interrelationship of yield and its components in cucumber, Indian J. Hort., 61:315-318
- Rao, N.P., Rao, V.P and Reddy, I.P. 2000. Character association and path co-efficient studies in ridge gourd (*Luffa acutangula* (Roxb.) L). Andra agric. J., 47:103-107.
- Rastogi, K.B. and Deep, A. 1990a. A note on inter-relationship between yield and important plant characters of cucumber (*Cucumis sativus* L). Veg.Sci., 17:102-104.

- Rastogi, K.B. and Deep, A. 1990 b. Variability studies in cucumber (*Cucumis sativus* L.). Veg. Sci., 17 : 224-226.
- Reddy, V.V.P., Rao, M.R. and Reddy, C.R. 1987. Heterosis and combining ability in water melon (*Citrullus lanatus* (Thunb.) Mansf.). Veg. Sci., 14: 152-160.
- Rehman, M.A., Hossain, M.D., Islam, M.S., Biswas, D.K. and Ahiduzzaman. 2002. Genetic variability, Heritability and path analysis in snake guard. Pakistan J. Biol. Sci., 5: 284-286.
- Robinson, H.F., Comstock, R.E. and Harvey, P.H. 1949. Estimation of heritability and the degree of dominance in corn, Agron. J., 14: 352-359.
- Sadasivam, S and Manickam, A. 1996. Biochemical Methods for Agricultural sciences. Wiley Eastern Ltd and TamilNadu Agricultural University, Coimbatore, p-388
- Saika, J., Shadeque, A. and Bora, G.C. 1995. Genetic studies in cucumber correlation and path co-efficient analysis. Haryana J. Hort. Sci., 24: 126-130
- Sarkar, S.K., Maity, T.K., Roy, K. and Som, M.G. 1990. Studies of genetic variability of popinted gourd (*Trichosanthes dioica* Roxb.). Exp. Genet., 6: 68-73
- Sarkar, S.K., Maity, T.K., Roy, K. and Som, M.G. 1999. Correlation and path co-efficient studies in pointed gourd (*Trichosanthes dioica* Roxb.). Indian J. Hort., 56: 252
- Sarnaik, D.A., Verma, S.K. and Sharma, G.L. 1999. Character association in ivy gourd (*Coccinia grandis*). A. agric.res., 20: 436-438
- Saroj Rolania, Fageria, M.S., Dhaka, R.S., and Iat, R.G 2003. Correlation and Path coefficient Analysis in water melon. Haryana J. hort. Sci., 32: 113-116

- Satyanarayana, N. 1991. Genetical studies in cucumber (*Cucumis sativus*L). Ph.D. (Ag) thesis, Kerala Agricultural University, Thrissur, p.168
- Seshadri, V. S., and More, T. A, 2004. History and antiquity of cucurbits in India. The 8<sup>th</sup> EUCARPIA- Meeting on Cucurbit Genetics and Breeding, Olomouc, Czech- Republic, 12-17 July' 2004: 81-90
- Sharma, N.K., Dhanka., B.S and Jowatia, A.S. 1995. Heterosis in bottle gourd. Haryana agric. Univ. J. Res., 23: 8-14
- Shibukumar, V.N. 1995. Variability studies in water melon (Citrullus lannatus Thunb. Mansf). M.Sc (Ag) thesis, Kerala Agricultural University, Thrissur, p.114
- Singh, A.K., Singh, R.D. and Singh, J.P. 1993. Correlation and path coefficient analysis in pointed gourd. Indian J. Hort., 50: 68-72
- Singh, A.K., Singh, R.D., and Singh, K. 1992. Genetic variability and genetic advance for some traits in pointed gourd (*Trichosanthes dioica* Roxb.). Haryana J. Hort. Sci., 21: 236-240
- Singh, R.P., Mohan, J and Singh, D. 2002. Studies on genetic variability and heritability in ridge gourd (*Luffa acutangula*). Agric. Sci. Digest, 22: 279-280
- Sirohi, P.S. 1993: Genetic diversity in cucurbits-pumpkin. Indian Hort., 38 : 35-37
- Sirohi, P.S. and Yayasani, S.R. 2001. Gene action of mineral elements and vitamins in pumpkin (*Cucurbita moschiata.*) Veg. Sci., 28: 127-129
- Suresh. S. 2004. Molecular characterization of ivy gourd (*Coccenia grandis* (L) voigt.) M.Sc. (Agri) thesis, KAU, Thrissur.
- Suribabu, B., Reddy, E.N. and Rao, M.R. 1986. Inheritance of certain quantitative and qualitative characters in bitter gourd (*Momordica* charantia L.). South Indian Hort., 34: 380-386

- Swamy, K.R.M. and Dutta, O.P. 1985. A diallel analysis of total soluble solids in musk melon (*Cucumis melo L.*). Madras Agric. J., 72: 399-403
- Thakur, j.C. and Nandpuri, K.S. 1974. Studies on variability and heritability of some important quantitative characters in water melon (*Citrullus lannatus* (Thumb.) Mansf. ). Veg. Sci., 1: 1- 8
- Tyagi, J.D. a972. variability and correlation studies in bottle gourd (*Lagenaria siceraria*). Indian J. Hort., 29: 219 222
- Vahab, M.A. 1989. Homeostatic analysis of components of genetic variance and inheritance of fruit colour, fruit shape and bitterness in bittergourd (*Memordica charantia* L.). Ph.D thesis, Kerala Agricultural University, Thrissur.
- Varalaksmi, B., Rao, P.V and Reddy, Y.N 1995. Genetic variability and heritability in ridge gourd (*Luffa acutangula*).Indian J. agric. Sci., 65: 608-610
- Varghese, P. 1991. Heterosis in snake gourd (*Trichosanthes anguina* L). M.Sc. (Ag) thesis, Kerala Agricultural University, Thrissur, p.106
- Varghese, P. and Rajan, S. 1993 b. Heterosis of growth characters and earliness in snakegourd (*Trichosanthes anguina* L.). J. Tropic Agric., 31: 18 - 23
- Varghese, P. and Rajan, S. 1993. Genetic variability and heritability studies in snake gourd (*Trichosanthes anguina* L). J. Trop.Agric., 31: 13-17
- Varghese, S. 2003. Genetic analysis of ivy gourd (*Coccinia grandis* (L) Voigt.). Ph.D thesis, Kerala Agricultural University, Thrissur, p.112
- Wasantwisut, E and Viriyapanich, C. 2003. Ivy gourd (*Coccinia grandis*L. Voigt.) in human nutrition and traditional applications, Plantsin- human health- and- nutrition- policy, 60-66

92

- Wehner, T.C. and Cramer, C. S. 1996. Ten cycles of recurrent selection for fruit yield, earliness and quality in three slicing cucumber populations. J. Amer. Soc. Hort. Sci., 121: 362 – 366
- Wilson D., Jiju J.K.K, Abdul Khader, K.M. and Sunny K Oommen. 2006
  Clonal selection and hybridization in ivy gourd (*Coccinia grandis*L. Voigt.). Proceedings of the 18<sup>th</sup> Kerala Science Congress,
  Extended Abstracts, 18-19
- Wilson D. Jiju J.K.K, Abdul Khader, K.M., Sunny K Oommen and Prabu, R. 2007. Performance of the F<sub>1</sub> progeny of ivy gourd (*Coccinia* grandis L. Voigt.). 19<sup>th</sup> Kerala Science Congress, Extended Abstracts, 505-507
- Wilson, D. and Deepthy, R. 2006. Heterosis for yield and fruit characters in melon (*Cucumis melo* var. conomon), Second National Plant Breeding Congress, Abstracts, p.168,169.
- Zalapa J.E., Staub J.E, Mc Creight J.D. Chng S.M., and Cuevas H, 2007. Detection of QTL for recombinant inbred lines derived from exotic and elite US Western Shipping melon germplasm, Theoritical and Applied Genetics, 114:7, 1185
- \* Zhang, M., Wang, X. and Cui, H.W. 1999. Genetic path analysis of early yield in cucumber, Report Cucurbit Genetic Co-operative, 22:3-4

# **DEVELOPMENT AND EVALUATION OF HYBRIDS IN IVY GOURD (Coccinia grandis L.Voigt.)**

.

R. PRABU

### Abstract of the thesis submitted in partial fulfilment of the requirement for the degree of

## Master of Science in Agriculture

Faculty of Agriculture Kerala Agricultural University, Thrissur

2008

Department of Plant Breeding and Genetics COLLEGE OF AGRICULTURE VELLAYANI, THIRUVANANTHAPURAM-695 522

#### ABSTRACT

The present investigation on "Development and evaluation of hybrids in ivy gourd (*Coccinia grandis* L.Viogt.) was carried out at the College of Agriculture, Vellayani during 2007-2008. The objectives of the study were to develop hybrids in ivy gourd, to estimate genetic parameters, variability among and within families and to identify superior hybrids over the check variety "Sulabha".

Hybridization was carried out between four selected gynoecious and three androecious lines. Based on the mean values of the twelve families and their progenies variability among the families and progenies were studied. Genetic parameters and standard heterosis were also estimated in this study.

The family 1(Royappanpatty 1 x Royappanatty local) revealed superiority over the other families for days to flower, first harvest, length, girth number of fruits and fruit yield per plant.

The magnitude of variation among the progenies of 12 families revealed the heterogeneous nature of the  $F_1$  families. Significant variation among the progenies for vitamin C content was expressed in eight families viz., family 3, 4, 5, 7, 9, 11 and 12. Seven families viz., family 2, 3, 4, 5, 6, 11 and 12 showed significant variations among progenies for days to first flowering. Variations among progenies for fruit length were expressed by the families 2, 3, 4, 6, 9, 10, and 11. The families 4, 5, 8 and 10 showed significant variations for average fruit girth among the progenies. The families 4, 5 and 7 showed variations among progenies for days to first fruit harvest. The families 2, 6 and 8 showed variations among the progenies for girth of fruits. Families 4, 6 and 11 recorded significant variations among the progenies for number of fruits. Families 4 and 6 showed significant variations among progenies for fruit yield per plant. Superior plants can be selected from the families showing significant variation for different characters.

Pooled analysis of the data showed significant differences among the progenies for all the characters. The minimum days to flowering was recorded in progeny 1 and 5 of family 1 (Royappanpatty 1 x Royappanatty local) followed by progeny 7 of the same family and progeny 1 of family 3. (Royappanpatty 1 x Karyavattam). The same trend was observed for days to first harvest of fruits. The progeny 1 of family 1 was found to be earliest to take harvest followed by three other progenies progeny 2, 5 and 6 of the same family.

Significant variation for average fruit length was observed among the progenies. Progeny 1 of family 1 recorded the longest fruits. The highest fruit girth was observed in progeny 1 of family 4. The progeny 3 of family 3 produced fruits having the maximum weight followed by progeny 3 of family 1. The maximum number of fruits was produced by progeny 4 of family 1. Progeny 1 of family 4 recorded the highest vitamin C content. The highest fruit yield was recorded by progeny 3 of family 1 followed by progeny 1 of family 3 and progeny 3 of the same family. Superior progenies identified for different characters from the hybrid population can be utilized for further improvement.

The results of the analysis of variance showed significant differences among the genotypes for all the characters studied.

The differences between phenotypic and genotypic coefficients of variation were found to be low in this study indicating the major contribution of genetic component.

The highest heritability was estimated for days to first harvest followed by fruit yield per plant, fruit weight, fruit length, fruit girth, number of fruits per plant, days to first flower and vitamin c content. High heritability and high genetic advance estimated for important traits like fruit yield per plant, number of fruits per plant, and fruit weight which indicated the effectiveness of selection. Higher genotypic correlation than the phenotypic correlation observed for most of the characters indicated the reduced influence of environment. The number of fruits per plant had significant positive correlation with yield per plant, fruit length and fruit weight. Average length of fruits, girth of fruits, average weight of fruits and number of fruits per plant exerted significant positive direct effect on fruit yield per plant. Therefore improvement of anyone of these characters would bring out simultaneous improvement of yield.

. 7

The families showed standard heterosis at varying levels for different characters. In this study mean values of the families were taken for comparison with the standard parent. Compared to the standard check "Sulabha" the hybrid population took more days to flower. However, among the hybrid progenies family 1 took the minimum days to flower and fruit harvest. The families 1 (Royappanpatty 1 x Royappanatty local) and 3 (Royappanpatty 1 x Karyavattam) manifested marginal positive heterosis for fruit length and families 1 (Royappanpatty 1 x Royappanatty local), 2 (Royappanpatty 1 x Vellayani local), 3 (Royappanpatty 1 x Karyavattam) 5 (NBPGR 1 x Vellayani local and) and 6 (NBPGR x Karyavattam) expressed heterosis for fruit girth. The positive heterosis for average fruit weight was observed in the families 1 and 3.

The maximum positive heterosis for number of fruits per plant was observed in the family 1 followed by 2 and 3. Nine hybrids expressed positive heterosis for vitamin C content. The heterosis for fruit yield per plant was the maximum in family 1 followed by 3 and 2. The superior progenies identified for different characters from the hybrid population can be utilized for further improvement.