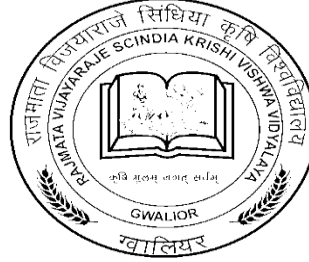


**Assessment of morphological, fruit yield and quality
characters in hybrid chilli (*Capsicum annum* L.)**

THESIS



Submitted to the

Rajmata Vijayaraje Scindia Krishi Vishwa Vidyalaya

In partial fulfillment of the requirements for the degree of

MASTER OF SCIENCE

In

HORTICULTURE

(VEGETABLE SCIENCE)

By

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2017

CERTIFICATE – I

This is to certify that the thesis entitled “**Assessment of morphological, fruit yield and quality characters in hybrid chilli (*Capsicum annum* L.)**” submitted in partial fulfillment of the requirement for the degree of **MASTER OF SCIENCE** in **Department of Vegetable Science** of Rajmata Vijayaraje Scindia Krishi Vishwa Vidyalaya, Gwalior, is a record of the bona-fide research work carried out by **Mr. Ravindra Patidar ID No 13559** under my guidance and supervision. The subject of the thesis has been approved by the Student’s Advisory Committee and the Director of Instruction.

No part of the thesis has been submitted for any other degree or diploma or has been published. All the assistance and help received during the course of investigation has been acknowledged by the scholar.

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CERTIFICATE – II

This is to certify that the thesis entitled “**Assessment of morphological, fruit yield and quality characters in hybrid chilli (*Capsicum annum* L.)**” submitted by **Mr. Ravindra Patidar ID No 13559** to the Rajmata Vijayaraje Scindia Krishi Vishwa Vidyalaya, Gwalior, in partial fulfillment of the requirements for the degree of Master of Science in **Agriculture** in the Department of **Vegetable Science** has been accepted after evaluation by the External Examiner and approved by the Student’s Advisory Committee after an oral examination of the same.

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ABBREVIATIONS USE IN TEXT

| | | |
|----------------|---|--|
| / | : | Per |
| @ | : | At the rate of |
| % | : | Percentage |
| ANOVA | : | Analysis of variance |
| B: C | : | Benefit cost ratio |
| C.D | : | Critical difference |
| cm | : | Centimeter |
| ^o C | : | Degree Celsius |
| cv. | : | Cultivar |
| D.F. | : | Degree of Freedom |
| DAP | : | Dia Ammonium Phosphate |
| DAS | : | Days after sowing |
| <i>et.al.</i> | : | and others |
| Fig. | : | Figure |
| g | : | Gram |
| ha | : | Hectare |
| HI | : | Harvest index |
| R.V.S.K.V.V | : | Rajmata Vijayaraje Scindia Krishi Vishwa Vidyalaya, Gwalior |
| K | : | Potassium |
| Kg/ha | : | Kilogram per hectare |
| Max. | : | Maximum |
| Min. | : | Minimum |
| mg | : | Milligram |
| MOP | : | Murate of potash |
| M.S.S | : | Mean sum of square |
| No. | : | Number |
| P | : | Phosphorus |
| R.H. | : | Relative humidity |
| Rs. | : | Rupees |
| S.Em. | : | Standard Error of Mean |
| Sq m | : | square meter |
| Sig. | : | Significant |
| q/ha | : | Quintal per hectare |
| Viz. | : | (Videlicet) Namely |

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Place:

Date:

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CHAPTER - I

INTRODUCTION

Chilli (*Capsicum annum* L., $2n=24$) is the most important warm season vegetable grown throughout the world. It belongs to the family Solanaceae and said to be the native of South America. Chilli was known to Indians about 400 years ago, when this crop was first introduced into India by Portuguese, towards the end of the 15th century. Its cultivation became popular in the 17th century. It is an important cash crop in India and is grown for its pungent fruits, which are used both green and ripe (the latter in the dried form) to impart pungency, spicy taste and colour to the food.

The composition of green chilli per 100 g of edible portion is moisture (85.7g), protein (2.9g), fat (0.6g), minerals(1.0g), fiber (6.8g), carbohydrates (3.0g), calcium (30mg), magnesium (24mg), riboflavin (0.39mg), nicotinic acid (0.9mg), phosphorous (80mg), iron (1.2mg), sodium (6.5mg), potassium (217mg), Vitamin A (292 I.U.) and Vitamin C (111mg). In fact, Vitamin C was first purified from *Capsicum* fruits in 1928 by Hungarian biochemist Albert Szent Gyorgyi which helped him to receive the Nobel Prize in physiology and medicine during 1937 (Kumar *et al.*, 2006). Green chillies are rich in Vitamin A and C, minerals and protein. Dry chillies are also rich in Vitamin A and D. As a condiment, it has become indispensable in every Indian home. It is also used medicinally, sauces, chutneys and pickles. The pungency is due to the oleoresin 'capsaicin' (a condensation product of 3 hydroxy-4-methoxy benzylamine and decylenic acid). It is secreted by the outer walls of the fruit. At ripening stage, fruit become red in colour due to presence of capsanthin pigment.

Chilli is extensively cultivated in Asia, Africa, Europe, and Central Northern part of America. In India, the major chillies growing states are Andhra Pradesh, Karnataka, Maharashtra, Orissa, Tamil Nadu, Madhya Pradesh, and Rajasthan. India occupies 0.794 million ha area and annual production 1.304 million tons,

while in Madhya Pradesh it occupies 0.054 million ha area and produce 0.093 million tons (NHB, 2013).

Chilli is one of the most important vegetable cum spice crops in India. In spite of its high nutritive value, well acceptability among consumers and wide range of genetic variability, the optimum productivity in chilli still remain to be achieved. Therefore, much concerted efforts are necessary to improve its yield and yield attributes. Fruit yield as well as quality improvement efforts continue to be the major objective of chilli improvement programme. Fruit yield is a complex inherited character influenced by several attributes of the plant. A wide range of variability is available in chilli genotypes which provide great scope for improving fruit yield through systematic breeding. Estimation of genetic variability present in the germplasm of a crop is a pre-requisite for designing effective breeding programme. However, utilization of this variability requires its systematic evaluation to understand and to estimate the genetic variability, heritability and genetic advance of various yield and physiochemical components (Amit *et al.* 2014).

The extent of genetic variation for major economic traits in a crop is pre-requisite for its improvement due to fact that the efficiency of selection depend mainly on it. Genetic improvement of traits depends upon nature and amount of genetic variability present in parents and their inheritance pattern are equally important for further upgrading yield level.

A comprehensive knowledge of the available variability within the breeding material of a crop species for desired characters enables the breeders to identify most potential genotype. The phenotypic variation arises as a result of phenotypic, genotypic and interaction between genotypic and environmental variations, but for making effective selections, the heritable unit *i.e.*, the genetic variation is most important. Burton (1952) suggested that genetic variability along with heritability should be considered for assessing the maximum and accurate effect of selection. Shift in the gene frequency towards desirable combinations under selection procedure is termed as genetic advance as percentage of mean .

Correlation and path analysis have been successfully used for architecting ideotypes for improvement to enhance productivity. However, under complex situation, correlations alone become insufficient to explain relationships among characters and thus path analysis of economic yield components with yield is most important the similar published information's is scanty in chilli (Krishna *et al.* 2007 and Hosamani and Shivkumar, 2008). Keeping in view these points the present investigation was carried out "Assessment of morphological, fruit yield & quality characters in hybrid chilli (*Capsicum annum* L.)" with the following objectives:

1. To estimate extent of genetic variation for growth and fruit yield traits.
2. To estimate heritability and genetic advance for growth and fruit yield traits.
3. To find out major fruit yield traits through correlation and path analyses.

CHAPTER – II

REVIEW OF LITERATURE

The literature pertinent to the various aspects of the present investigation entitled “**Assessment of morphological, fruit yield & quality characters in hybrid chilli (*Capsicum annum* L.)**” is reviewed under the following heads:

2.1 Genetic Variability

2.2 Heritability and Genetic Advance

2.3 Correlation and path analysis

2.1 Genetic Variability

Genetic variability is a measure of the tendency of individual genotypes in a population to vary from one another. Variability is the amount of variation seen in a particular population.

Munshi and Behera (2000) indicated existence of considerable amount of genetic variability for all the characters except fruit girth. The number of fruits per plant exhibited highest values of GCV and PCV.

Manju and Sreelathakumary (2002) recorded high coefficients of phenotypic (PCV) and genotypic (GCV) variation for several characters, the highest being for fruits per plant, followed by yield per plant, seeds per fruit and fruit weight. The lowest PCV and GCV were for days to first flowering. The GCV was near to PCV for most of the characters, indicating a highly significant effect of genotype on phenotypic expression with very little effect of environment.

Ananthi *et al.* (2006) indicated that the genetic material in the present investigation possessed high variability. High estimates of PCV and GCV were obtained for green fruit yield per plant, vulnerability index, average fruit weight and number of fruits per plant indicated a good deal of variability for these characters.

Bharadwaj *et al.* (2007) planted twenty seven genotypes of diverse origin of chilli [*Capsicum annum* (L.)] to work out variability for fruit yield and nine other related characters. High GCV observed for number of fruits per plant.

Farhad *et al.* (2008) observed fourteen quantitative characters viz. days to 50% flowering, time between fruit set and maturity, plant height, primary branches per plant, number of secondary branches per plant, fruit length, fruit girth, fruit weight, number of fruits per plant, seeds per fruit, 100 seed weight, fruit pedicel length, vitamin-C content and dry fruit yield per plant. The analysis of variance revealed remarkable variation among the genotypes for the characters concerned. The magnitudes of phenotypic coefficient of variations were higher than corresponding genotypic coefficients of variations. Higher genotypic as well as phenotypic coefficient of variations were computed for vitamin-C content, number of secondary branches per plant, number of primary branches per plant and yield per plant.

Tembhurne *et al.* (2008) reported the differences between phenotypic and genotypic coefficient of variation were low for days to 50% flowering, plant height, fruit diameter and number of fruits per plant.

Gupta *et al.* (2009) recorded maximum genotypic & phenotypic coefficient of variations for fruits per plant, fruit yield, fruit length and low values for plant height, primary branches per plant, number of secondary branches per plant.

Sarkar *et al.* (2009) evaluated forty-nine genotypes of chilli to study the genetic variability for 12 growth and fruit characters. There was significant variation among the genotypes. Fruit yield (g) per plant, number of fruits per plant, fruit length (cm), placenta length (cm), fruit weight (g), number of seeds per fruit and plant height (cm) showed high values of GCV and PCV.

Sharma *et al.* (2009) reported that the phenotypic coefficient of variation (PCV) and genotypic coefficient of variation (GCV) were high for fruit yield per plant and ascorbic acid content indicating that these traits had wide genetic variability and would respond better to selection.

Chattopadhyay *et al.* (2011) recorded phenotypic and genotypic coefficient of variation values for green fruit weight (119.95%, 111.26%), green fruit girth (89.76%, 48.93%), weight of red ripe fruit (112.02%, 111.93%), weight of dry fruit (111.63%, 110.97%) and number of fruits per plant (86.05%, 85.02%).

Ullah *et al.* (2011) studied genetic variability for morphological traits in chilli genotypes. Statistically, significant variation was observed among tested materials for all the characters studied. The higher genotypic coefficient of variations was found in case of fruit yield per plant followed by fruits per plant, average fruit weight and fruit length.

Singh *et al.* (2013) reported that the average of mean data reflects maximum variability for yield per plant itself followed by number of fruits per plant, fruit length, fruit width, plant height, days to 50% flowering and days to 50% fruiting. On the basis of mean performance and range of varietal mean of yield and their components, these genetic materials have enough variability.

2.2 Heritability and Genetic Advance

Heritability is an index of transmissibility of traits from one generation to another generation. The concept of heritability is important to determine whether the phenotypic differences observed among various individuals are due to differences in their genetic make-up or simply as result of environmental factors.

According to Comstock and Robinson (1952) the genetic advance is an improvement in the genetic value in the new population as compared to original one and depends upon the following three factors namely heritability, phenotypic and the intensity of selection.

Manju and Sreelathakumary (2002) estimated heritability for most of the characters ranged from 39.35 (primary branches per plant) to 99.38 per cent (yield per plant). Higher magnitude of heritability (>90 per cent) resulting from high GCV was also registered for fruits per plant, fruit length, fruit girth, fruit weight, seeds per fruit, 1000-seed weight, pollen viability, days to maturity and number of harvests. The study revealed high heritability coupled with high genetic advance for several biometric characters including yield per plant, fruit per plant, fruit weight, fruit girth and fruit length.

Shreelathakumary and Rajamony (2004) noticed high estimates for heritability with high genetic advance for branches per plant, fruit yield, seeds per plant, plant weight while low for fruit dry weight.

Abu and Uguru (2006) recorded high heritability for plant height, length, fruit diameter, fruit weight and number of fruit per plant for 10 genotypes of chilli.

Bharadwaj *et al.* (2007) planted twenty seven genotypes of diverse origin of chilli [*Capsicum annuum* (L.)] to work out heritability for fruit yield and nine other related characters. High heritability along with genetic advance observed for number of fruits per plant indicated that number of fruits per plant may be governed by additive genes.

Shirshat *et al.* (2007) estimated genetic variability in seventy-two germplasm lines and three commercial cultivars. The analysis of variance and other genetic parameters indicated considerable genetic variability for different characters among the genotypes. The phenotypic coefficient of variation was higher than genotypic coefficient of variation for all characters indicating the influence of environment on these characters. Fruit attributes viz., fruit length, fruit surface area, weight of dry fruit, pericarp weight of fruit, number of seeds per fruit, weight of seeds per fruit, stalk length, ascorbic acid and sugar content showed very narrow differences between phenotypic and genotypic coefficient of variation, indicating lesser sensitivity to environmental influence.

Shirshat *et al.* (2007) estimated heritability in respect of fruit length, fruit surface area, number of seeds per fruit, weight of seeds per fruit, weight of dry fruit, pericarp weight of fruit, ascorbic acid content and sugar content were high ranging from 74.00 per cent to 99.40 per cent. Moderate genetic advance was observed for the characters like number of fruits per plant, number of seeds per fruit and sugar content of the fruit. Heritability was high in these characters except for number of fruits per plant. In case of attributes like fruit length, fruit surface area, weight of dry fruit, pericarp weight of fruit, number of seeds per fruit and weight of seeds per fruit, the genetic advance was low to moderate coupled with high heritability. Yield per plant, the complex trait, which is dependent on several component characters showed moderate heritability with low genetic advance.

Vani *et al.* (2007) reported that high heritability coupled with high genetic advance was noticed through number of fruits per plant, number of seeds per

fruit, seed yield, and fruit length. The phenotypic and genotypic coefficients of variation were highest for dry fruit yield per plant, number of fruits per plant and seed yield per plant.

Ukkund *et al.* (2007) estimated high heritability for plant height (93.40%), days to first flowering (83.50%), percent fruit set (70.70%), number of fruits per plant (81.10%), fruit length (92.40%), ten fruit weight (92.40%) and total green fruits per plant (88.40%). Most of these characters also had moderate to high estimates of genetic advances as a percent over mean except days to first flowering.

Farhad *et al.* (2008) observed fourteen quantitative characters viz. days to 50% flowering, time between fruit set and maturity, plant height, primary branches per plant, number of secondary branches per plant, fruit length, fruit girth, fruit weight, number of fruits per plant, seeds per fruit, 100 seed weight, fruit pedicel length, vitamin-C content and dry fruit yield per plant. Among them, heritability estimates were higher for all the characters except days to 50% flowering and pedicel length. High heritability coupled with high genetic advances was estimated for seeds per fruits and plant height.

Singh and Yadav (2008) investigated high to moderate heritability with high to moderate genetic advance for number of seeds per fruit, ascorbic acid content, number of fruits per plant, fresh fruit weight per plant, fresh fruit yield (qperha), seed weight per plant and seed yield (qperha) showing dominance of additive gene effects.

Tembhurne *et al.* (2008) observed high heritability and high genetic advance values for fruits per plant.

Pandit *et al.* (2009) recorded the high heritability and genetic advance for number of seeds per fruit, fruits yield per plant, fruit length, days to 50% flowering and plant height.

Sarkar *et al.* (2009) reported high heritability in broad sense coupled with high Genetic Advance in % grand mean was recorded for fruit yield per plant, number of fruits per plant, fruit length, days to 50% flowering and plant height indicating such characters were controlled by additive gene action

Singh *et al.* (2009) noted high heritability coupled with high genetic advance for marketable fresh and dry yield per plant, average fruit weight, numbers of marketable fruit, fruit diameter, and oleoresin and capsaicin content, which indicated the role of additive gene action for the inheritance of these traits. These traits are likely to respond better to selection

Wilson and Philip (2009) reported the maximum heritability with maximum genetic advance for number of fruits and fruits yield per plant.

Sharma *et al.* (2010) recorded high heritability and high genetic advance for average fruit weight, fruit yield per plant, fruit diameter, number of lobes per fruit, days to first harvest, leaf area and ascorbic acid content. They indicated the role of additive gene action for the inheritance of these traits.

Chattopadhyay *et al.* (2011) reported that Green fruit yield per plant, ascorbic acid content, and number of fruits per plant showed very high broad-sense heritability and genetic advance.

Ullah *et al.* (2011) observed high heritability for all the tested characters except fruit diameter. High heritability with high genetic advance in percentage of mean was recorded for the characters fruit yield per plant, fruits per plant, plant height and days to 50% flowering indicating role of additive gene action in the expression of these traits.

Amit *et al.* (2014) reported that all the characters showed high heritability estimates. Number of the fruits per plant, green fruit yield per plant, dry (red) yield per plant, number of seeds per plant and plant height exhibited high genetic advance as percentage of mean indicating additive gene effect.

Anandhi and Anand (2014) recorded High heritability (above 80%) assisted with high genetic advance as percent of mean (above 20%) was observed for green fruit yield per plant, duration of crop, number of seeds per fruit, days to first flowering, duration of flowering, pedicel fruit ratio, number of fruits per plant and average fruit weight.

Pandit *et al.* (2014) recorded high genetic advance(as % of mean) in fruit yield per plant and moderately high in days to 50% flowering, placenta length,

number of fruits per plant and number of seeds per plant, indicating that these characters are most likely governed by additive gene action.

2.3 Correlation and Path Analysis

Correlation study gives the relationship of one character with the other. Yield, itself is not a unitary character but the end product of various component characters either jointly or singly. Therefore, knowledge of genetic association of yield and various component characters are of economic worth in formulating and executing the breeding programme. Hence, the prime requirement is to have precise and clear-cut information on the strength and direction of association of these traits with any worth character and also *inter se* relation among themselves. The selection efficiency is improved by making judicious combinations of the characters. The degree of associations between any character and its contributors can be estimated by correlation coefficient at genotypic and phenotypic levels. Correlation coefficient measure the association between any two characters. The literature available on correlation studies in chilli is reviewed as follows:

Ben-Chaim *et al.* (2003) founded the highest genetic correlation coefficients among pairs of traits between fruit weight and each of the three width characters: fruit diameter, pericarp thickness and pedicel diameter. In contrast, fruit weight had a low correlation coefficient with fruit length, indicating that the size of the pepper fruit in this cross was determined primarily by its width.

Mishra *et al.* (2003) reported the number of fruits per plant showed maximum positive direct effect on fruit yield per plant, followed by fruit weight and fruit diameter at phenotypic level and at genotypic levels

Ajjapplavara *et al.* (2005) studied the correlation and path coefficient analysis in 36 genotypes in chilli for 18 different quantitative characters. The correlation study indicated that significant and desirable correlation between dry fruit yield per plant with all other characters except number of primary and secondary branches, fruit diameter, fruit volume, powdery mildew disease incidence and leaf curl complex incidence. Path analysis revealed that importance should be given to fruit weight and fruits per plant.

Smitha and Basavaraja (2006) studied correlation and revealed that importance should be given to number of fruits per plant, fruit weight, number of primary branches, fruit length, fruit diameter and plant height during selection process, because these characters are going to contribute directly towards the yield.

Vani *et al.* (2007) observed the high positive direct effect of yield attributing characters such as fruit length and fruit weight resulted in significant correlation with yield. Number of fruits per plant and average fruit weight also contributed indirectly through all characters, which made the correlation significant.

Bharadwaj *et al.* (2007) planted twenty seven genotypes of diverse origin of chilli [*Capsicum annuum* (L.)] to work correlation association for fruit yield and nine other related characters. Fruit yield was positively associated with no. of branches per plant and no. of fruits per plant. Number of branches per plant is also positively correlated with fruit width, no. of fruits per plant, capsaicin content and fruits yield. They reported that selection for improvement of fruits yield is possible through positively associated yield contributing characters. However, these characters are influenced by environment. Therefore, selection of genotypes for improvement of fruit yield through no. of branches per plant and no. of fruits per plant is appropriate in the present set of material since these characters are not much influenced by environment.

Farhad *et al.* (2008) reported that dry fruit yield per plant showed significant and positive correlation with number of secondary branches per plant, fruit girth, fruit weight, fruits per plant, seeds per fruit and 100-seed weight. Fruits per plant and fruit weight exerted highest positive direct effect both at genotypic and phenotypic levels.

Tembhurne *et al.* (2008) reported that fruit yield exhibited positive association with most of the growth traits and high between fruit per plant with fruit diameter

Gupta *et al.* (2009) reported that the fruit yield per plant showed positively associated with fruits per plant and fruit length in chilli.

Patel *et al.* (2009) reported that the green fruit yield per plant showed highly significant and positive correlation with primary branches per plant, secondary branches per plant, number of fruits per plant, fruit length, fruit girth and fruit weight. Path coefficient analysis showed highly positive direct effect of number of fruits per plant, fruit weight, fruit length and number of primary branches per plant on green fruit yield per plant.

Sarkar *et al.* (2009) revealed that number of fruits per plant, fruit weight and 1000 seed weight had positive and high direct effect on fruit yield indicating their reliability as selection criteria to improve yield of chilli

Misra *et al.* (2010) performed genetic associations and path-coefficient analysis in 38 elite genetic stocks per accessions of chilli (*Capsicum annuum*). The capsaicin content was significantly associated with number of primary branches per plant and days to first flowering followed by weak positive association with days to fruit initiation and negative association with fruit weight dry and fruit weight fresh. In path-coefficient analysis, the number of primary branches per plant was found to be the premium direct contributor to the capsaicin content followed by fruit length. Fruit diameter exhibited the third highest direct effects. Primary branches per plant and days to first flowering emerged as a reliable component in selection criterion in breeding high capsaicin content yielding chilli (*Capsicum annuum*).

Sharma *et al.* (2010) revealed that at genotypic levels, the traits fruit length, fruit diameter and number of fruits per plant had significant positive correlation with fruit yield per plant. Number of fruits per plant exhibited the highest positive direct effect followed by average fruit weight, number of branches per plant, pedicel length and harvest duration at genotypic level.

Chattopadhyay *et al.* (2011) studied correlation and path coefficient analyses, the number of fruits per plant, green fruit length for green chilli, weight of dry fruit and the number of fruits per plant for dry chilli were found to be the most important selection indices.

Ullah *et al.* (2011) studied association of correlation and cause effect analysis for morphological traits in chilli genotypes. Fruits per plant, fruit length

and fruit diameter were the major characters contributing to yield as these traits were significantly and positively associated with yield per plant. Maximum contribution of fruits per plant to yield was observed in path analysis, which was followed by average fruit weight, days to first flowering and fruit length through higher direct effect. They revealed that the increasing fruit yield per plant a chilli genotype should have higher number of fruits per plant, coupled with large fruit length, high fruit diameter and high average fruit weight.

Amit *et al.* (2014) recorded that fruit yield (green and red) plant per plant was positively and significantly correlated with number of fruits plant per plant and fruit length. It revealed that the characters viz., plant height, fruit length, number of fruits plant per plant, fruit weight and fruit yield (green & red) are the most important traits for genetic improvement of chilli.

Vikram *et al.* (2014) studied the genetic correlations and path coefficients among twenty traits under study. Green yield per plant showed positive and significant correlation with average green fruit weight, fruit length and fruit breadth at middle while dry fruit yield exhibited the same with fruit length, green yield per plant and β -carotene content indicating the effective improvement in yield (green and dry) through above characters. Path analysis towards dry yield per plant revealed the importance of average dry fruit weight, number of fruits per plant, fruit length, green yield per plant and β -carotene content in improvement of dry yield per plant.

CHAPTER-III

MATERIAL AND METHODS

This chapter comprises the details about the materials used and the methods adopted during the course of present investigation entitled “Assessment of Morphological, Fruit Yield & Quality Characters in Hybrid Chilli (*Capsicum annum L.*)” was carried out in *Kharif* season 2014-15.

3.1 Experimental site

The present experiment was laid out at the field of Horticulture Research Farm, Department of Horticulture, R.A.K. College of Agriculture, Rajmata Vijayaraje Scindia Krishi Vishwa Vidyalyaya, Sehore, (M.P.).

3.2 Location and Climate:

The experimental site, R.A.K. College of Agriculture, Sehore is situated on 27⁰12 north latitude and 77⁰0 east longitude at an altitude of 498.77 meters from mean sea level in Vindhyan Plateau of Madhya Pradesh and enjoy sub tropical climate. The average rainfall varies from 689.3 mm concentrated mostly from the month of last July to January (Crop period) less rainfall occurs during the winter season also. The maximum temperature is 46⁰C and minimum temperature 6.8⁰C. The average annual relative humidity is 74%.

The meteorological parameter during the crop season such as minimum and maximum temperature, rainfall and relative humidity were recorded at Meteorological observatory R.A.K. College of Agriculture, Sehore (M.P.) are presented in Table 3.1

The data (Table 3.1) indicate that the total rainfall received during crop growth period was 783.4 mm in 43 rainy days. There were no rains during standard weather week 40 to 49. The minimum and maximum temperature during crop growth period varied 11.37⁰C to 25.28⁰C and from 20.94⁰C to 32.70⁰C, with season's average values of 18.88⁰C and 28.76⁰ C respectively. The relative humidity ranged between 68.71 to 98.71 % with season's average of 87.39%.

Table 3.1: Meteorological data (week wise) during the crop season of 2014-15 at R.A.K. College of Agriculture, Sehore (M.P.).

| SMW | Month and date | Temp (⁰ C) | | Rainfall (mm) | RH (%) |
|----------------|--------------------------|------------------------|-------|---------------|--------|
| | | Max. | Min. | | |
| 28 | 09.07.2014- 15.07.2014 | 28.8 | 23.8 | 14.0 | 65.9 |
| 29 | 16.07.2014- 22.07.2014 | 23.8 | 23 | 77.5 | 64.5 |
| 30 | 23.07.2014- 29.07.2014 | 29.3 | 22.9 | 7.0 | 65.6 |
| 31 | 30.07.2014- 05.08.2014 | 32.3 | 23.6 | 30.0 | 67.6 |
| 32 | 06.08.2014- 12.08.2014 | 30.7 | 23 | 143.0 | 66.3 |
| 33 | 13.08.2014- 19.08.2014 | 28.2 | 23.2 | 79.5 | 65.4 |
| 34 | 20.08. 2014- 26.08. 2014 | 28.1 | 22.5 | 5.5 | 64.8 |
| 35 | 27.08. 2014- 02.09. 2014 | 30.9 | 21.3 | 0.5 | 65.7 |
| 36 | 03.09. 2014- 09.09. 2014 | 32.1 | 20.8 | 1.0 | 66.1 |
| 37 | 10.09. 2014- 16.09. 2014 | 34.1 | 20.3 | 0.0 | 66.7 |
| 38 | 17.09. 2014- 23.09. 2014 | 32.3 | 19.1 | 0.0 | 68.5 |
| 39 | 24.09. 2014- 30.09. 2014 | 31.8 | 19.8 | 1.5 | 65.5 |
| 40 | 01.10. 2014- 07.10. 2014 | 30.8 | 17.3 | 0.0 | 63.7 |
| 41 | 08.10. 2014- 14.10. 2014 | 31.1 | 15.8 | 0.0 | 63.0 |
| 42 | 15.10. 2014- 21.10. 2014 | 30.4 | 16.8 | 0.0 | 63.1 |
| 43 | 22.10. 2014- 28.10. 2014 | 28.4 | 18.2 | 0.0 | 62.0 |
| 44 | 29.10. 2014- 04.11. 2014 | 28.3 | 11.2 | 0.0 | 58.4 |
| 45 | 05.11. 2014- 11.11. 2014 | 28.7 | 13 | 0.0 | 54.7 |
| 46 | 12.11. 2014- 18.11. 2014 | 26.3 | 9.5 | 0.0 | 57.4 |
| 47 | 19.11. 2014- 25.11. 2014 | 24.7 | 9.8 | 26.0 | 56.9 |
| 48 | 26.11. 2014- 02.12. 2014 | 21 | 6.2 | 0.0 | 52.2 |
| 49 | 03.12.2014- 09.12. 2014 | 21.7 | 5.4 | 10.5 | 53.7 |
| 50 | 10.12.2014- 16.12. 2014 | 17.6 | 7.6 | 25.0 | 59.2 |
| 51 | 17.12.2014- 23.12. 2014 | 28.8 | 23.8 | 14.0 | 65.9 |
| 52 | 24.12.2014- 31.12. 2014 | 23.8 | 23 | 77.5 | 64.5 |
| 1 | 01.01.2015- 07.01. 2015 | 29.3 | 22.9 | 7.0 | 65.6 |
| 2 | 08.01.2015- 14.01. 2015 | 21.77 | 10.94 | 0 | 55.93 |
| 3 | 15.01.2015- 21.01. 2015 | 21.44 | 6.20 | 0 | 61.14 |
| 4 | 22.01.2015- 28.01. 2015 | 17.14 | 7.58 | 1.0 | 61.74 |
| Total | | --- | --- | 422.0 | --- |
| Average | | 27.36 | 16.84 | 17.94 | 62.47 |

Source: Meteorological Observatory, R.A.K. College of Agriculture

3.3 Soil type

The soil of the experimental field was medium black with good drainage and uniform texture with low, medium and high NPK status, respectively.

3.3.1 Soil

The soil of the experimental field was medium black with 37% clay, 38% silt and 25% sand with pH ranging from 7.2. The soil was low in available nitrogen, medium in available phosphorus and high in available potassium.

Table 3.2: Chemical properties of experimental field

| S. No. | Composition | Content | Category | Method used |
|----------------------------------|---|---------------|--------------|---|
| A. Mechanical composition | | | | |
| 1 | Sand (%) | 25 | - | Bouyoucos Hydrometer method (Piper, 1967) |
| 2 | Silt (%) | 38 | - | |
| 3 | Clay (%) | 37 | - | |
| 4 | Textural class | | Medium black | |
| B. Chemical composition | | | | |
| S. No. | Particulars | Content | Level | Method adopted by |
| 1. | Organic carbon (%) | 0.47 | Low | Walkey & Black method (1934) |
| 2. | Available nitrogen N (kg per ha) | 152.6 | Low | Alkaline potassium permanganate method (Subbaih and Asija, 1956) |
| 3. | Available phosphorus P (kg per ha) | 21.2 | Medium | Olsen's method (Olsen <i>et al.</i> , 1954) |
| 4. | Available potassium K (kg per ha) | 308.0 | High | Flame photometer (Jackson, 1967) |
| 5. | Soil pH | 7.2 | Normal | pH meter (Jackson, 1967) |
| 6. | Electrical conductivity (mmhospercm) | 0.5 dsperm | Normal | Conductivity meter at 25°C (Jackson, 1967) |

In order to determine the textural class and fertility status of the experimental area, the soil samples were collected randomly from each plot with the help of soil auger before sowing from the experimental field. Primary samples were mixed to prepare and composite soil sample from each replication was drawn to study physico-chemical properties of the experimental field. The data pertaining to various physico-chemical properties have been presented in Table 3.2.

3.4 Cropping history of the experimental field

The following crops, as given in Table 3.3, were grown during the past four years.

Table 3.3: Previous history of the experimental field

| Year | Kharif | Rabi | Summer |
|---------|--------|-------|--------|
| 2012-13 | Onion | Gram | Fallow |
| 2013-14 | Cowpea | Gram | Fallow |
| 2014-15 | Cowpea | | |

3.5 Experimental details

The experiment was laid out in the Randomized block design with three replications. Each replication was comprised of 18 treatments. The details are given below.

Treatments Detail:

| | | | | | |
|-----|------------|------|--------------|------|-----------------|
| T1 | Smriti | T 7 | ARCH 930 | T 13 | S-586 |
| T 2 | Aujasvi | T 8 | US 612 | T 14 | Amrita |
| T 3 | Shourya | T 9 | US 611Tara | T 15 | P. jyoti |
| T 4 | Albeli-333 | T 10 | Marshal 1857 | T 16 | Jwala garima-12 |
| T 5 | Ujjala | T 11 | Saniya (03) | T 17 | TS-0017 |
| T 6 | Tiger | T 12 | Jalsa | T 18 | Tara |

Table 3.4: Details of layout

| | | |
|------------------------------|---|--|
| Experimental site | : | Horticulture Research Farm, R.A.K.College of Agriculture, Sehore (M.P.) |
| Crop | : | Chilli (<i>Capsicum annum</i> L.) |
| Design | : | Randomized Block Design |
| Treatments | : | 18 |
| Replications | : | 03 |
| Total no. of plots | : | 54 |
| Distance between replication | : | 1 m |
| Distance between plot | : | 0.5 m |
| Gross plot size | : | 3.0 m x 2.5 m |
| Total area | : | 405 m ² |
| Row to row Distance | : | 60 cm |
| Plant to plant Distance | : | 50 cm |
| Season | : | Late Kharif, 2014-5 |
| Date of sowing | : | 25.08.2014 |

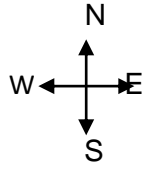
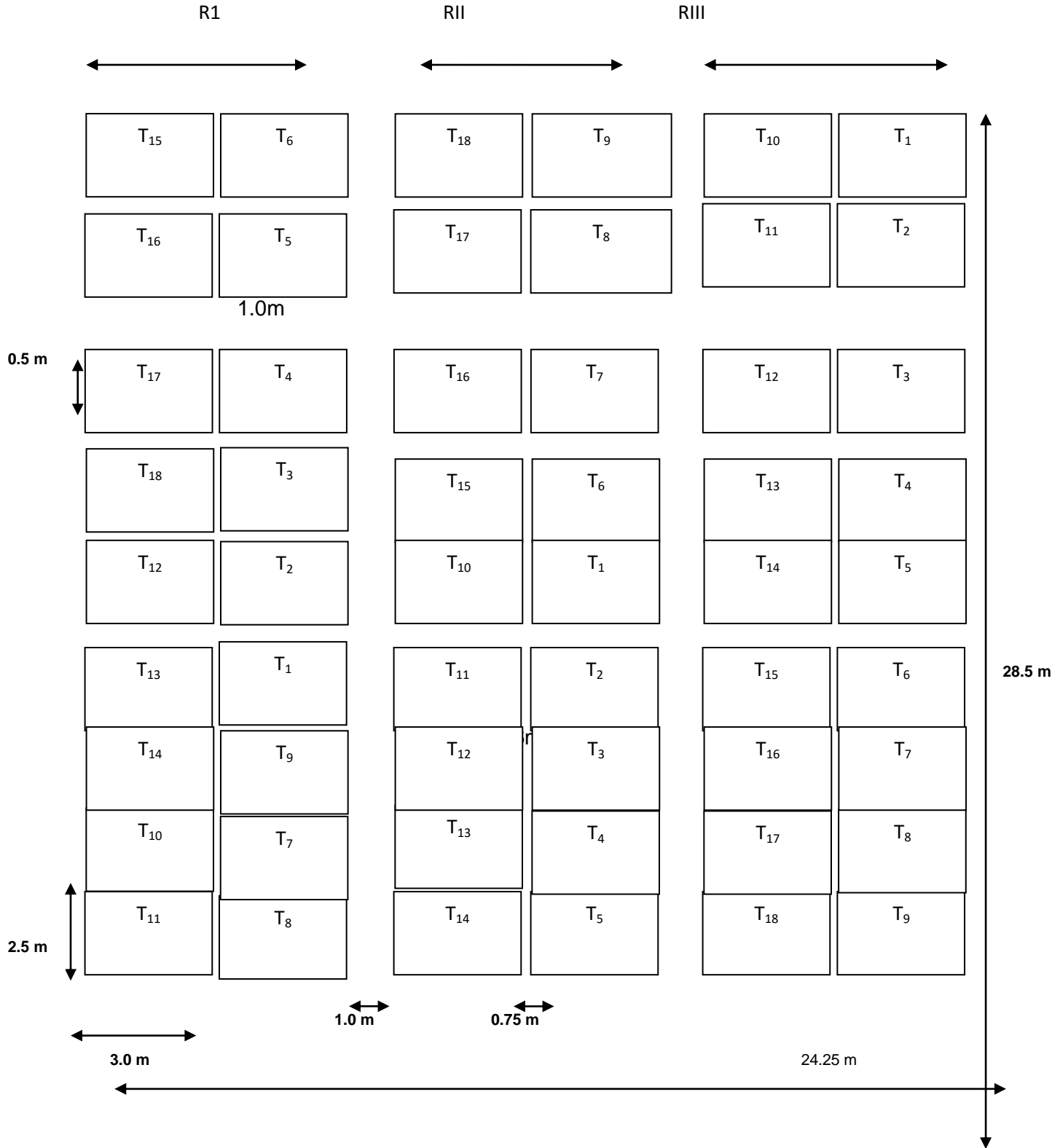


Fig.2 Layout Plan of Experiment

RANDOMIZED BLOCK DESIGN



3.6: Field preparation

The field was prepared with 2 ploughing and a harrowing so as to make the soil well pulverized and free from weeds. The field was laid out for the trial as per given in plan of layout.

3.6.1: Application of manure and chemical fertilizers

The treatments of manure and chemical fertilizers were applied as per respective plot. Full dose of phosphorus, potash, organic manures and 1/3 nitrogen were given to the plot before sowing as basal dose at the time of 2nd ploughing. Remaining 2/3 quantity of nitrogen was applied in two split doses i.e. 30 and 60 DAS.

3.6.2: Nursery:

The 200-250 g per ha seed was used for sowing and before sowing it was treated with 3 g Vitavax per kg culture of seed.

3.6.3: Irrigation

A light irrigation was applied for establishment of the seedlings after transplanting. Subsequent irrigations were provided as and when required for growth and development of plants. The irrigation at weekly interval was given as and when required subsequently after one month the irrigation interval was increased up to 10 days as per requirement.

3.6.4: Gap filling and thinning

Healthy and uniform seedlings were selected for planting. Light irrigation was given before uprooting seedlings from the nursery beds so that minimum damage may occur to roots of the seedling. Transplanting was done in the afternoon at 60 x 50 cm distance and immediately followed by light irrigation for proper establishment of the seedlings. A week after transplanting gap filling was done. All the other recommended package of practices was followed to raise a healthy crop.

3.6.5: Weeding

In order to control the weeds and avoid competition between plants and weeds, two hand weedings were also done by manual labour, at 20 and 45 days after sowing.

3.6.6: Plant protection

The crop was sprayed with Diafenthiuron 50% WP broad spectrum insecticide to keep the crop free from pest during crop growth period.

3.6.7: Harvesting

Harvesting of chilli was done at marketable stage at regular interval.

Table 3.5: Schedule of cultural operations carried out during the course of investigation

| S.No. | | Name of operations | Date |
|-------|---|---|------------|
| 1 | | Field operation | |
| | A | Ploughing by tractor | 20.08.2014 |
| | B | Two Harrowing | 22.08.2014 |
| | C | Planking | 22.08.2014 |
| 2 | | Field layout | 23.08.2014 |
| 3 | | Fertilizer application | 24.08.2014 |
| 4 | | Seed treatment | 20.07.2012 |
| 5 | | Seed sowing | 21.07.2014 |
| 6 | | Transplanting | 25.08.2014 |
| 7 | | Gap filling | 30.08.2014 |
| 8 | | Thinning | 30.08.2014 |
| 9 | | Dusting of Malathion dust | 30.08.2014 |
| 10 | | Spray of insecticide (1 st) | 20.09.2014 |
| 11 | | Spray of insecticide (2 nd) | 05.10.2014 |
| 12 | | Interculture and weeding (1 st) | 10.09.2014 |
| 13 | | Interculture and weeding (2 nd) | 25.09.2014 |
| 14 | | Top dressing of urea (1 st) | 25.09.2014 |
| 15 | | Top dressing of urea (2 nd) | 25.10.2014 |
| 16 | | Irrigation schedule | |
| | A | 1 st | 25.08.2014 |
| | B | 2 nd | 15.10.2014 |
| | C | 3 rd | 30.10.2014 |
| | D | 4 th | 12.11.2014 |
| | E | 5 th | 24.11.2014 |
| | F | 6 th | 08.12.2014 |

3.7: Observation and its procedure

Sampling was done at 30 day's interval for growth parameters. Five plants were randomly selected from each treatments and replication for the study.

Observation recorded

A. Growth / Morphological parameters

1. Plant height (cm) at an equal interval 30, 60, 90, 120 DAT and at maturity
2. Number of primary branches per plant at 30, 60, 90, 120 DAT and at maturity.
3. Number of secondary branches per plant at 30, 60, 90, 120 DAT and at maturity.
4. Days to first flower initiation.
5. Days of 50% flowering
6. Days to first picking of fruits.

B. Yield parameters

- | | |
|---|---|
| 1. Days to first fruit initiation | 2. Number of fruits per cluster |
| 3. Fruit width (cm) | 4. Fruit length (cm) |
| 5. Number of fruits per plant | 6. Days to maturity |
| 7. Fresh weight of fruits per plant (g) | 8. Fresh weight of fruits per plot (kg) |
| 9. Fresh weight of fruits per ha. (q.) | 10. Dry weight of fruits per plant (g) |
| 11. Dry fruit weight per plot (kg) | 12. Dry fruits weight per ha. (q.) |
| 13. Number of seeds per fruit | 14. 1000 seed weight (g) |

C. Phenological parameter

1. Leaf shape and size.
2. Shape and size of fruits.
3. Fruits colour
4. Fruit position per habit
5. Plant type

3.7.1 Growth parameters

3.7.1.1 Plant height (cm)

The height of five randomly selected plants was measured at 30, 60, 90, 120 DAT and at maturity from the base of plant to the tip of the plant with the help of meter scale. The average was worked out.

3.8.1.2 Number of primary branches per plant

Numbers of primary branches per plant were recorded under each treatment at 30, 60, 90, 120 and at maturity DAT from randomly selected five plants and the average was calculated.

3.7.1.3 Number of secondary branches per plant

Numbers of secondary branches per plant were recorded under each treatment at 30, 60, 90, 120 and at maturity DAT from randomly selected five plants and the average was calculated.

3.7.1.4 Days to first flower initiation.

Number of days taken from transplanting to the appearance of first flower on the plant in a plot was recorded. Data was recorded on plot basis.

3.7.1.5 Days of 50% flowering

Number of days required for 50% flowering after sowing were recorded under each treatment and replications.

3.7.1.6 Days to first picking of fruits.

The number of days taken from sowing to the date of first picking was calculated for each treatment and the mean values were determined.

3.7.2 Yield parameters

3.7.2.1 Days to first fruit initiation

The days taken from sowing to the date of first fruit initiation was calculated for each treatment and the mean values were determined.

3.7.2.2 Number of fruits per cluster

Number of fruits was counted on each cluster, 5 clusters were taken on each selected plants.

3.7.2.3 Fruit width (cm)

The fruit width of the selected five mature fruits was measured by the Vernier calipers in centimeter taken from each treatment and the average girth of the fruit was calculated.

3.7.2.4 Fruit length (cm)

The fruit length of five mature fruits of a plant from each plot was recorded and then average fruit length was tabulated.

3.7.2.5 Number of fruits per plant

Number of Mature green fruits that harvested from each picking in the tagged plants were counted and recorded till the final harvest. The

average number of fruits per plant was calculated and expressed as number of fruits per plant.

3.7.2.6 Days to maturity

Number of days from the date of transplanting to the date of physiological maturity of the plant in a plot.

3.7.2.7 Fresh weight of fruits per plant (g)

The fresh fruit chilli of selected plants were picked and weighed and fresh five fruits weight was tabulated.

3.7.2.8 Fresh weight of fruits per plot (kg)

The fresh fruit chilli of selected plants were picked and weighed and fresh five plants fruits weight was tabulated.

3.7.2.9 Fresh weight of fruits per ha. (q.)

The fresh fruit chilli of each plots were picked and weighed and calculated from 1 hac. Plant density and tabulated them.

3.7.2.10 Dry weight of fruits per plant (g)

The dry weight of fruit per plant of all the harvests per plant was taken and expressed in grams.

3.7.2.11 Dry fruits weight per plot (kg)

The dry weight of fruit per plot of all the harvests fruits randomly selected plants was taken and expressed in kilo grams.

3.7.2.12 Dry fruits weight per ha. (q.)

The dry yield of fruits per plant recorded was expressed in quintal per hectare.

3.7.2.13 Number of seeds per fruit

The fully matured fruits were harvested and counted their seeds from randomly selected fruits.

3.7.2.14 1000 seed weight (g)

The 1000 seeds were taken from each genotype of selected plants were weighed.

3.7.3 Phenological parameter

3.7.3.1 Leaf shape and size

Recorded at full foliage stage (see figure on page 49)

- Deltoid
- Ovate
- Lanceolate
- Others

Recorded at full foliage stage

- Small
- Medium
- Large
- Others

3.7.3.2 Shape and size of fruits

The shape of fruit was recorded in following categories.

- Straight
- Slightly curved
- Indefinitely curved

The size of fruit was recorded in following categories.

- Big
- Medium
- Small

3.7.3.3 Fruits colour

The colour of the fruits (fresh) was recorded at marketable stage in following categories.

- Dark green
- Green
- Light green
- Purple
- Yellowish green

3.7.3.4 Fruit position per habit

The fruit position per habit of the fruits (fresh) was recorded at harvesting stage in following categories.

3.7.3.5 Plant type

- Bushy
- Tall

3.8 Statistical analysis

The data obtained in respect of all the characters have been subjected to the following statistical analysis.

3.8.1 Mean: It was calculated by using following formula.

$$\text{Mean} = \frac{\sum x}{N}$$

Where,

$\sum x$ = The sum of all the observation

N = Number of observation

3.8.2 Analysis of variance

The data based on the mean of individual plants selected for observation were statistically analysed described by Panse and Sukhatme (1985) to find out overall total variability present in the material under study for each character and for all the populations. The first and foremost step is to carry out analysis of variance to test the significance of differences among the populations. The skeleton of analysis of variance used is given below.

Table 3.6: ANOVA for completely randomized block design

| Source of Variation | d.f. | Sum of Square | Mean sum of square | F value | F _t 5% or 1% table value |
|---------------------|-------------|---------------|--------------------|-----------|-------------------------------------|
| Replication | r-1 | RSS | RMS | RMSperEMS | - |
| Hybrids | g-1 | GSS | GMS | GMSperEMS | - |
| Error | (r-1)(g-1) | ESS | EMS | - | - |
| Total | rg-1 | TSS | - | - | - |

Where,

r = Number of replications

g = Number of hybrids

D.F. = Degree of freedom

RSS = Replication sum of square

GSS = Hybrid sum of square

ESS = Error sum of square

TSS = Total sum of square

RMS = Replication mean sum of square

GMS = Hybrid mean sum of square

EMS = Error mean sum of square

A significant value of F test indicates that the test entries differ significantly among themselves, which requires computing C.D.

$$C.V. = \frac{\sqrt{EMS}}{GM} \times 100$$

$$SE_{m \pm} = \sqrt{\frac{EMS}{r}}$$

$$SE_{diff} = \sqrt{\frac{2 EMS}{r}}$$

$$CD \text{ at } 5\% \text{ prob. Level} = SE_{diff} \times t_{5\%}^r \text{ table value}$$

Where,

C.V. % = Coefficient of variation

SE_{m±} = Standard error of means

S E diff = Standard error of difference

GM = Grand mean

C.D. = Critical difference

t 5% = t, table value 5% probability level at error d.f.

Estimation of mean, components of variance, phenotypic, genotypic and environmental coefficient of variation, heritability, genetic advance and genetic advance as percentage of mean:

The mean of different characters were calculated by conventional method:

$$\text{Mean} = \frac{\sum x_i}{n}$$

Where,

$\sum x_i$ = The sum of all the observation for i^{th} character.

N = Number of observations.

Range was recorded by observing the lowest and the highest mean values for each character.

Table 3.7: The component of variance was calculated as follows:

| S. No. | Source | M.S.S. | Expected M.S.S. |
|--------|-------------|--------|----------------------------------|
| 1. | Replication | - | - |
| 2. | Genotypes | M_i | $\sigma^2 e_i + r. \sigma^2 g_i$ |
| 3. | Error | E_i | $\sigma^2 e_i$ |

$$\sigma^2 g_i = \frac{M_i - E_i}{r}$$

$$\sigma^2 e_i = E_i$$

$$\sigma^2 p_i = \sigma^2 g_i + \sigma^2 e_i$$

Where,

$\sigma^2 g_i$ = Genotypic variance for i^{th} character.

$\sigma^2 e_i$ = Environmental variance for i^{th} character.

$\sigma^2 p_i$ = Phenotypic variance for i^{th} character.

Phenotypic and genotypic coefficient of variation (expressed in %) were calculated by using the formula given by Burton (1952). Genotypic coefficient of variation (GCV) was calculated as below:

$$\text{GCV\%} = \frac{\sqrt{\sigma^2 g_i}}{\bar{X}_i} \times 100$$

Phenotypic coefficient of variation (PCV)

$$\text{PCV\%} = \frac{\sqrt{\sigma^2 p_i}}{\bar{X}_i} \times 100$$

Where,

\bar{X}_i = General mean of the i^{th} character under consideration

$\sigma^2 g_i$ and $\sigma^2 p_i$ = Genotypic and phenotypic standard deviation of the i^{th} character respectively

Heritability and genetic advance

Heritability (broad sense) which is ratio of genotypic variance to the total phenotypic variance is symbolized as h^2 (BS) and expressed in percentage. Estimation of heritability was done as per the formula given by Hanson *et al.* (1956).

$$h^2 \text{ (BS)} = \frac{\sigma^2 g_i}{\sigma^2 p_i} \times 100$$

or

$$= \frac{\text{Genotypic variance of the } i^{\text{th}} \text{ character}}{\text{Phenotypic variance of the } i^{\text{th}} \text{ character}}$$

Expected genetic advance was calculated by using the method suggested by Johnson *et al.* (1955) at 5% selection intensity.

$$\text{Genetic advance (GA)} = K \cdot P_i \cdot h_i^2$$

Genetic advance as percentage of mean was calculated as follows:

$$\frac{\text{Genetic advance}}{\bar{X}_i}$$

Where,

K= Selection intensity its value at 5% selection level is 2.06

P_i = Phenotypic standard deviation of the i^{th} character

h^2_i = Broad sense heritability (fraction) of the i^{th} character

\bar{X}_i = General mean of the i^{th} character under consideration

Correlation coefficients

Correlation coefficients were calculated in all possible combinations taking all the characters in to consideration at genotypic, phenotypic and environmental levels by using the formula as proposed by Miller *et al.* (1958).

$$\Sigma xy - \frac{\Sigma x \times \Sigma y}{n}$$

$$r = \frac{\Sigma xy - \frac{\Sigma x \times \Sigma y}{n}}{\sqrt{(\Sigma x^2 - \frac{(\Sigma x)^2}{n}) (\Sigma y^2 - \frac{(\Sigma y)^2}{n})}}$$

Where,

r = Correlation coefficient

n = Number of treatments

X and Y = Character under study

Genotypic, phenotypic and environmental correlations were computed by substituting corresponding variance and covariance in the above formula, e.g.

$$r_G (X_i X_j) = \frac{G \text{ Cov } (X_i X_j)}{\sqrt{V_G (X_i) \cdot V_G (X_j)}}$$

$$r_P (X_i X_j) = \frac{P \text{ Cov } (X_i X_j)}{\sqrt{V_P (X_i) \cdot V_P (X_j)}}$$

$$r_E (X_i X_j) = \frac{E \text{ Cov } (X_i X_j)}{\sqrt{V_E (X_i) \cdot V_E (X_j)}}$$

Testing of correlations

The phenotypic correlations are tested for their significance by following formula based on "t" test:

$$t_c = \frac{\sqrt{n-2}}{\sqrt{(1-R^2)}} \quad \text{at } (n-2) \text{ d.f.}$$

Where,

N= Number of treatments.

R= phenotypic correlations coefficient.

The calculated value of “t” is compared with table of “t” at (n-2) d.f. If the calculated value is equal to or greater than table value, it is significant at given probability level. If $t_c < t_T$, it is non significant.

Path coefficient analysis:

The proportion of direct and indirect contributions of various characteristics to the total correlation coefficients with seed yield was estimated through path coefficient analysis as suggested by Wright (1921, 1934) and elaborated by Dewey and Lu (1959).

Path coefficient is a standardized partial regression, which measures the direct influence of one variable upon another and allows partition of correlation coefficient into components of direct and indirect effects.

To estimate various direct and indirect effects, the following set of simultaneous equations were formed and solved.

$$r_{1y} = P_{1y} + r_{12}P_{2y} + r_{13}P_{3y} + \dots + r_{1l}P_{ly}$$

$$r_{2y} = r_{2y}P_{1y} + P_{2y} + r_{23}P_{3y} + \dots + r_{2l}P_{ly}$$

$$r_{ly} = r_{l1}P_{1y} + r_{l2}P_{2y} + r_{l3}P_{3y} + \dots + P_{ly}$$

Where,

r_{1y} to r_{ly} = Coefficient of correlation between causal factor 1 to l and dependent character y,

r_{12} to $r_{l-1,l}$ = Coefficient of correlation among causal factors themselves, and

P_{1y} to P_{ly} = Direct effects of characters 1 to l on character y.

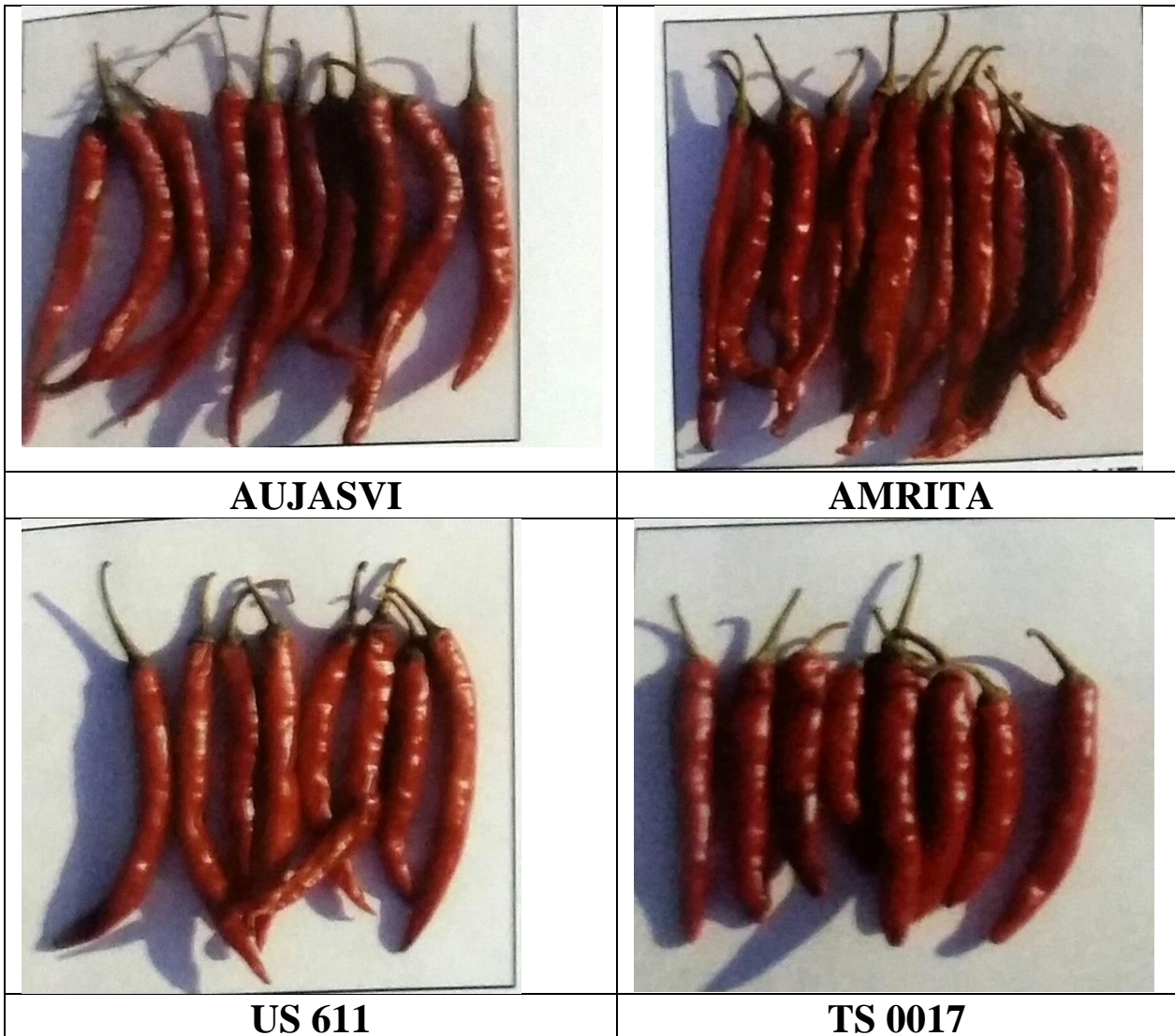
Residual effect, which measures the contribution of the characters not considered in the causal scheme, was obtained as:

$$\text{Residual effect } (P_{RY}) = \sqrt{1 - R^2}$$

Where,

$$R^2 = \sum_{iy} P_i^2 Y + 2 \sum_{\substack{i \neq j \\ i > j}} P_{iy} P_{jy} R_{ij}$$

Varieties of Chilli





TIGER



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CHAPTER - IV

RESULTS

The results obtained from the present investigation entitle “Assessment of Morphological, Fruit Yield and Quality Characters in Hybrid Chilli (*Capsicum annum* L.)” are presented under the following heads:

4.1 Analysis of variance

The analysis of variances for all the characters studied has been presented in Appendix – I to III. Mean squares due to genotypes were highly significantly for all the characters, indicating that the presence of genetic diversity in the existing material.

4.1.1 Mean and range performance of the genotypes

Range and mean performance (Minimum and Maximum) of the ten genotypes of chilli for all the thirty three characters are presented in the table 4.1, 4.2,4.3, 4.4,4.5 and 4.6

4.1.1.1 Plant height (cm)

Plant height varied from 21.97 to 33.33 cm, 38.33 to 64.67 cm, 60.67 to 85.00 cm, 62.33 to 97.00 cm and 64.67 to 97.00 cm with overall mean performance of 28.32, 49.70, 71.13, 76.34 and 76.98 cm at 30, 60, 90, 120 DAT and at maturity, respectively. Hybrid Amrita was observed the tall plant 33.33, 64.67, 85.00, 97.00, and 97.00 cm and genotype S-586 was recorded the dwarf plant with height of 21.97, 38.33, 60.67, 62.33, and 64.67 cm at 30, 60, 90, 120 DAT and at maturity, respectively.

4.1.1.2 Number of primary branches per plant

Range of number of primary branches per plant was 3.67 to 6.00, 7.33 to 12.67, 8.67 to 16.67, 13.33 to 23.00 and 13.33 to 23.00 with an a mean performance of 4.80, 10.30, 13.11, 18.39 and 18.39 at 30, 60, 90, 120 and at maturity, respectively. Maximum number of primary branches 6.00, 12.67, 16.67, 23.00 and 23.00 per plant was observed in hybrid Amrita, while it as recorded minimum in hybrid (S-586) 3.67, 7.33, 8.67, 13.33 and 13.33 at 30, 60, 90, 120 DAT and at maturity, respectively.

4.1.1.3 Number of secondary branches per plant

Range of number of secondary branches per plant was 7.33 to 18.00, 16.00 to 30.67, 25.00 to 41.00, 33.67 to 51.00 and 35.33 to 51.00 with an a mean performance of 14.99, 24.91, 34.89, 40.78 and 42.50 at 30, 60, 90, 120 and at maturity, respectively. Maximum number of secondary branches 18.00, 30.67, 41.00, 51.00 and 51.00 per plant was observed in hybrid Amrita, while it as recorded minimum in hybrid (S-586) 7.33, 16.00, 25.00, 33.67 and 35.33 at 30, 60, 90, 120 DAT and at maturity, respectively.

4.1.1.4 Days to first flower initiation

Early first flower initiation was recorded in Amrita (32.33 days) and hybrid S-586 exhibited in late initiation of first flowering (48.00 days). The overall mean performance of 39.22 days and it was range from 32.33 to 48.00 days.

4.1.1.5 Days to 50% flowering

Days to 50% flowering varied from 62.19 to 70.19 days with overall mean performance to 66.14 days. Hybrid Amrita was recorded the early days to 50% flowering 62.19 days, therefore, hybrid S-586 exhibited in late group which is maximum days to 50% flowering 70.19 days.

4.1.1.6 Days to first picking of fruits

Days to first picking of fruits ranged from 71.19 to 79.19 days with a mean of 75.14 days. Hybrid Amrita was recorded the early first picking in 71.19 days, therefore, hybrid S-586 exhibited in late picking of fruits 79.19 days.

4.1.1.7 Days to first fruit initiation

Days to first fruit initiation varied from 67.52 to 75.52 days with overall mean performance to 71.47 days. Hybrid Amrita was recorded the early fruit initiation in 67.52 days, therefore, hybrid S-586 exhibited in late group which fruit initiation in 75.52 days.

4.1.1.8 Number of fruits per clusters

Number of fruits per cluster varied from 8.10 to 18.47 fruits per cluster and average was 13.12 fruits. The maximum 18.47 fruits per cluster were recorded in hybrid Amrita, while it was found minimum (8.10 fruits) in hybrid S-586.

4.1.1.9 Fruit width (cm)

Hybrid Amrita showed maximum fruit width (0.39 cm) and it was found minimum in hybrid S-586 (0.16 cm). The average for fruit width was 0.22 cm and it varied from 0.16 to 0.39 cm.

4.1.1.10 Fruit length (cm)

Hybrid Amrita showed maximum fruit length (7.69 cm) and it was found minimum in hybrid S-586 (3.37 cm). The average for fruit length was 5.16 cm and it varied from 3.37 to 7.69 cm.

4.1.1.11 Number of fruits per plant

Number of fruits per plant varied from 86.00 to 146.00 fruits per plant and average was 108.41 fruits. The maximum 146.00 fruits per plant were recorded in hybrid Amrita, while it was found minimum (86.00 fruit) in hybrid S-586.

4.1.1.12 Days to maturity

Days to maturity range between 145.71 to 164.21 days with a overall mean performance of 154.78 days. Early maturity was obtained under the hybrid Smriti (145.71 days) and late in the Tara (164.21 days).

4.1.1.13 Fresh fruit yield per plant (g)

Hybrid Tara was recorded maximum fresh fruit yield per plant (692.39 g) followed by TS-0017 (635.51 g) and Jwala garima-12 (616.55 g). Whereas, it was recorded minimum in Smriti (385.58 g). The average yield per plant was 501.41 g and it ranged from 385.58 to 692.39 g.

4.1.1.14 Fresh fruits yield per plot (kg)

hybrid Tara was recorded maximum fresh fruits yield per plot (20.77 kg) followed by TS-0017 (19.07 kg) and Jwala garima- 12 (18.50 kg). Whereas, it was recorded minimum in Smriti (11.57 kg). The average yield per plot was 15.04 kg and it ranged from 11.57 to 20.77 kg.

4.1.1.15 Fresh fruits yield per ha (q)

Hybrid Tara was recorded maximum fresh fruits yield per ha (276.95 q) followed by TS-0017 (254.20 q) and Jwala garima- 12 (246.62 q). Whereas, it was recorded minimum in Smriti (154.23 q). The average yield per ha was 200.57 q and it ranged from 154.23 to 276.95 q.

4.1.1.16 Dry fruits yield per plant (g)

Hybrid Tara was recorded maximum dry fruit yield per plant (230.80 g) followed by TS-0017 (211.84 g) and Jwala Garima- 12 (205.51 g). Whereas, it was recorded minimum in Smriti (128.53 g). The average dry fruit yield per plant was 167.14 g and it ranged from 128.53 to 230.80 g.

4.1.1.17 Dry fruits yield per plot (kg)

Dry fruits yield per plot varied from 3.86 to 6.92 kg with an overall mean performance of 5.01 kg. Highest 6.92, 6.36, 6.17 and 5.74 kg dry fruits yield per plot were recorded in Hybrid Tara, TS-0017, Jwala garima-12 and P. jyoti, respectively and which were at par with each other. However lowest 3.86 kg dry fruits yield per plot exhibited in Smriti.

4.1.1.18 Dry fruit yield per ha (q)

Highest 92.32 q per ha, 84.73 q per ha, 82.21 q per ha and 76.51 q per ha were recorded in hybrid Tara, TS-0017, Jwala garima-12 and P. jyoti, respectively and which were at par with each other. However lowest 51.41 q per ha dry fruits yield per ha exhibited in Smriti.

4.1.1.19 Number of seeds per fruit

Number of seed per fruit varied from 22.67 to 37.00 with the average performance of 28.37 seed. Hybrid Tiger was observed the maximum seeds per fruit (37.00), while it was noted minimum (22.67) in hybrid Marshal 1857 and Shourya.

4.1.1.20 1000 seed weight (g)

1000 seed weight varied from 7.56 to 12.33 g with an overall mean performance of 9.46 g. Hybrid Tiger was recorded maximum 1000 seed weight (12.33 g), while was showed minimum in hybrid Shourya (7.56 g).

Table 4.1 Mean performance of Chilli hybrids for growth characters

| Character | Plant Height (cm) 30 DAT | Plant Height (cm) 60 DAT | Plant Height (cm) 90 DAT | Plant Height (cm) 120 DAT | Plant Height (cm) Maturity |
|------------------|---------------------------------|---------------------------------|---------------------------------|----------------------------------|-----------------------------------|
| Smriti | 22.67 | 42.27 | 63.00 | 68.33 | 70.00 |
| Aujasvi | 28.67 | 45.40 | 65.50 | 69.67 | 70.50 |
| Shourya | 26.17 | 46.27 | 65.67 | 70.00 | 70.67 |
| Albeli-333 | 29.67 | 46.50 | 66.90 | 70.50 | 71.67 |
| Ujjala | 27.67 | 47.33 | 68.00 | 73.00 | 73.00 |
| Tiger | 29.00 | 49.67 | 70.33 | 73.67 | 73.67 |
| ARCH 930 | 30.33 | 54.33 | 78.00 | 83.00 | 83.00 |
| US 612 | 32.33 | 59.00 | 80.67 | 89.00 | 89.00 |
| US 611Tara | 33.33 | 62.33 | 82.00 | 91.00 | 91.00 |
| Marshal 1857 | 24.50 | 40.50 | 61.00 | 65.00 | 66.67 |
| Saniya (03) | 29.33 | 49.83 | 73.00 | 75.33 | 75.33 |
| Jalsa | 30.67 | 53.33 | 77.33 | 82.67 | 82.67 |
| S-586 | 21.97 | 38.33 | 60.67 | 62.33 | 64.67 |
| Amrita | 33.33 | 64.67 | 85.00 | 97.00 | 97.00 |
| P. jyoti | 30.00 | 51.67 | 76.00 | 81.33 | 81.33 |
| Jwala garima-12 | 30.00 | 58.00 | 80.00 | 85.67 | 85.67 |
| TS-0017 | 24.67 | 40.67 | 62.33 | 67.00 | 69.60 |
| Tara | 25.50 | 44.40 | 65.00 | 69.60 | 70.33 |
| Mean | 28.32 | 49.69 | 71.13 | 76.34 | 76.99 |
| S.Em. | 0.78 | 0.81 | 1.05 | 1.61 | 1.17 |
| C.D. at 5% | 2.25 | 2.33 | 3.02 | 4.63 | 3.37 |
| Range Lowest | 21.97 | 38.33 | 60.67 | 62.33 | 64.67 |
| Range Highest | 33.33 | 64.67 | 85.00 | 97.00 | 97.00 |

Table 4.2 Mean performance of Chilli hybrids for growth characters

| Character | Primary Branches per Plant 30 DAT | Primary Branches per Plant 60 DAT | Primary Branches per Plant 90 DAT | Primary Branches per Plant 120 DAT | Primary Branches per Plant Maturity |
|------------------|--|--|--|---|--|
| Smriti | 4.33 | 8.67 | 10.67 | 15.00 | 15.00 |
| Aujasvi | 4.33 | 9.00 | 11.33 | 16.33 | 16.33 |
| Shourya | 4.67 | 10.00 | 12.33 | 18.00 | 18.00 |
| Albeli-333 | 4.67 | 10.33 | 12.33 | 18.00 | 18.00 |
| Ujjala | 4.67 | 10.33 | 12.67 | 18.33 | 18.33 |
| Tiger | 4.67 | 10.67 | 14.00 | 18.67 | 18.67 |
| ARCH 930 | 5.33 | 11.67 | 15.00 | 21.67 | 21.67 |
| US 612 | 5.33 | 12.00 | 16.00 | 22.00 | 22.00 |
| US 611Tara | 5.67 | 12.33 | 16.67 | 22.00 | 22.00 |
| Marshal 1857 | 4.00 | 7.67 | 9.33 | 14.00 | 14.00 |
| Saniya (03) | 5.00 | 10.67 | 14.33 | 19.00 | 19.00 |
| Jalsa | 5.00 | 11.33 | 14.67 | 21.00 | 21.00 |
| S-586 | 3.67 | 7.33 | 8.67 | 13.33 | 13.33 |
| Amrita | 6.00 | 12.67 | 16.67 | 23.00 | 23.00 |
| P. jyoti | 5.00 | 11.33 | 14.67 | 20.00 | 20.00 |
| Jwala garima-12 | 5.33 | 11.67 | 15.67 | 21.67 | 21.67 |
| TS-0017 | 4.33 | 8.67 | 10.00 | 14.00 | 14.00 |
| Tara | 4.33 | 9.00 | 11.00 | 15.00 | 15.00 |
| Mean | 4.80 | 10.30 | 13.11 | 18.39 | 18.39 |
| S.Em. | 0.32 | 0.79 | 0.55 | 0.66 | 0.66 |
| C.D. 5% | 0.93 | 2.27 | 1.58 | 1.90 | 1.90 |
| Range Lowest | 3.67 | 7.33 | 8.67 | 13.33 | 13.33 |
| Range Highest | 6.00 | 12.67 | 16.67 | 23.00 | 23.00 |

Table 4.3 Mean performance of Chilli hybrids for growth characters

| Character | Secondary Branches per Plant 30 DAT | Secondary Branches per Plant 60 DAT | Secondary Branches per Plant 90 DAT | Secondary Branches per Plant 120 DAT | Secondary Branches per Plant Maturity |
|------------------|--|--|--|---|--|
| Smriti | 14.33 | 19.00 | 27.67 | 35.33 | 36.67 |
| Aujasvi | 15.00 | 19.33 | 32.67 | 36.33 | 39.00 |
| Shourya | 15.00 | 25.33 | 35.00 | 37.67 | 40.67 |
| Albeli-333 | 15.33 | 27.00 | 35.33 | 39.00 | 41.67 |
| Ujjala | 15.67 | 27.00 | 37.00 | 40.33 | 42.67 |
| Tiger | 15.67 | 27.67 | 37.67 | 41.00 | 43.00 |
| ARCH 930 | 16.67 | 28.67 | 39.00 | 46.00 | 47.00 |
| US 612 | 17.33 | 29.00 | 39.67 | 46.67 | 48.33 |
| US 611Tara | 17.33 | 30.67 | 40.00 | 48.33 | 49.00 |
| Marshal 1857 | 9.00 | 17.00 | 26.33 | 33.67 | 36.00 |
| Saniya (03) | 16.00 | 28.00 | 37.67 | 41.67 | 43.33 |
| Jalsa | 16.33 | 28.67 | 38.33 | 45.33 | 46.67 |
| S-586 | 7.33 | 16.00 | 25.00 | 33.67 | 35.33 |
| Amrita | 18.00 | 30.67 | 41.00 | 51.00 | 51.00 |
| P. jyoti | 16.33 | 28.00 | 38.00 | 41.67 | 43.67 |
| Jwala garima-12 | 17.00 | 29.00 | 39.67 | 46.00 | 47.67 |
| TS-0017 | 12.33 | 18.00 | 27.33 | 34.67 | 36.33 |
| Tara | 15.00 | 19.33 | 30.67 | 35.67 | 37.00 |
| Mean | 14.98 | 24.91 | 34.89 | 40.78 | 42.50 |
| S.Em. | 0.66 | 0.96 | 0.68 | 0.93 | 1.42 |
| C.D. 5% | 1.90 | 2.76 | 1.96 | 2.67 | 4.07 |
| Range Lowest | 7.33 | 16.00 | 25.00 | 33.67 | 35.33 |
| Range Highest | 18.00 | 30.67 | 41.00 | 51.00 | 51.00 |

Table 4.4 Mean performance of Chilli hybrids for yield characters

| Character | Days to First Picking of Fruits | Days of 50% Flowering | Days to First Fruit Initiation | Fruits per Cluster | Days to First Flower Initiation |
|-----------------|---------------------------------|-----------------------|--------------------------------|--------------------|---------------------------------|
| Smriti | 77.19 | 68.19 | 73.52 | 10.47 | 44.67 |
| Aujasvi | 76.19 | 67.19 | 72.52 | 12.43 | 43.33 |
| Shourya | 75.69 | 66.69 | 72.02 | 12.47 | 43.00 |
| Albeli-333 | 75.58 | 66.58 | 71.91 | 12.47 | 39.67 |
| Ujjala | 75.19 | 66.19 | 71.52 | 12.50 | 37.00 |
| Tiger | 75.19 | 66.19 | 71.52 | 13.47 | 36.67 |
| ARCH 930 | 74.03 | 65.03 | 70.37 | 14.48 | 35.33 |
| US 612 | 72.03 | 63.03 | 68.37 | 16.47 | 34.33 |
| US 611Tara | 71.58 | 62.58 | 67.91 | 17.80 | 33.33 |
| Marshal 1857 | 77.58 | 68.58 | 73.91 | 8.37 | 46.67 |
| Saniya (03) | 75.03 | 66.03 | 71.37 | 13.47 | 36.33 |
| Jalsa | 74.58 | 65.58 | 70.91 | 14.47 | 35.67 |
| S-586 | 79.19 | 70.19 | 75.52 | 8.10 | 48.00 |
| Amrita | 71.19 | 62.19 | 67.52 | 18.47 | 32.33 |
| P. jyoti | 75.03 | 66.03 | 71.37 | 13.47 | 35.67 |
| Jwala garima-12 | 73.19 | 64.19 | 69.52 | 16.43 | 35.33 |
| TS-0017 | 77.58 | 68.58 | 73.91 | 9.47 | 45.33 |
| Tara | 76.53 | 67.53 | 72.87 | 11.40 | 43.33 |
| Mean | 75.14 | 66.14 | 71.47 | 13.12 | 39.22 |
| S.Em. | 0.54 | 0.54 | 1.82 | 0.90 | 1.46 |
| C.D. 5% | 1.55 | 1.55 | 5.22 | 2.58 | 4.19 |
| Range Lowest | 71.19 | 62.19 | 67.52 | 8.10 | 32.33 |
| Range Highest | 79.19 | 70.19 | 75.52 | 18.47 | 48.00 |

Table 4.5 Mean performance of Chilli hybrids for yield characters

| Character | Fruit Width (cm) | Fruit Length (cm) | Fruits per Plant | Days to Maturity | Fruit Fresh Weight Plant (g) |
|------------------|-------------------------|--------------------------|-------------------------|-------------------------|-------------------------------------|
| Smriti | 0.17 | 4.27 | 92.00 | 145.71 | 385.58 |
| Aujasvi | 0.18 | 4.38 | 98.00 | 146.61 | 394.52 |
| Shourya | 0.19 | 4.40 | 100.00 | 147.41 | 403.46 |
| Albeli-333 | 0.19 | 4.71 | 103.00 | 148.51 | 412.40 |
| Ujjala | 0.19 | 5.38 | 105.00 | 149.61 | 421.34 |
| Tiger | 0.19 | 5.43 | 108.00 | 149.91 | 439.22 |
| ARCH 930 | 0.24 | 5.75 | 119.00 | 151.61 | 448.16 |
| US 612 | 0.29 | 6.17 | 130.00 | 152.11 | 461.57 |
| US 611Tara | 0.30 | 7.15 | 134.00 | 153.41 | 470.51 |
| Marshal 1857 | 0.16 | 3.40 | 88.00 | 155.81 | 512.27 |
| Saniya (03) | 0.20 | 5.51 | 109.00 | 157.61 | 517.01 |
| Jalsa | 0.22 | 5.75 | 115.00 | 158.51 | 531.23 |
| S-586 | 0.16 | 3.37 | 86.00 | 159.81 | 545.45 |
| Amrita | 0.39 | 7.69 | 146.00 | 160.61 | 564.41 |
| P. jyoti | 0.21 | 5.61 | 112.00 | 161.01 | 573.89 |
| Jwala garima-12 | 0.29 | 5.92 | 121.00 | 161.61 | 616.55 |
| TS-0017 | 0.16 | 3.72 | 90.00 | 161.91 | 635.51 |
| Tara | 0.17 | 4.38 | 95.33 | 164.21 | 692.39 |
| Mean | 0.22 | 5.16 | 108.41 | 154.78 | 501.41 |
| S.Em. | 0.03 | 0.07 | 0.76 | 2.02 | 54.17 |
| C.D. 5% | 0.10 | 0.20 | 2.18 | 5.81 | 155.70 |
| Range Lowest | 0.16 | 3.37 | 86.00 | 145.71 | 385.58 |
| Range Highest | 0.39 | 7.69 | 146.00 | 164.21 | 692.39 |

Table 4.6 Mean performance of Chilli hybrids for yield characters

| Character | Fruit Fresh Weight per Plot (kg) | Fruit Fresh Weight per ha. (q.) | Fruit Dry Weight per Plant (g) | Fruit Dry Weight per Plot (kg) | Fruit Dry Weight per ha. (q.) | Seeds per Fruit | 1000 Seed Weight (g) |
|------------------|---|--|---------------------------------------|---------------------------------------|--------------------------------------|------------------------|-----------------------------|
| Smriti | 11.57 | 154.23 | 128.53 | 3.86 | 51.41 | 27.67 | 9.22 |
| Aujasvi | 11.83 | 157.81 | 131.50 | 3.94 | 52.60 | 26.33 | 8.78 |
| Shourya | 12.10 | 161.38 | 134.49 | 4.04 | 53.80 | 22.67 | 7.55 |
| Albeli-333 | 12.37 | 164.96 | 137.47 | 4.12 | 54.99 | 26.67 | 8.89 |
| Ujjala | 12.64 | 168.53 | 140.44 | 4.21 | 56.18 | 34.67 | 11.55 |
| Tiger | 13.17 | 175.69 | 146.41 | 4.39 | 58.56 | 37.00 | 12.33 |
| ARCH 930 | 13.44 | 179.26 | 149.38 | 4.48 | 59.75 | 30.33 | 10.11 |
| US 612 | 13.85 | 184.63 | 153.85 | 4.62 | 61.54 | 32.33 | 10.78 |
| US 611Tara | 14.12 | 188.20 | 156.84 | 4.71 | 62.74 | 28.00 | 9.33 |
| Marshal 1857 | 15.37 | 204.91 | 170.76 | 5.12 | 68.30 | 22.67 | 7.55 |
| Saniya (03) | 15.51 | 206.80 | 172.33 | 5.17 | 68.93 | 23.33 | 7.78 |
| Jalsa | 15.94 | 212.49 | 177.07 | 5.31 | 70.83 | 25.33 | 8.44 |
| S-586 | 16.36 | 218.18 | 181.81 | 5.45 | 72.73 | 27.00 | 9.00 |
| Amrita | 16.93 | 225.76 | 188.14 | 5.64 | 75.25 | 29.00 | 9.67 |
| P. jyoti | 17.22 | 229.55 | 191.29 | 5.74 | 76.52 | 30.00 | 10.00 |
| Jwala garima-12 | 18.50 | 246.62 | 205.51 | 6.17 | 82.21 | 27.00 | 9.00 |
| TS-0017 | 19.07 | 254.20 | 211.84 | 6.36 | 84.73 | 26.67 | 8.89 |
| Tara | 20.77 | 276.95 | 230.80 | 6.92 | 92.32 | 34.00 | 11.33 |
| Mean | 15.04 | 200.56 | 167.14 | 5.01 | 66.85 | 28.37 | 9.46 |
| S.Em. | 1.63 | 21.67 | 18.06 | 0.54 | 7.22 | 0.69 | 0.23 |
| C.D. 5% | 4.67 | 62.28 | 51.90 | 1.56 | 20.76 | 1.98 | 0.66 |
| Range Lowest | 11.57 | 154.23 | 128.53 | 3.86 | 51.41 | 22.67 | 7.55 |
| Range Highest | 20.77 | 276.95 | 230.80 | 6.92 | 92.32 | 37.00 | 12.33 |

4.2 Genetic Variability :

Environmental coefficient of variation (ECV), Genotypic coefficient of variation (GCV) and phenotypic coefficient of variation (PCV)

The PCV was higher in its corresponding GCV and ECV for all the parameters under present study. Data pertaining to genetic parameters are presented in Table 4.7.

4.2.1 Genotypic coefficient of variation (GCV)

The highest GCV value was recorded for fruits width cm (23.99 %), while moderate for fruit length cm (23.10 %), fruits per cluster (21.54 %), secondary branches per plant at 60 DAT (20.30 %), primary branches per plant 90 DAT (18.90 %), secondary branches per plant 30 DAT (18.25 %), primary branches per plant maturity (16.83 %), primary branches per plant 120 DAT (16.83 %), plant height (cm) 60 DAT (15.37 %), fruits per plant (15.45 %), secondary branches per plant 90 DAT (14.98 %), fruit fresh weight plant (g), Fruit fresh weight per plot (kg), fruit fresh weight per ha. (q.), fruit dry weight per plant (g), fruit dry weight per plot (kg), fruit dry weight per ha. (q.) (14.37 %), seeds per fruit, 1000 seed weight (g) (14.16 %), primary branches per plant 60 DAT (13.46 %), secondary branches per plant 120 DAT (13.11 %), plant height (cm) 120 DAT (12.76 %), days to first flower initiation (12.26 %), plant height (cm) maturity (11.90 %), plant height (cm) 30 DAT (11.74 %), plant height (cm) 90 DAT (11.09 %), secondary branches per plant maturity (11.32), primary branches per plant 30 DAT (10.42), days to maturity (3.68 %), days of 50% flowering (3.16 %), days to first picking of fruits (2.78 %) and days to first fruit initiation (1.64 %).

4.2.2 Phenotypic coefficient of variation (PCV)

The highest PCV value was recorded for fruits width cm (36.00 %), while moderate for fruits per cluster (24.59 %), fruit fresh weight per plot (kg), fruit dry weight per plot (kg) (23.60 %), fruit fresh weight plant (g), fruit fresh weight per ha. (q.), fruit dry weight per plant (g), fruit dry weight per ha. (q.) (23.59 %), Fruit length cm (23.22 %),

Low value for PCV was recorded for days to first fruit initiation (4.69 %), days to maturity (4.32 %), days to 50 % flowering (3.46 %) and days to first fruit picking (3.05 %).

4.2.3 Environmental coefficient of variation (PCV)

The highest ECV value was recorded for fruits width cm (26.85 %), while moderate for fruit fresh weight per plot (kg), fruit dry weight per plot (kg) (18.72 %), fruit fresh weight per plant (g), fruit fresh weight per ha. (q.), fruit dry weight per plant (g), fruit dry weight per ha (q) (18.71 %).

Low value for ECV was recorded for plant height (cm) at 60 DAT (2.83 %), plant height (cm) at 90 DAT (2.56 %), days to maturity (2.26 %), Days to 50 % flowering (1.41 %), days to first fruit picking (1.24 %) and fruits per plant (1.21 %).

4.3. Heritability

Estimates with regard to heritability (broad sense) have been presented in Table 4.7. Heritability estimates varied from parameter to parameter and also with the change in environment for the same parameters. Therefore heritability estimates in broad sense were classified into three groups: - High more than 80, medium 50 to 80 and low less than 50.

Heritability estimates in broad sense were high for plant height (cm) 30 DAT (86 %), plant height (cm) 60 DAT (97 %), plant height (cm) 90 DAT (95 %), plant height (cm) 120 DAT (92 %), plant height (cm) maturity (95 %), primary branches per plant 90 DAT (87 %), primary branches per plant 120 DAT (88 %), primary branches per plant maturity (88 %), secondary branches per plant 30 DAT (85 %), secondary branches per plant 60 DAT (90 %), secondary branches per plant 90 DAT (95 %), secondary branches per plant 120 DAT (92 %), days to first picking of fruits (83.5), fruit length (cm) (99 %), fruits per plant (99 %), seeds per fruit (92 %), 1000 Seed Weight (g) (92 %).

Heritability estimates in broad sense were medium for Primary Branches per Plant 60 DAT (51 %), Secondary Branches per Plant Maturity (79 %), Days to First Flower Initiation (78 %), Fruits per cluster (77 %), Days to maturity (73 %).

Heritability estimates in broad sense were low for primary branches per plant 30 DAT (44 %), fruit width (44 %), days to maturity, fruit fresh weight plant

(g), fruit fresh weight per plot (kg), fruit fresh weight per ha. (q.), fruit dry weight per plant (g), fruit dry weight per plot (kg), fruit dry weight per ha. (q.) (37 %).

4.4 Genetic advance

High value of genetic advance was recorded for Fruit Fresh weight plant (g) (90.39) followed by Fruit Fresh weight per ha. (q) (36.16), Fruit per plant (34.40), Fruit Dry weight per plant (g) (30.13), Plant Height (cm) 120 DAT (19.29), Plant Height (cm) Maturity (18.42), Plant Height (cm) 90 DAT (15.83) and Plant Height (cm) 60 DAT (15.47).

4.5 Genetic advance as percentage of mean

Genetic advance was computed as percentage as mean of the parameter over environment. It has been classified as high (more than 40), medium (25 to 40) and low (less than 25).

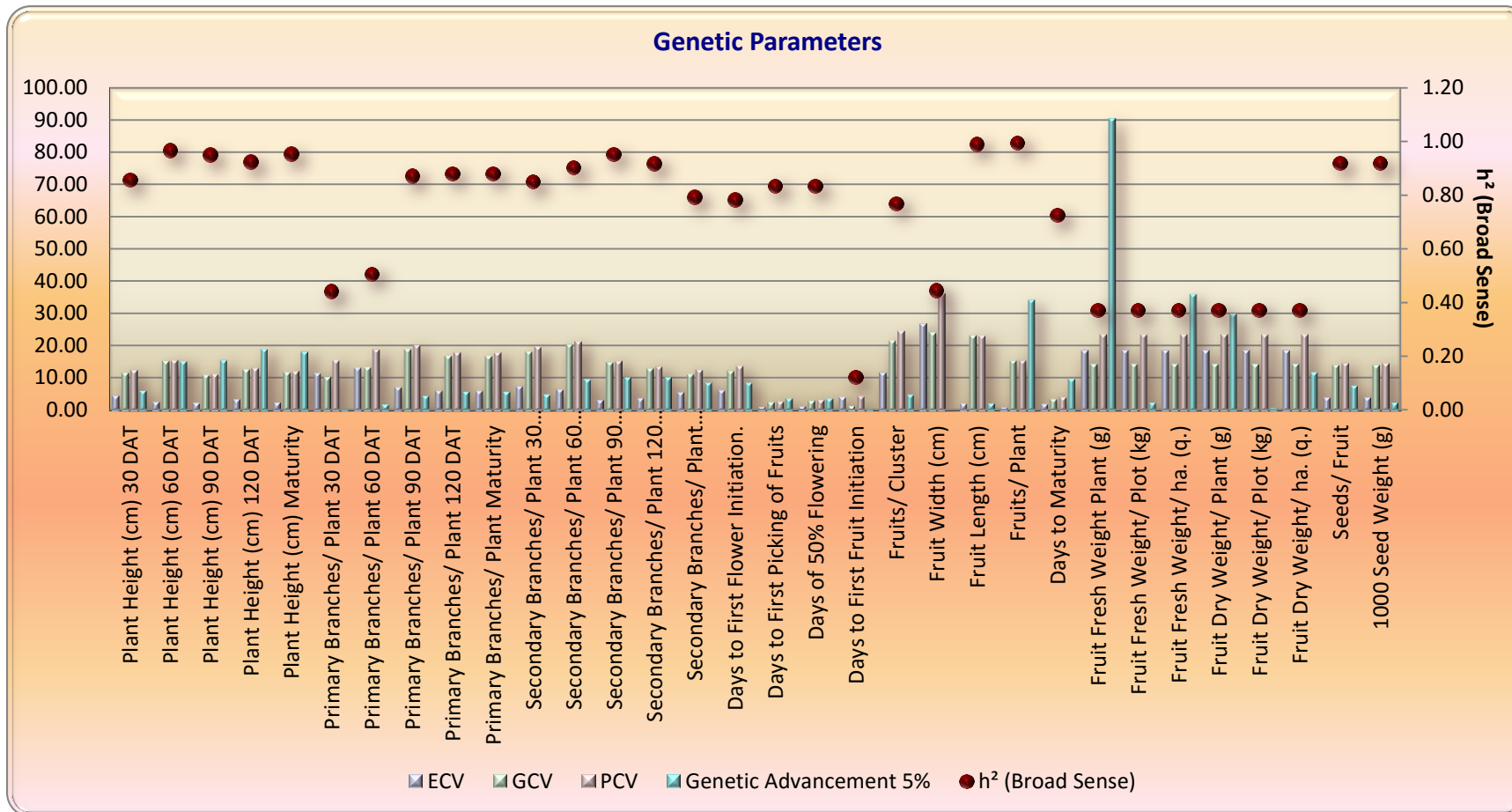
Medium value of genetic advance as percentage of mean was recorded for plant height (cm) 60 DAT (31.13 %), plant height (cm) 120 DAT (25.27 %), primary branches per plant 90 DAT (36.34 %), primary branches per plant 120 DAT (32.51 %), primary branches per plant maturity (32.51 %), secondary branches per plant 30 DAT (34.68 %), secondary branches per plant 60 DAT (39.73 %), secondary branches per plant 90 DAT (30.05 %), secondary branches per plant 120 DAT (25.85 %), fruits per cluster (38.88 %), fruit width (cm) (32.92 %), fruit length (cm) (47.34 %), fruits per plant (31.73 %), seeds per fruit (27.96 %) and 1000 seed weight (g) (27.97 %).

Low value of genetic advance as percentage of mean was recorded for plant height (cm) 30 DAT (22.38 %), plant height (cm) 90 DAT (22.25 %), plant height (cm) maturity (23.93 %), primary branches per plant 30 DAT (14.27 %), primary branches per plant 60 DAT (19.72 %), secondary branches per plant maturity (20.76 %), days to first flower initiation (22.35 %), days to first picking of fruits (5.24 %), days of 50% flowering (5.95 %), days to first fruit initiation (1.18 %), days to maturity (6.45 %), fruits fresh weight per plant (g) (18.03 %), fruits fresh weight per plot (kg) (18.02 %), fruits fresh weight per ha. (q.) (18.03 %), fruits dry weight per plant (g) (18.03 %), fruit dry weight per plot (kg) (18.02 %) and fruit dry weight per ha. (q.) (18.03 %).

Table 4.7: Genetic parameters of fruit yield and its attributing characters in chilli hybrids

| | General Mean | PCV | GCV | ECV | Heritability (Broad Sense) % | Genetic Advance 5% | Gen. Adv as % of Mean 5% |
|--|---------------------|------------|------------|------------|-------------------------------------|---------------------------|---------------------------------|
| Plant Height (cm) 30 DAT | 28.32 | 12.68 | 11.74 | 4.80 | 86 | 6.34 | 22.38 |
| Plant Height (cm) 60 DAT | 49.69 | 15.62 | 15.37 | 2.83 | 97 | 15.47 | 31.13 |
| Plant Height (cm) 90 DAT | 71.13 | 11.38 | 11.09 | 2.56 | 95 | 15.83 | 22.25 |
| Plant Height (cm) 120 DAT | 76.34 | 13.28 | 12.76 | 3.66 | 92 | 19.29 | 25.27 |
| Plant Height (cm) Maturity | 76.99 | 12.19 | 11.90 | 2.64 | 95 | 18.42 | 23.93 |
| Primary Branches per Plant 30 DAT | 4.80 | 15.67 | 10.42 | 11.70 | 44 | 0.68 | 14.27 |
| Primary Branches per Plant 60 DAT | 10.30 | 18.93 | 13.46 | 13.31 | 51 | 2.03 | 19.72 |
| Primary Branches per Plant 90 DAT | 13.11 | 20.25 | 18.90 | 7.27 | 87 | 4.76 | 36.34 |
| Primary Branches per Plant 120 DAT | 18.39 | 17.95 | 16.83 | 6.23 | 88 | 5.98 | 32.51 |
| Primary Branches per Plant Maturity | 18.39 | 17.95 | 16.83 | 6.23 | 88 | 5.98 | 32.51 |
| Secondary Branches per Plant 30 DAT | 14.98 | 19.79 | 18.25 | 7.65 | 85 | 5.20 | 34.68 |
| Secondary Branches per Plant 60 DAT | 24.91 | 21.37 | 20.30 | 6.68 | 90 | 9.89 | 39.73 |
| Secondary Branches per Plant 90 DAT | 34.89 | 15.33 | 14.95 | 3.38 | 95 | 10.48 | 30.05 |
| Secondary Branches per Plant 120 DAT | 40.78 | 13.69 | 13.11 | 3.95 | 92 | 10.54 | 25.85 |
| Secondary Branches per Plant Maturity | 42.50 | 12.71 | 11.32 | 5.78 | 79 | 8.82 | 20.76 |
| Days to First Flower Initiation | 39.22 | 13.85 | 12.26 | 6.44 | 78 | 8.77 | 22.35 |
| Days to First Picking of Fruits | 75.14 | 3.05 | 2.78 | 1.24 | 83 | 3.94 | 5.24 |
| Days of 50% Flowering | 66.14 | 3.46 | 3.16 | 1.41 | 83 | 3.94 | 5.95 |
| Days to First Fruit Initiation | 71.47 | 4.69 | 1.64 | 4.40 | 12 | 0.84 | 1.18 |
| Fruits per Cluster | 13.12 | 24.59 | 21.54 | 11.85 | 77 | 5.10 | 38.88 |
| Fruit Width (cm) | 0.22 | 36.00 | 23.99 | 26.85 | 44 | 0.07 | 32.92 |
| Fruit Length (cm) | 5.16 | 23.22 | 23.10 | 2.38 | 99 | 2.44 | 47.34 |
| Fruits per Plant | 108.41 | 15.50 | 15.45 | 1.21 | 99 | 34.40 | 31.73 |
| Days to Maturity | 154.78 | 4.32 | 3.68 | 2.26 | 73 | 9.98 | 6.45 |
| Fruit Fresh Weight Plant (g) | 501.41 | 23.59 | 14.37 | 18.71 | 37 | 90.39 | 18.03 |
| Fruit Fresh Weight per Plot (kg) | 15.04 | 23.60 | 14.37 | 18.72 | 37 | 2.71 | 18.02 |
| Fruit Fresh Weight per ha. (q.) | 200.56 | 23.59 | 14.37 | 18.71 | 37 | 36.16 | 18.03 |
| Fruit Dry Weight per Plant (g) | 167.14 | 23.59 | 14.37 | 18.71 | 37 | 30.13 | 18.03 |
| Fruit Dry Weight per Plot (kg) | 5.01 | 23.60 | 14.37 | 18.72 | 37 | 0.90 | 18.02 |
| Fruit Dry Weight per ha. (q.) | 66.85 | 23.59 | 14.37 | 18.71 | 37 | 12.05 | 18.03 |
| Seeds per Fruit | 28.37 | 14.77 | 14.16 | 4.20 | 92 | 7.93 | 27.96 |
| 1000 Seed Weight (g) | 9.46 | 14.77 | 14.16 | 4.20 | 92 | 2.65 | 27.97 |

Fig 4.1: Genetic parameter of fruit yield and its attributing characters in chilli hybrids.



4.6 Correlation coefficients

Correlation coefficients were worked out at phenotypic, genotypic and environmental levels for all possible combination of eleven yield and its attributing characters (Table 4.8). Results indicated that genotypic correlation coefficient, in general, were of higher magnitude than the corresponding phenotypic correlation coefficient for all the characters.

4.6.1 Genotypic Correlation Coefficients

4.6.1.1 Plant height (cm) at maturity

Plant height expressed positive correlation with primary branches per plant maturity (0.9436), secondary branches per plant maturity (1.0106), fruits per cluster (0.9971), fruit width (cm) (1.1327), fruit length (cm) (0.9616), fruits per plant (0.9906), days to maturity (0.2000) and seeds per fruit (0.1812). Negative association of this character was recorded with days of 50% flowering (-0.9985) and days to first fruit initiation (-1.6590)

4.6.1.2 Number of primary branches per plant at maturity

Correlation coefficient of number of primary branches per plant at maturity was observed positive with number of secondary branches per plant at maturity (1.0839), fruits per cluster (1.0033), fruit width (cm) (1.0036), fruit length (cm) (0.9634), fruits per plant (0.9696) and seeds per fruit (0.1862). Negative association of this character was recorded with days of 50% flowering (-1.0275), days to first fruit initiation (-1.6935) and days to maturity (-0.0377).

4.6.1.3 Number of secondary branches per plant at maturity

Correlation coefficient of number of secondary branches per plant at maturity was observed), fruits per cluster (1.0196), fruit width (cm) (1.1226), fruit length (cm) (0.9968), fruits per plant (1.0151), days to maturity (0.1013) and seeds per fruit (0.1871). Negative association of this character was recorded with days of 50% flowering (-1.0025), days to first fruit initiation (-2.1715)

4.6.1.4 Days to 50% flowering

Association of days to 50% flowering was exhibited positive with days to 1st first fruit initiation (1.9904). Negative association of this character was recorded with fruit per clusters (-1.0861), fruit width (cm) (-1.1540), fruit length

(cm) (-0.9898), fruits per plant (-1.0063), days to maturity (-0.0689) and seeds per fruit (-0.2415).

4.6.1.5 Days to first fruit initiation

Correlation coefficient of days to 1st fruit initiation was found positive and significant with days to maturity (0.2100). Significant and negative association of this character was recorded with fruit per clusters (-2.1356), fruit width (cm) (-1.8653), fruit length (cm) (-1.7986), fruits per plant (-1.7835) and seeds per fruit (-0.5256).

4.6.1.6 Number of Fruits per cluster

Correlation coefficient of fruits per cluster was found positive and significant with fruit width (cm) (1.0558), fruit length (cm) (1.0200), fruits per plant (1.0257) and seeds per fruit (0.1890). Significant and negative association of this character was recorded with days to maturity (-0.0144).

4.6.1.7 Fruit width (cm)

Fruit width expressed a significant and positive association with fruit length (cm) (1.1012), fruits per plant (1.1485), days to maturity (0.0904) and seeds per fruit (0.0889).

4.6.1.8 Fruit length (cm)

Fruit length expressed a significant and positive association with fruits per plant (0.9797), days to maturity (0.1030) and seeds per fruit (0.2930).

4.6.1.9 Fruits per plant

Fruits per plant expressed a significant and positive association with days to maturity (0.1193) and seeds per fruit (0.2271).

4.6.1.10 Days to maturity

Days to maturity expressed a negative association with and seeds per fruit (-0.0453).

4.6.2 Phenotypic Correlation Coefficients

4.6.2.1 Plant height (cm) at maturity

Plant height expressed positive phenotypic correlation with primary branches per plant maturity (0.8860), secondary branches per plant maturity (0.8381), fruits per cluster (0.8567), fruit width (cm) (0.7645), fruit length (cm)

(0.9291), fruits per plant (0.9644), days to maturity (0.2145) and seeds per fruit (0.1700). Negative association of this character was recorded with days of 50% flowering (-0.8712) and days to first fruit initiation (-0.6800)

4.6.2.2 Number of primary branches per plant at maturity

Phenotypic correlation coefficient of number of primary branches per plant at maturity was observed positive with number of secondary branches per plant at maturity (0.8360), fruits per cluster (0.8769), fruit width (cm) (0.6912), fruit length (cm) (0.8916), fruits per plant (0.9032), days to maturity (0.0952) and seeds per fruit (0.1922). Negative association of this character was recorded with days of 50% flowering (-0.8097) and days to first fruit initiation (-0.6459).

4.6.2.3 Number of secondary branches per plant at maturity

Phenotypic correlation coefficient of number of secondary branches per plant at maturity was observed), fruits per cluster (0.8750), fruit width (cm) (0.6410), fruit length (cm) (0.8911), fruits per plant (0.8944) and seeds per fruit (0.2091). Negative association of this character was recorded with days of 50% flowering (-0.8942), days to first fruit initiation (-0.3741) and days to maturity (-0.0087).

4.6.2.4 Days to 50% flowering

Association of days to 50% flowering was exhibited positive with days to 1st first fruit initiation (0.5528) and days to maturity (0.0639). Negative association of this character was recorded with fruit per clusters (-0.8174), fruit width (cm) (-0.6701), fruit length (cm) (-0.9096), fruits per plant (-0.9300) and seeds per fruit (-0.1911).

4.6.2.5 Days to first fruit initiation

Phenotypic correlation coefficient of negative association of this character was recorded with fruit per clusters (-0.4388), fruit width (cm) (-0.5489), fruit length (cm) (-0.6000), fruits per plant (-0.6450), days to maturity (-0.1543) and seeds per fruit (-0.0675).

4.6.2.6 Fruits per cluster

Phenotypic correlation coefficient of fruits per cluster was found positive and significant with fruit width (cm) (0.7609), fruit length (cm) (0.8778), fruits per plant (0.8789), days to maturity (0.1034) and seeds per fruit (0.1034).

4.6.2.7 Fruit width (cm)

Fruit width expressed a phenotypic correlation positive association with fruit length (cm) (0.6969), fruits per plant (0.7566), days to maturity (0.2785) and seeds per fruit (0.1730).

4.6.2.8 Fruit length (cm)

Fruit length expressed a significant and positive association with fruits per plant (0.9724), days to maturity (0.0596) and seeds per fruit (0.2741).

4.6.2.9 Fruits per plant

Fruits per plant expressed a significant and positive association with days to maturity (0.0939) and seeds per fruit (0.2075).

4.6.2.10 Days to maturity

Days to maturity expressed a positive association with seeds per fruit (0.230).

Table 4.8 : Genotypic correlation coefficient of fruit yield and its component characters of chilli hybrids.

| Character | Plant Height (cm) Maturity | Primary Branches per Plant Maturity | Secondary Branches per Plant Maturity | Days of 50% Flowering | Days to First Fruit Initiation | Fruits per Cluster | Fruit Width (cm) | Fruit Length (cm) | Fruit yield per Plant | Days to Maturity | Seeds per Fruit |
|---------------------------------------|----------------------------|-------------------------------------|---------------------------------------|-----------------------|--------------------------------|--------------------|------------------|-------------------|-----------------------|------------------|-----------------|
| Plant Height (cm) Maturity | 1.0000 | 0.9436** | 1.0106** | -0.9985 | -1.6590 | 0.9971** | 1.1372** | 0.9616** | 0.9906** | 0.2000* | 0.1812* |
| Primary Branches per Plant Maturity | | 1.0000 | 1.0839** | -1.0275 | -1.6935 | 1.0033** | 1.0036** | 0.9634** | 0.9696** | -0.0377 | 0.1862* |
| Secondary Branches per Plant Maturity | | | 1.0000 | -1.0025 | -2.1715 | 1.0196** | 1.1226** | 0.9968** | 1.0151** | 0.1013* | 0.1871* |
| Days of 50% Flowering | | | | 1.0000 | 1.9904** | -1.0861 | -1.1540 | -0.9898 | -1.0063 | -0.0689 | -0.2415 |
| Days to First Fruit Initiation | | | | | 1.0000 | -2.1356 | -1.8653 | -1.7986 | -1.7835 | 0.2100* | -0.5256 |
| Fruitsper Cluster | | | | | | 1.0000 | 1.0558** | 1.0200** | 1.0257** | -0.0144 | 0.1890* |
| Fruit Width (cm) | | | | | | | 1.0000 | 1.1012** | 1.1485** | 0.0904* | 0.0889* |
| Fruit Length (cm) | | | | | | | | 1.0000 | 0.9797** | 0.1030* | 0.2935* |
| Fruits per Plant | | | | | | | | | 1.0000 | 0.1193* | 0.2271* |
| Days to Maturity | | | | | | | | | | 1.0000 | -0.0453 |
| Seeds per Fruit | | | | | | | | | | | 1.0000 |

* 5% level of significance

** 1% level of significance

Table 4.9 : Phenotypic correlation coefficient of fruit yield and its component characters of chilli hybrids.

| Character | Plant Height (cm) Maturity | Primary Branches per Plant Maturity | Secondary Branches per Plant Maturity | Days of 50% Flowering | Days to First Fruit Initiation | Fruits per Cluster | Fruit Width (cm) | Fruit Length (cm) | Fruits per Plant | Days to Maturity | Seeds per Fruit |
|--|----------------------------|-------------------------------------|---------------------------------------|-----------------------|--------------------------------|--------------------|------------------|-------------------|------------------|------------------|-----------------|
| Plant Height (cm) Maturity | 1.0000 | 0.8860** | 0.8381** | -0.8712 | -0.6800 | 0.8567** | 0.7645** | 0.9291** | 0.9644** | 0.2145** | 0.1700** |
| Primary Branches per Plant Maturity | | 1.0000 | 0.8360** | -0.8097 | -0.6459 | 0.8769** | 0.6912** | 0.8916** | 0.9032** | 0.0952* | 0.1922* |
| Secondary Branches per Plant Maturity | | | 1.0000 | -0.8942 | -0.3741 | 0.8750** | 0.6410** | 0.8911** | 0.8944** | -0.0087 | 0.2091* |
| Days of 50% Flowering | | | | 1.0000 | 0.5528** | -0.8174 | -0.6701 | -0.9096 | -0.9300 | 0.0639* | -0.1911 |
| Days to First Fruit Initiation | | | | | 1.0000 | -0.4388 | -0.5489 | -0.6000 | -0.6450 | -0.1543 | -0.0675 |
| Fruits per Cluster | | | | | | 1.0000 | 0.7609** | 0.8778** | 0.8789** | 0.1034* | 0.2760* |
| Fruit Width (cm) | | | | | | | 1.0000 | 0.6969** | 0.7566** | 0.2785* | 0.1730* |
| Fruit Length (cm) | | | | | | | | 1.0000 | 0.9724** | 0.0596* | 0.2741* |
| Fruits per Plant | | | | | | | | | 1.0000 | 0.0939* | 0.2075* |
| Days to Maturity | | | | | | | | | | 1.0000 | 0.0230* |
| Seeds per Fruit | | | | | | | | | | | 1.0000 |

* 5% level of significance

** 1% level of significance

E. Path coefficient

A path coefficient is a standardized partial regression coefficient and as such measures the direct influence of one variable upon another and permits partitioning of the correlation coefficients into components of direct and indirect effects. The genotypic and phenotypic correlation coefficients were further partitioned into direct and indirect effects and the results are presented in (Table 4.10).

4.7 Genotypic path coefficient

4.7.1 Direct and Indirect effects

4.7.1.1 Plant height (cm):

Plant height (1.0321) revealed that positive direct effect on fruit dry weight per plant (g) (0.1036) while, it expressed positive indirect correlation primary branches per plant at maturity (0.9739), secondary branches per plant at maturity (1.0431), fruits per cluster (1.0291), fruit width (cm) (1.1737), fruit length (cm) (0.9924), fruits per plant (1.0223), days to maturity (0.2064) and seeds per fruit (0.1870), while it negative indirect effect on days of 50% flowering (-1.0306) and days to first fruit initiation (-1.7122).

4.7.1.2 Number of primary branches per plant

Primary branches per plant (-0.1606) revealed that high negative direct effect on fruit dry weight per plant (g) (-0.2132) while, it expressed positive indirect correlation days to 50% flowering (0.1650), days to first fruit initiation (0.2719) and days to maturity (0.0061), while it negative indirect effect on plant height (cm) at maturity (-0.1515), primary branches per plant at maturity (-0.1606), secondary branches per plant at maturity (-0.1740), fruits per cluster (-0.1611), fruit width (cm) (-0.1612), fruit length (cm) (-0.1547), fruit yield per plant (-0.1557) and seeds per fruit (-0.0299).

4.7.1.3 Number of secondary branches per plant

Secondary branches per plant (-0.4960) revealed that high positive direct effect on fruit dry weight per plant (g) (0.0399) while, it expressed

positive indirect correlation days to 50% flowering (0.4973) and days to first fruit initiation (1.0771), while it negative indirect effect on plant height (cm) at maturity (-0.5013), primary branches per plant at maturity (-0.5377), secondary branches per plant at maturity (-0.4960), fruits per cluster (-0.5058), fruit width (cm) (-0.5569), fruit length (cm) (-0.4944), fruit yield per plant (-0.5035), days to maturity (-0.0503) and seeds per fruit (-0.0928).

4.7.1.4 Days to 50% flowering

Days to 50% flowering (-0.3085) revealed that high negative direct effect on fruit dry weight per plant (g) (-0.0474) while, it expressed positive indirect correlation plant height (cm) at maturity (0.3081), primary branches per plant at maturity (0.3170), secondary branches per plant at maturity (0.3093), fruits per cluster (0.3351), fruit width (cm) (0.3561), fruit length (cm) (0.3054), fruits yield per plant (0.3105), days to maturity (0.0213) and seeds per fruit (0.0745), while it negative indirect effect on days to first fruit initiation (-0.6141).

4.7.1.5 Days to first fruit initiation

Days to first fruit initiation (-0.1642) revealed that high positive direct effect on fruit dry weight per plant (g) (0.7547) while, it expressed positive indirect correlation plant height (cm) at maturity (0.2724), primary branches per plant at maturity (0.2781), secondary branches per plant at maturity (0.3566), fruits per cluster (0.3507), fruit width (cm) (0.3063), fruit length (cm) (0.2953), fruits yield per plant (0.2929) and seeds per fruit (0.0863), while it negative indirect effect on days to 50% flowering (-0.3268) and days to maturity (-0.0345).

4.7.1.6 Fruits per cluster

Fruits per cluster (0.8128) revealed that high negative direct effect on fruit dry weight per plant (g) (-0.1212) while, it expressed positive indirect correlation plant height (cm) at maturity (0.8104), primary branches per plant at maturity (0.8155), secondary branches per plant at maturity (0.8287), fruit width (cm) (0.8581), fruit length (cm) (0.8291), fruits yield

per plant (0.8337) and seeds per fruit (0.1536), while it negative indirect effect on days to 50% flowering (-0.8828) and days to first fruit initiation (-1.7358) and days to maturity (-0.0117).

4.7.1.7 Fruit width (cm)

Fruit width (cm) (0.0043) revealed that high negative direct effect on fruit dry weight per plant (g) (-0.1305) while, it expressed positive indirect correlation plant height (cm) at maturity (0.0049), primary branches per plant at maturity (0.0044), secondary branches per plant at maturity (0.0049), fruits per cluster (0.0046), fruit length (cm) (0.0048), fruits yield per plant (0.0050), days to maturity (0.0004) and seeds per fruit (0.0004), while it negative indirect effect on days to 50% flowering (-0.0050) and days to first fruit initiation (-0.0081).

4.7.1.8 Fruit length (cm)

Fruit length (cm) (-0.3004) revealed that high positive direct effect on fruit dry weight per plant (g) (0.0206) while, it expressed positive indirect correlation days to 50% flowering (0.2973) and days to first fruit initiation (0.5402), while it negative indirect effect on plant height (cm) at maturity (-0.2888), primary branches per plant at maturity (-0.2894), secondary branches per plant at maturity (-0.2994), fruits per cluster (-0.3064), fruit width (cm) (-0.3307), fruit length (cm) (-0.3004), fruits yield per plant (-0.2942), days to maturity (-0.0309) and seeds per fruit (-0.0882).

4.7.1.9 Fruits per plant

Fruits per plant (-1.6500) revealed that high positive direct effect on fruit dry weight per plant (g) (0.0290) while, it expressed positive indirect correlation days to 50% flowering (1.6604) and days to first fruit initiation (2.9428), while it negative indirect effect on plant height (cm) at maturity (-1.6345), primary branches per plant at maturity (-1.5999), secondary branches per plant at maturity (-1.6750), fruits per cluster (-1.6924), fruit width (cm) (-1.8951), fruit length (cm) (-1.6165), days to maturity (-0.1969) and seeds per fruit (-0.3747).

4.7.1.10 Days to maturity

Days to maturity (1.1234) revealed that high positive direct effect on fruit dry weight per plant (g) (1.0264) while, it expressed positive indirect correlation plant height (cm) at maturity (0.2246), secondary branches per plant at maturity (0.1138), days to first fruit initiation (0.2359), fruit width (cm) (0.1016), fruit length (cm) (0.1157), fruits yield per plant (0.1341), while it negative indirect effect on primary branches per plant at maturity (-0.0424), days to 50% flowering (-0.0774), fruits per cluster (-0.0162) and seeds per fruit (-0.0509).

4.2.1.11 Seeds per fruit

Seeds per fruit (0.1500) revealed that high positive direct effect on fruit dry weight per plant (g) (0.0155) while, it expressed positive indirect correlation plant height (cm) at maturity (0.0272), primary branches per plant at maturity (0.0279), secondary branches per plant at maturity (0.0281), fruits yield per plant (0.0284), fruit width (cm) (0.0133), fruit length (cm) (0.0440), fruits per plant (0.0341), while it negative indirect effect on days to 50% flowering (-0.0362), days to first fruit initiation (-0.0788) and days to maturity (-0.0068).

4.7 Phenotypic path coefficient

4.7.1 Direct and Indirect effects

4.7.1.1 Plant height (cm):

Plant height (-0.0044) revealed that positive direct effect on fruit dry weight per plant (g) (0.1458) while, it expressed positive indirect correlation days of 50% flowering (0.0039) and days to first fruit initiation (0.0030), while it negative indirect effect on primary branches per plant at maturity (-0.0039), secondary branches per plant at maturity (-0.0037), fruits per cluster (-0.0038), fruit width (cm) (-0.0034), fruit length (cm) (-0.0041), fruits per plant (-0.0043), days to maturity (-0.0010) and seeds per fruit (-0.0008).

4.7.1.2 Number of primary branches per plant

Primary branches per plant (0.0086) revealed that high positive direct effect on fruit dry weight per plant (g) (0.0663) while, it expressed negative indirect correlation days to 50% flowering (-0.0070), days to first fruit initiation (-0.0056), while it positive indirect effect on plant height (cm) at maturity (0.0077), secondary branches per plant at maturity (0.0072), fruits per cluster (0.0076), fruit width (cm) (0.0060), fruit length (cm) (0.0077), fruits per plant (0.0078), days to maturity (0.0008) and seed per fruit (0.0017).

4.7.1.3 Number of secondary branches per plant

Secondary branches per plant (-0.1490) revealed that high negative direct effect on fruit dry weight per plant (g) (-0.1332) while, it expressed positive indirect correlation days to 50% flowering (0.1333), days to first fruit initiation (0.0557) and days to maturity (0.0013), while it negative indirect effect on plant height (cm) at maturity (-0.1249), primary branches per plant at maturity (-0.1246), fruits per cluster (-0.1304), fruit width (cm) (-0.0955), fruit length (cm) (-0.1328), fruits per plant (-0.1333) and seeds per fruit (-0.0312).

4.7.1.5 Days to 50% flowering

Days to 50% flowering (-0.0522) revealed that high positive direct effect on fruit dry weight per plant (g) (0.1453) while, it expressed positive indirect correlation plant height (cm) at maturity (0.0454), primary branches per plant at maturity (0.0422), secondary branches per plant at maturity (0.0466), fruits per cluster (0.0426), fruit width (cm) (0.0350), fruit length (cm) (0.0474), fruits per plant (0.0485) and seeds per fruit (0.0100), while it negative indirect effect on days to first fruit initiation (-0.0288) and days to maturity (0.0033).

4.7.1.7 Days to first fruit initiation

Days to first fruit initiation (-0.3229) revealed that high negative direct effect on fruit dry weight per plant (g) (-0.2576) while, it expressed positive indirect correlation plant height (cm) at maturity (0.2196), primary

branches per plant at maturity (0.2086), secondary branches per plant at maturity (0.1208), fruits per cluster (0.1417), fruit width (cm) (0.1772), fruit length (cm) (0.1938), fruits per plant (0.2083), days to maturity (0.0218), while it negative indirect effect on days to 50% flowering (-0.1785).

4.7.1.8 Fruits per cluster

Fruits per cluster (0.4140) revealed that high positive direct effect on fruit dry weight per plant (g) (0.0733) while, it expressed positive indirect correlation plant height (cm) at maturity (0.3546), primary branches per plant at maturity (0.3630), secondary branches per plant at maturity (0.3622), fruit width (cm) (0.3150), fruit length (cm) (0.3634), fruits per plant (0.3638), days to maturity (0.0428) and seeds per fruit (0.1143), while it negative indirect effect on days to 50% flowering (-0.3384) and days to first fruit initiation (-0.1816).

4.7.1.9 Fruit width (cm)

Fruit width (cm) (0.0404) revealed that high positive direct effect on fruit dry weight per plant (g) (0.2836) while, it expressed positive indirect correlation plant height (cm) at maturity (0.0309), primary branches per plant at maturity (0.0279), secondary branches per plant at maturity (0.0259), fruits per cluster (0.0308), fruit length (cm) (0.0282), fruits per plant (0.0306), days to maturity (0.0113) and seeds per fruit (0.0070), while it negative indirect effect on days to 50% flowering (-0.0271) and days to first fruit initiation (-0.0222).

4.7.1.10 Fruit length (cm)

Fruit length (cm) (-0.2607) revealed that high negative direct effect on fruit dry weight per plant (g) (-0.0300) while, it expressed positive indirect correlation days to 50% flowering (0.2371) and days to first fruit initiation (0.1564), while it negative indirect effect on plant height (cm) at maturity (-0.2422), primary branches per plant at maturity (-0.2325), secondary branches per plant at maturity (-0.2323), fruits per cluster (-0.2289), fruit width (cm) (-0.1817), fruits per plant (-0.2535), days to maturity (-0.0155) and seeds per fruit (-0.0714).

4.7.1.11 Fruits per plant

Fruits per plant (-0.3580) revealed that high positive direct effect on fruit dry weight per plant (g) (0.0102) while, it expressed positive indirect correlation days to 50% flowering (0.3330) and days to first fruit initiation (0.2309), while it negative indirect effect on plant height (cm) at maturity (-0.3453), primary branches per plant at maturity (-0.3234), secondary branches per plant at maturity (-0.3202), fruits per cluster (-0.3147), fruit width (cm) (-0.2709), fruit length (cm) (-0.3481), days to maturity (-0.0336) and seeds per fruit (-0.0743).

4.7.1.12 Days to maturity

Days to maturity (0.8880) revealed that high positive direct effect on fruit dry weight per plant (g) (0.9425) while, it expressed positive indirect correlation plant height (cm) at maturity (0.1905), primary branches per plant at maturity (0.0846), days to 50% flowering (0.0568), fruits per cluster (0.0918), fruit width (cm) (0.2473), fruit length (cm) (0.0529), fruits per plant (0.0834) and seeds per fruit (0.0204), while it negative indirect effect on secondary branches per plant at maturity (-0.0077) and days to first fruit initiation (-0.1370).

4.7.1.13 Seeds per fruit

Seeds per fruit (0.0816) revealed that high positive direct effect on fruit dry weight per plant (g) (0.0791) while, it expressed positive indirect correlation plant height (cm) at maturity (0.0139), primary branches per plant at maturity (0.0157), secondary branches per plant at maturity (0.0171), fruits per cluster (0.0225), fruit width (cm) (0.0141), fruit length (cm) (0.0224), fruits per plant (0.0169) and days to maturity (0.0019), while it negative indirect effect on days to 50% flowering (-0.0156) and days to first fruit initiation (-0.0055).

Table 4.10 : Genotypic path coefficient of fruit yield and its component characters of chilli hybrids.

| Character | Plant Height (cm) Maturity | Primary Branches per Plant Maturity | Secondary Branches per Plant Maturity | Days of 50% Flowering | Days to First Fruit Initiation | Fruits per Cluster | Fruit Width (cm) | Fruit Length (cm) | Fruits per Plant | Days to Maturity | Seeds per Fruit |
|---------------------------------------|----------------------------|-------------------------------------|---------------------------------------|-----------------------|--------------------------------|--------------------|------------------|-------------------|------------------|------------------|-----------------|
| Plant Height (cm) Maturity | 1.0321 | 0.9739 | 1.0431 | -1.0306 | -1.7122 | 1.0291 | 1.1737 | 0.9924 | 1.0223 | 0.2064 | 0.1870 |
| Primary Branches per Plant Maturity | -0.1515 | -0.1606 | -0.1740 | 0.1650 | 0.2719 | -0.1611 | -0.1612 | -0.1547 | -0.1557 | 0.0061 | -0.0299 |
| Secondary Branches per Plant Maturity | -0.5013 | -0.5377 | -0.4960 | 0.4973 | 1.0771 | -0.5058 | -0.5569 | -0.4944 | -0.5035 | -0.0503 | -0.0928 |
| Days of 50% Flowering | 0.3081 | 0.3170 | 0.3093 | -0.3085 | -0.6141 | 0.3351 | 0.3561 | 0.3054 | 0.3105 | 0.0213 | 0.0745 |
| Days to First Fruit Initiation | 0.2724 | 0.2781 | 0.3566 | -0.3268 | -0.1642 | 0.3507 | 0.3063 | 0.2953 | 0.2929 | -0.0345 | 0.0863 |
| Fruits per Cluster | 0.8104 | 0.8155 | 0.8287 | -0.8828 | -1.7358 | 0.8128 | 0.8581 | 0.8291 | 0.8337 | -0.0117 | 0.1536 |
| Fruit Width (cm) | 0.0049 | 0.0044 | 0.0049 | -0.0050 | -0.0081 | 0.0046 | 0.0043 | 0.0048 | 0.0050 | 0.0004 | 0.0004 |
| Fruit Length (cm) | -0.2888 | -0.2894 | -0.2994 | 0.2973 | 0.5402 | -0.3064 | -0.3307 | -0.3004 | -0.2942 | -0.0309 | -0.0882 |
| Fruits per Plant | -1.6345 | -1.5999 | -1.6750 | 1.6604 | 2.9428 | -1.6924 | -1.8951 | -1.6165 | -1.6500 | -0.1969 | -0.3747 |
| Days to Maturity | 0.2246 | -0.0424 | 0.1138 | -0.0774 | 0.2359 | -0.0162 | 0.1016 | 0.1157 | 0.1341 | 1.1234 | -0.0509 |
| Seeds per Fruit | 0.0272 | 0.0279 | 0.0281 | -0.0362 | -0.0788 | 0.0284 | 0.0133 | 0.0440 | 0.0341 | -0.0068 | 0.1500 |
| Fruit Dry Weight per Plant (g) | 0.1036 | -0.2132 | 0.0399 | -0.0474 | 0.7547 | -0.1212 | -0.1305 | 0.0206 | 0.0290 | 1.0264 | 0.0155 |
| Partial R ² | 0.1069 | 0.0342 | -0.0198 | 0.0146 | -0.1239 | -0.0985 | -0.0006 | -0.0062 | -0.0478 | 1.1530 | 0.0023 |

R SQUARE = 1.0142 RESIDUAL EFFECT =SQRT(1- 1.0142)

Table 4.11: Categorization of chilli hybrids based on quality parameters

| Genotypes | Shape of fruits | Size of fruits | Flowers colour | Fruits colour | Breeding habit | Plant type |
|------------------------|------------------------|-----------------------|-----------------------|----------------------|-----------------------|-------------------|
| Smriti | Straight | Long | White | Dark green | Pendent | Bushy |
| Aujasvi | Straight | Medium | White | Green | Pendent | Bushy |
| Shourya | Straight | Medium | White | Dark green | Pendent | Bushy |
| Albeli-333 | Straight | Medium | White | Dark green | Pendent | Bushy |
| Ujjala | Straight | Long | White | Green | Pendent | Tall |
| Tiger | Straight | Long | White | Dark green | Pendent | Bushy |
| ARCH 930 | Straight | Long | White | Light green | Pendent | Bushy |
| US 612 | Straight | Long | White | Light green | Pendent | Tall |
| US 611Tara | Straight | Medium | White | Dark green | Pendent | Tall |
| Marshal 1857 | Straight | Medium | White | Dark green | Pendent | Bushy |
| Saniya (03) | Straight | Medium | White | Light green | Pendent | Bushy |
| Jalsa | Straight | Medium | White | Light green | Pendent | Bushy |
| S-586 | Straight | Medium | White | Dark green | Pendent | Bushy |
| Amrita | Straight | Long | White | Green | Pendent | Tall |
| P. jyoti | Straight | Long | White | Green | Pendent | Bushy |
| Jwala garima-12 | Straight | Medium | White | Dark green | Upright | Tall |
| TS-0017 | Straight | Medium | White | Green | Pendent | Bushy |
| Tara | Straight | Medium | White | Dark green | Pendent | Bushy |

CHAPTER - V

DISCUSSION

The results obtained from the present investigation entitle “Assessment of Morphological, Fruit Yield & Quality Characters in Hybrid Chilli (*Capsicum annum* L.)” are presented under the following heads:

In any crop improvement programme the success of selection depends on genetic variability and correlation among the characters in the population. The genetic variability estimated in terms of genotypic coefficient of variation and phenotypic coefficients of variation only is not adequate for the estimation of heritable variation. The heritability values, in broad sense are also helpful in selection on the basis of phenotypic performance of the quantitative characters. However, heritability estimates alone are not of any use in predicting the results of selection unless the correlation studies which indicate degree of relationship between yield and its components is done. Correlation studies provide information on nature, extent and direction of selection.

The findings obtained under the study has been analyzed and elaborated in the light of published research work as follows.

4.1 Analysis of variance

The analysis of variances for all the characters studied has been presented in Appendix – I to III. Mean squares due to genotypes were highly significantly for all the characters, indicating that the presence of genetic diversity in the existing material.

4.1.1 Mean and range performance of the genotypes

Range and mean performance (Minimum and Maximum) of the ten genotypes of chilli for all the thirty three characters are presented in the table 4.1a, b and c.

Hybrid Amrita was observed the tall plant and hybrid S-586 was recorded the dwarf plant, Maximum number of primary branches per plant was observed in hybrid Amrita, while it as recorded minimum in hybrid (S-586), Maximum number of secondary branches per plant was observed in hybrid Amrita, while it as recorded minimum in hybrid (S-586), Early first flower initiation was recorded in Amrita and hybrid S-586 exhibited in late

initiation of first flowering. Hybrid Amrita was recorded the early days to 50% flowering, therefore, hybrid S-586 exhibited in late group which is maximum days to 50% flowering days. Hybrid Amrita was recorded the early first picking in 71.19 days, therefore, hybrid S-586 exhibited in late picking of fruits 79.19 days. Hybrid Amrita was recorded the early fruit initiation in 67.52 days, therefore, hybrid S-586 exhibited in late group which fruit initiation in 75.52 days. The maximum fruits per cluster were recorded in hybrid Amrita, while it was found minimum in hybrid S-586. Hybrid Amrita showed maximum fruit width and it was found minimum in hybrid S-586. Hybrid Amrita showed maximum fruit length (7.69 cm) and it was found minimum in hybrid S-586 (3.37 cm). The maximum 146.00 fruits per plant were recorded in hybrid Amrita, while it was found minimum (86.00 fruit) in hybrid S-586. Early maturity was obtained under the hybrid Smriti and late in the hybrid Tara. Hybrid Tara was recorded maximum fresh fruit yield per plant followed by TS-0017 and Jwala garima-12. Whereas, it was recorded minimum in Smriti. Highest dry fruit yield per plot were recorded in hybrid Tara, TS-0017, Jwala garima-12 and P. jyoti, respectively and which were at par with each other. Highest dry fruit yield were recorded in genotype Tara, TS-0017, Jwala garima-12 and P. jyoti, respectively and which were at par with each other. However lowest dry fruit yield per ha exhibited in Smriti. Tinger was observed the maximum seeds per fruit, while it was noted minimum in hybrid Marshal1857 and Shourya. Hybrid Tinger was recorded maximum 1000 seed weight, while was showed minimum in hybrid Shourya.

The findings were quite similar to as reported by Dutonde *et al.* (2006), Tembhone *et al.* (2008), Jabeen *et al.* (2012) for fresh fruit yield per ha, Dutonde *et al.* (2006), Tembhone *et al.* (2008) for number of fruits per plant, Dutonde *et al.* (2006), Tembhone *et al.* (2008), Jabeen *et al.* (2012) for fresh fruit yield per plant.

5.2 Variability studies:

In the present findings PCV were observed higher than the corresponding GCV for all the characters studied, however the differences was narrow which implied their relative resistance to environmental variation. It also described that genetic factors were predominantly responsible for expression of those attributes and selection could be made effectively on the

basis of phenotypic performance. The finding of Datta and Jana (2010), Diwaker *et al.* (2012) is similar to that of the present findings.

The highest GCV value was recorded for Fruits width cm (23.99 %), while moderate for fruit length cm (23.10 %), fruits per cluster (21.54 %), secondary branches per plant at 60 DAT (20.30 %), primary branches per plant 90 DAT (18.90 %), primary branches per plant maturity (16.83 %), fruits per plant (15.45 %), secondary branches per plant 90 DAT (14.98 %), fruit fresh weight plant (g), fruit fresh weight per plot (kg), fruit fresh weight per ha. (q.), fruit dry weight per plant (g), fruit dry weight per plot (kg), fruit dry weight per ha. (q.) (14.37 %), seeds per fruit, 1000 seed weight (g) (14.16 %), primary branches per plant 60 DAT (13.46 %), secondary branches per plant 120 DAT (13.11 %), plant height (cm) 120 DAT (12.76 %), days to first flower initiation (12.26 %), plant height (cm) maturity (11.90 %), plant height (cm) 30 DAT (11.74 %), plant height (cm) 90 DAT (11.09 %), secondary branches per plant maturity (11.32), primary branches per plant 30 DAT (10.42), days to maturity (3.68 %), days of 50% flowering (3.16 %), days to first picking of fruits (2.78 %) and days to first fruit initiation (1.64 %).

The highest PCV value was recorded for fruits width cm (36.00 %), while moderate for fruits per cluster (24.59 %), fruit fresh weight per plot (kg), fruit dry weight per plot (kg) (23.60 %), fruit fresh weight plant (g), fruit fresh weight per ha. (q.), fruit dry weight per plant (g), fruit dry weight per ha. (q.) (23.59 %), fruit length cm (23.22 %), while low value for PCV was recorded for Days to First Fruit Initiation (4.69 %), Days to maturity (4.32 %), Days to 50 % flowering (3.46 %) and Days to first fruit picking (3.05 %). The high values of GCV suggested greater phenotypic and genotypic variability among the genotypes and responsiveness of the attributes for making further improvement by selection. The findings are in close harmony with the result of Mohammed *et al.* (2001) for branches per plant, Rathod *et al.* (2002), Manju and Sreelathakumary (2002a), Mini and Khader (2004), Sreelathakumary and Rajamony (2004), Prabhakaran *et al.* (2004), Varkey *et al.* (2005), Bharadwaj *et al.* (2007), Vani *et al.* (2007), Samadia (2007), Yadwad *et al.* (2008), Gupta *et al.* (2009), Sarkar *et al.* (2009), Sharma *et al.* (2009), Datta and Jana (2010) for number of fruits per plant, Ukkund *et al.* (2006) for number of secondary branches per plant.

The highest ECV value was recorded for fruits width cm (26.85 %), while moderate for fruit fresh weight per plot (kg), fruit dry weight per plot (kg) (18.72 %), fruit fresh weight per plant (g), fruit fresh weight per ha. (q.), fruit dry weight per plant (g), fruit dry weight per ha (q) (18.71 %). Low value for ECV was recorded for plant height (cm) at 60 DAT (2.83 %), plant height (cm) at 90 DAT (2.56 %), days to maturity (2.26 %), days to 50 % flowering (1.41 %), days to first fruit picking (1.24 %) and fruits per plant (1.21 %).

5.3 Heritability

Heritability which denotes the proportion of genetically controlled variability expressed by a programme for a particular character or a set of character is very important biometrical tool for guiding plant breeders for adoption of appropriate breeding procedures. High heritability in broad sense is helpful in identifying appropriate character for selection and enables the breeder to select superior genotypes on the basis of phenotypic expression of quantitative characters. The estimated values of heritability in broad sense were classified as very high (more than 80%), medium (50-70%) and low (less than 50%).

Heritability estimates in broad sense were high for plant height (cm) 30 DAT (86 %), plant height (cm) 60 DAT (97 %), plant height (cm) 90 DAT (95 %), plant height (cm) 120 DAT (92 %), plant height (cm) maturity (95 %), primary branches per plant 90 DAT (87 %), primary branches per plant 120 DAT (88 %), primary branches per plant maturity (88 %), secondary branches per plant 30 DAT (85 %), secondary branches per plant 60 DAT (90 %), secondary branches per plant 90 DAT (95 %), secondary branches per plant 120 DAT (92 %), days to first picking of fruits (83.5), fruit length (cm) (99 %), fruits yield per plant (99 %), seeds per fruit (92 %), 1000 seeds weight (g) (92 %). The results were in close proximity to that of Mohammed *et al.* (2001), Gogoi and Gautam (2002), Ukkund *et al.* (2006) for plant height, Mohammed *et al.* (2001), Gogoi and Gautam (2002), Rathod *et al.* (2002), Ukkund *et al.* (2006) for fruit length, Mohammed *et al.* (2001), Gogoi and Gautam (2002), Rathod *et al.* (2002), Ukkund *et al.* (2006), Bharadwaj *et al.* (2007) for number of fruits per plant, Rathod *et al.* (2002) for 1000 seed weight, Rathod *et al.* (2002) for number of primary branches per plant, Rathod

et al. (2002), Ukkund *et al.* (2006) for fresh fruit yield, Rathod *et al.* (2002) for dry fruit yield. High values of broad sense heritability for these characters expressed that they were least influenced by environmental modification. It reflected that the phenotypes were the true representative of their genotypes and selection based on phenotypic performance would be reliable.

Heritability estimates in broad sense were medium for primary branches per plant 60 DAT (51 %), secondary branches per plant maturity (79 %), days to first flower initiation (78 %), fruits per cluster (77 %), days to maturity (73 %). The findings were in agreement to Gogoi and Gautam (2002).

Heritability estimates in broad sense were low for primary branches per plant 30 DAT (44 %), fruit width (44 %), days to maturity, fruit fresh weight plant (g), fruit fresh weight per plot (kg), fruit fresh weight per ha. (q.), fruit dry weight per plant (g), fruit dry weight per plot (kg), fruit dry weight per ha. (q.) (37 %). Heritability however indicates only the effectiveness with which selection of a genotype can be based on phenotypic performance, but fails to indicate the genetic progress. Heritability estimates along with genetic gains are more effective and reliable in predicting the improvement through selection (Johnson *et al.*, 1955). Estimates of genetic advance helps to predict the extent of improvement that can be achieved for improving the different characters. The estimated values of genetic advance as percent of mean were classified as high (more than 40%), moderate (20-35%) and low (less than 20%).

5.4 Genetic advance

High value of genetic advance was recorded for fruit fresh weight plant (g) (90.39) followed by fruit fresh weight per ha. (q) (36.16), fruit per plant (34.40), fruit dry weight per plant (g) (30.13), plant height (cm) 120 DAT (19.29), plant height (cm) maturity (18.42), plant height (cm) 90 DAT (15.83) and plant height (cm) 60 DAT (15.47). The results were in consonance with Mohammed *et al.* (2001), Jabeen *et al.* (2012) for branches per plant, Mohammed *et al.* (2001), Gogoi and Gautam (2002), Varkey *et al.* (2005), Sharma *et al.* (2009), Singh and Singh (2012) for dry fruit yield, Gogoi and Gautam (2002), Rathod *et al.* (2002), Manju and Sreelathakumary (2002a), Mini and Khader (2004), Prabhakaran *et al.* (2004), Dutonde *et al.*

(2006), Kaur and Singh (2007), Gupta *et al.* (2009), Sarkar *et al.* (2009), Diwaker *et al.* (2012), Singh and Singh (2012), Jabeen *et al.* (2012) for fresh fruit yield per plant, Gogoi and Gautam (2002), Rathod *et al.* (2002), Manju and Sreelathakumary (2002a), Mini and Khader (2004), Varkey *et al.* (2005), Bharadwaj *et al.* (2007), Vani *et al.* (2007), Kaur and Singh (2007), Tembhurne *et al.* (2008), Gupta *et al.* (2009), Sarkar *et al.* (2009), Datta and Jana (2010), Diwaker *et al.* (2012), Jabeen *et al.* (2012) for number of fruits plant⁻¹, Gogoi and Gautam (2002), Mini and Khader (2004), Dutonde *et al.* (2006), Vani *et al.* (2007), Gupta *et al.* (2009), Sarkar *et al.* (2009), Singh and Singh (2012), Jabeen *et al.* (2012) for fruit length, Manju and Sreelathakumary (2002a), Varkey *et al.* (2005), Vani *et al.* (2007), Kaur and Singh (2007), Singh and Singh (2012), Jabeen *et al.* (2012) for number of seeds per fruit, Mini and Khader (2004) for secondary branches per plant, Jabeen *et al.* (2012) for fresh fruit yield ha.

Genetic advance was computed as percentage as mean. It has been classified as high (more than 40), medium (25 to 40) and low (less than 25). Medium value of genetic advance as percentage of mean was recorded for plant height (cm) 60 DAT (31.13 %), plant height (cm) 120 DAT (25.27 %), primary branches per plant 90 DAT (36.34 %), primary branches per plant 120 DAT (32.51 %), primary branches per plant maturity (32.51 %), secondary branches per plant 30 DAT (34.68 %), secondary branches per plant 60 DAT (39.73 %), secondary branches per plant 90 DAT (30.05 %), secondary branches per plant 120 DAT (25.85 %), fruits per cluster (38.88 %), fruit width (cm) (32.92 %), fruit length (cm) (47.34 %), fruits per plant (31.73 %), seeds per fruit (27.96 %) and 1000 seed weight (g) (27.97 %). The results were in consonance with Ukkund *et al.* (2006).

Low value of genetic advance as percentage of mean was recorded for plant height (cm) 30 DAT (22.38 %), plant height (cm) 90 DAT (22.25 %), plant height (cm) maturity (23.93 %), primary branches per plant 30 DAT (14.27 %), primary branches per plant 60 DAT (19.72 %), secondary branches per plant maturity (20.76 %), days to first flower initiation (22.35 %), days to first picking of fruits (5.24 %), days of 50% flowering (5.95 %), days to first fruit initiation (1.18 %), days to maturity (6.45 %), fruit fresh weight plant (g) (18.03 %), fruit fresh weight per plot (kg) (18.02 %), fruit fresh weight per

ha. (q.) (18.03 %), fruit dry weight per plant (g) (18.03 %), fruit dry weight per plot (kg) (18.02 %) and fruit dry weight per ha. (q.) (18.03 %).

5.5 Association studies

Correlation coefficient analysis

A wide range of variation in quantitative characters provides the basis for selection in plant breeding programme. The knowledge of association among the characters is useful to the breeder for improving the efficiency of selection. Correlation coefficient analysis measures the mutual relationship between plant characters and determines the component character on which selection can be made for genetic improvement of yield. Investigation regarding the presence of component and nature of association among themselves is essential and pre-requisite for improvement in yield. Correlation coefficient provides a clear picture of the extent of association between a pair of traits and indicates whether simultaneous improvement of the correlated traits may be possible or not. The knowledge of genetic association between yield and its component characters help in improving the efficiency of selection for yield by making proper choice and balancing one component with another.

The magnitude of genotypic correlation was higher than the phenotypic correlation for all the traits that indicated inherent association between various characters. The findings were in agreement to Singh and Singh (2012).

Plant height expressed positive correlation with primary branches per plant maturity (0.9436), secondary branches per plant maturity (1.0106), fruits per cluster (0.9971), fruit width (cm) (1.1327), fruit length (cm) (0.9616), fruits per plant (0.9906), days to maturity (0.2000) and seeds per fruit (0.1812). Negative association of this character was recorded with days of 50% flowering (-0.9985) and days to first fruit initiation (-1.6590). Smitha and Basavaraja (2006) for fresh fruits yield per plant and Padhar and Zaveri (2010) for dry fruits yield per plant.

Correlation coefficient of number of primary branches per plant at maturity was observed positive with number of secondary branches per plant at maturity (1.0839), fruits per cluster (1.0033), fruit width (cm) (1.0036), fruit length (cm) (0.9634), fruits per plant (0.9696) and seeds per fruit (0.1862). Negative association of this character was recorded with days of 50%

flowering (-1.0275), days to first fruit initiation (-1.6935) and days to maturity (-0.0377).

Correlation coefficient of number of secondary branches per plant at maturity was observed), fruits per cluster (1.0196), fruit width (cm) (1.1226), fruit length (cm) (0.9968), fruits per plant (1.0151), days to maturity (0.1013) and seedsperfruit (0.1871). Negative association of this character was recorded with days of 50% flowering (-1.0025), days to first fruit initiation (-2.1715)

Association of days to 50% flowering was exhibited positive with days to 1st first fruit initiation (1.9904). Negative association of this character was recorded with fruitper clusters (-1.0861), fruit width (cm) (-1.1540), fruit length (cm) (-0.9898), fruitsperplant (-1.0063), days to maturity (-0.0689) and seeds per fruit (-0.2415).

Correlation coefficient of days to 1st fruit initiation was found positive and significant with days to maturity (0.2100). Significant and negative association of this character was recorded with fruit per clusters (-2.1356), fruit width (cm) (-1.8653), fruit length (cm) (-1.7986), fruits per plant (-1.7835) and seeds per fruit (-0.5256). These findings corroborated the earlier findings of Diwaker *et al.* (2012) for number of fruits per plant.

Correlation coefficient of fruits per cluster was found positive and significant with fruit width (cm) (1.0558), fruit length (cm) (1.0200), fruits per plant (1.0257) and seeds per fruit (0.1890). Significant and negative association of this character was recorded with days to maturity (-0.0144).

Fruit width expressed a significant and positive association with fruit length (cm) (1.1012), fruits per plant (1.1485), days to maturity (0.0904) and seed per fruit (0.0889).

Fruit length expressed a significant and positive association with fruits per plant (0.9797), days to maturity (0.1030) and seeds per fruit (0.2930). These results are in close harmony with the findings of Mohammed *et al.* (2001) for dry fruits yield per plant, Smitha and Basavaraja (2006), Vani *et al.* (2007), Gupta *et al.* (2009), Singh *et al.* (2009), Patel *et al.* (2009), Diwaker *et al.* (2012) and Singh and Singh (2012) for fresh fruit yield plant⁻¹,

Fruits per plant expressed a significant and positive association with days to maturity (0.1193) and seeds per fruit (0.2271). These results are in

close harmony with the findings of Mohammed *et al.* (2001) and Padhar and Zaveri (2010) for dry fruits yield per plant, Nehru *et al.* (2003), Nandadevi and Hosamani (2003), Smitha and Basavaraja (2006), Bharadwaj *et al.* (2007), Bharadwaj *et al.* (2007a), Tembhurne *et al.* (2008), Gupta *et al.* (2009), Sarkar *et al.* (2009), Singh *et al.* (2009), Patel *et al.* (2009), Padhar and Zaveri (2010), Diwaker *et al.* (2012), Singh and Singh (2012) and Chaudhary *et al.* (2013) for fresh fruits yield per plant.

Days to maturity expressed a negative association with and seeds per fruit (-0.0453).

Phenotypic Correlation Coefficient

Plant height expressed positive phenotypic correlation with primary branches per plant maturity (0.8860), secondary branches per plant maturity (0.8381), fruits per cluster (0.8567), fruit width (cm) (0.7645), fruit length (cm) (0.9291), fruits per plant (0.9644), days to maturity (0.2145) and seeds per fruit (0.1700). Negative association of this character was recorded with days of 50% flowering (-0.8712) and days to first fruit initiation (-0.6800)

Phenotypic correlation coefficient of number of primary branches per plant at maturity was observed positive with number of secondary branches per plant at maturity (0.8360), fruits per cluster (0.8769), fruit width (cm) (0.6912), fruit length (cm) (0.8916), fruits per plant (0.9032), days to maturity (0.0952) and seeds per fruit (0.1922). Negative association of this character was recorded with days of 50% flowering (-0.8097) and days to first fruit initiation (-0.6459).

Phenotypic correlation coefficient of number of secondary branches per plant at maturity was observed), fruits per cluster (0.8750), fruit width (cm) (0.6410), fruit length (cm) (0.8911), fruits per plant (0.8944) and seeds per fruit (0.2091). Negative association of this character was recorded with days of 50% flowering (-0.8942), days to first fruit initiation (-0.3741) and days to maturity (-0.0087).

Association of days to 50% flowering was exhibited positive with days to 1st first fruit initiation (0.5528) and days to maturity (0.0639). Negative association of this character was recorded with fruit per clusters (-0.8174), fruit width (cm) (-0.6701), fruit length (cm) (-0.9096), fruits per plant (-0.9300) and seeds per fruit (-0.1911).

Phenotypic correlation coefficient of negative association of this character was recorded with fruit per clusters (-0.4388), fruit width (cm) (-0.5489), fruit length (cm) (-0.6000), fruits per plant (-0.6450), days to maturity (-0.1543) and seeds per fruit (-0.0675).

Phenotypic correlation coefficient of fruits per cluster was found positive and significant with fruit width (cm) (0.7609), fruit length (cm) (0.8778), fruits per plant (0.8789), days to maturity (0.1034) and seeds per fruit (0.1034).

Fruit width expressed a phenotypic correlation positive association with fruit length (cm) (0.6969), fruits per plant (0.7566), days to maturity (0.2785) and seeds per fruit (0.1730).

Fruit length expressed a significant and positive association with fruits per plant (0.9724), days to maturity (0.0596) and seeds per fruit (0.2741).

Fruits per plant expressed a significant and positive association with days to maturity (0.0939) and seeds per fruit (0.2075).

Days to maturity expressed a positive association with and seeds per fruit (0.230).

5.4 Path coefficient

5.4.1 Genotypic path coefficient

Plant height (1.0321) revealed that positive direct effect on fruit dry weight per plant (g) (0.1036) while, it expressed positive indirect correlation primary branches per plant at maturity (0.9739), secondary branches per plant at maturity (1.0431), fruits per cluster (1.0291), fruit width (cm) (1.1737), fruit length (cm) (0.9924), fruits per plant (1.0223), days to maturity (0.2064) and seeds per fruit (0.1870), while it negative indirect effect on days of 50% flowering (-1.0306) and days to first fruit initiation (-1.7122).

Primary branches per plant (-0.1606) revealed that high negative direct effect on fruit dry weight per plant (g) (-0.2132) while, it expressed positive indirect correlation days to 50% flowering (0.1650), days to first fruit initiation (0.2719) and days to maturity (0.0061), while it negative indirect effect on plant height (cm) at maturity (-0.1515), primary branches per plant at maturity (-0.1606), secondary branches per plant at maturity (-0.1740), fruits per cluster (-0.1611), fruit width

(cm) (-0.1612), fruit length (cm) (-0.1547), fruits per plant (-0.1557) and seeds per fruit (-0.0299).

Secondary branches per plant (-0.4960) revealed that high positive direct effect on fruit dry weight per plant (g) (0.0399) while, it expressed positive indirect correlation days to 50% flowering (0.4973) and days to first fruit initiation (1.0771), while it negative indirect effect on plant height (cm) at maturity (-0.5013), primary branches per plant at maturity (-0.5377), secondary branches per plant at maturity (-0.4960), fruits per cluster (-0.5058), fruit width (cm) (-0.5569), fruit length (cm) (-0.4944), fruits per plant (-0.5035), days to maturity (-0.0503) and seeds per fruit (-0.0928).

Days to 50% flowering (-0.3085) revealed that high negative direct effect on fruit dry weight per plant (g) (-0.0474) while, it expressed positive indirect correlation plant height (cm) at maturity (0.3081), primary branches per plant at maturity (0.3170), secondary branches per plant at maturity (0.3093), fruits per cluster (0.3351), fruit width (cm) (0.3561), fruit length (cm) (0.3054), fruits per plant (0.3105), days to maturity (0.0213) and seeds per fruit (0.0745), while it negative indirect effect on days to first fruit initiation (-0.6141).

Days to first fruit initiation (-0.1642) revealed that high positive direct effect on fruit dry weight per plant (g) (0.7547) while, it expressed positive indirect correlation plant height (cm) at maturity (0.2724), primary branches per plant at maturity (0.2781), secondary branches per plant at maturity (0.3566), fruits per cluster (0.3507), fruit width (cm) (0.3063), fruit length (cm) (0.2953), fruits per plant (0.2929) and seeds per fruit (0.0863), while it negative indirect effect on days to 50% flowering (-0.3268) and days to maturity (-0.0345).

Fruits per cluster (0.8128) revealed that high negative direct effect on fruit dry weight per plant (g) (-0.1212) while, it expressed positive indirect correlation plant height (cm) at maturity (0.8104), primary branches per plant at maturity (0.8155), secondary branches per plant at maturity (0.8287), fruit width (cm) (0.8581), fruit length (cm) (0.8291), fruits per plant (0.8337) and seeds per fruit (0.1536), while it

negative indirect effect on days to 50% flowering (-0.8828) and days to first fruit initiation (-1.7358) and days to maturity (-0.0117).

Fruit width (cm) (0.0043) revealed that high negative direct effect on fruit dry weight per plant (g) (-0.1305) while, it expressed positive indirect correlation plant height (cm) at maturity (0.0049), primary branches per plant at maturity (0.0044), secondary branches per plant at maturity (0.0049), fruits per cluster (0.0046), fruit length (cm) (0.0048), fruits per plant (0.0050), days to maturity (0.0004) and seeds per fruit (0.0004), while it negative indirect effect on days to 50% flowering (-0.0050) and days to first fruit initiation (-0.0081).

Fruit length (cm) (-0.3004) revealed that high positive direct effect on fruit dry weight per plant (g) (0.0206) while, it expressed positive indirect correlation days to 50% flowering (0.2973) and days to first fruit initiation (0.5402), while it negative indirect effect on plant height (cm) at maturity (-0.2888), primary branches per plant at maturity (-0.2894), secondary branches per plant at maturity (-0.2994), fruits per cluster (-0.3064), fruit width (cm) (-0.3307), fruit length (cm) (-0.3004), fruits per plant (-0.2942), days to maturity (-0.0309) and seeds per fruit (-0.0882).

Fruits per plant (-1.6500) revealed that high positive direct effect on fruit dry weight per plant (g) (0.0290) while, it expressed positive indirect correlation days to 50% flowering (1.6604) and days to first fruit initiation (2.9428), while it negative indirect effect on plant height (cm) at maturity (-1.6345), primary branches per plant at maturity (-1.5999), secondary branches per plant at maturity (-1.6750), fruits per cluster (-1.6924), fruit width (cm) (-1.8951), fruit length (cm) (-1.6165), days to maturity (-0.1969) and seeds per fruit (-0.3747).

Days to maturity (1.1234) revealed that high positive direct effect on fruit dry weight per plant (g) (1.0264) while, it expressed positive indirect correlation plant height (cm) at maturity (0.2246), secondary branches per plant at maturity (0.1138), days to first fruit initiation (0.2359), fruit width (cm) (0.1016), fruit length (cm) (0.1157), fruits per plant (0.1341), while it negative indirect effect on primary branches per

plant at maturity (-0.0424), days to 50% flowering (-0.0774), fruits per cluster (-0.0162) and seeds per fruit (-0.0509).

Seeds per fruit (0.1500) revealed that high positive direct effect on fruit dry weight per plant (g) (0.0155) while, it expressed positive indirect correlation plant height (cm) at maturity (0.0272), primary branches per plant at maturity (0.0279), secondary branches per plant at maturity (0.0281), fruits per plant (0.0284), fruit width (cm) (0.0133), fruit length (cm) (0.0440), fruits per plant (0.0341), while it negative indirect effect on days to 50% flowering (-0.0362), days to first fruit initiation (-0.0788) and days to maturity (-0.0068).

5.4.2 Phenotypic path coefficient

Plant height (-0.0044) revealed that positive direct effect on fruit dry weight per plant (g) (0.1458) while, it expressed positive indirect correlation days of 50% flowering (0.0039) and days to first fruit initiation (0.0030), while it negative indirect effect on primary branches per plant at maturity (-0.0039), secondary branches per plant at maturity (-0.0037), fruits per cluster (-0.0038), fruit width (cm) (-0.0034), fruit length (cm) (-0.0041), fruits per plant (-0.0043), days to maturity (-0.0010) and seeds per fruit (-0.0008).

Primary branches per plant (0.0086) revealed that high positive direct effect on fruit dry weight per plant (g) (0.0663) while, it expressed negative indirect correlation days to 50% flowering (-0.0070), days to first fruit initiation (-0.0056), while it positive indirect effect on plant height (cm) at maturity (0.0077), secondary branches per plant at maturity (0.0072), fruits per cluster (0.0076), fruit width (cm) (0.0060), fruit length (cm) (0.0077), fruits per plant (0.0078), days to maturity (0.0008) and seeds per fruit (0.0017).

Secondary branches per plant (-0.1490) revealed that high negative direct effect on fruit dry weight per plant (g) (-0.1332) while, it expressed positive indirect correlation days to 50% flowering (0.1333), days to first fruit initiation (0.0557) and days to maturity (0.0013), while it negative indirect effect on plant height (cm) at maturity (-0.1249),

primary branches per plant at maturity (-0.1246), fruits per cluster (-0.1304), fruit width (cm) (-0.0955), fruit length (cm) (-0.1328), fruits per plant (-0.1333) and seeds per fruit (-0.0312).

Days to 50% flowering (-0.0522) revealed that high positive direct effect on fruit dry weight per plant (g) (0.1453) while, it expressed positive indirect correlation plant height (cm) at maturity (0.0454), primary branches per plant at maturity (0.0422), secondary branches per plant at maturity (0.0466), fruits per cluster (0.0426), fruit width (cm) (0.0350), fruit length (cm) (0.0474), fruits per plant (0.0485) and seeds per fruit (0.0100), while it negative indirect effect on days to first fruit initiation (-0.0288) and days to maturity (0.0033).

Days to first fruit initiation (-0.3229) revealed that high negative direct effect on fruit dry weight per plant (g) (-0.2576) while, it expressed positive indirect correlation plant height (cm) at maturity (0.2196), primary branches per plant at maturity (0.2086), secondary branches per plant at maturity (0.1208), fruits per cluster (0.1417), fruit width (cm) (0.1772), fruit length (cm) (0.1938), fruits per plant (0.2083), days to maturity (0.0218), while it negative indirect effect on days to 50% flowering (-0.1785).

Fruits per cluster (0.4140) revealed that high positive direct effect on fruit dry weight per plant (g) (0.0733) while, it expressed positive indirect correlation plant height (cm) at maturity (0.3546), primary branches per plant at maturity (0.3630), secondary branches per plant at maturity (0.3622), fruit width (cm) (0.3150), fruit length (cm) (0.3634), fruits per plant (0.3638), days to maturity (0.0428) and seeds per fruit (0.1143), while it negative indirect effect on days to 50% flowering (-0.3384) and days to first fruit initiation (-0.1816).

Fruit width (cm) (0.0404) revealed that high positive direct effect on fruit dry weight per plant (g) (0.2836) while, it expressed positive indirect correlation plant height (cm) at maturity (0.0309), primary branches per plant at maturity (0.0279), secondary branches per plant at maturity (0.0259), fruits per cluster (0.0308), fruit length (cm) (0.0282), fruits per plant (0.0306), days to maturity (0.0113) and seeds

per fruit (0.0070), while it negative indirect effect on days to 50% flowering (-0.0271) and days to first fruit initiation (-0.0222).

Fruit length (cm) (-0.2607) revealed that high negative direct effect on fruit dry weight per plant (g) (-0.0300) while, it expressed positive indirect correlation days to 50% flowering (0.2371) and days to first fruit initiation (0.1564), while it negative indirect effect on plant height (cm) at maturity (-0.2422), primary branches per plant at maturity (-0.2325), secondary branches per plant at maturity (-0.2323), fruits per cluster (-0.2289), fruit width (cm) (-0.1817), fruits per plant (-0.2535), days to maturity (-0.0155) and seeds per fruit (-0.0714).

Fruits per plant (-0.3580) revealed that high positive direct effect on fruit dry weight per plant (g) (0.0102) while, it expressed positive indirect correlation days to 50% flowering (0.3330) and days to first fruit initiation (0.2309), while it negative indirect effect on plant height (cm) at maturity (-0.3453), primary branches per plant at maturity (-0.3234), secondary branches per plant at maturity (-0.3202), fruits per cluster (-0.3147), fruit width (cm) (-0.2709), fruit length (cm) (-0.3481), days to maturity (-0.0336) and seeds per fruit (-0.0743).

Days to maturity (0.8880) revealed that high positive direct effect on fruit dry weight per plant (g) (0.9425) while, it expressed positive indirect correlation plant height (cm) at maturity (0.1905), primary branches per plant at maturity (0.0846), days to 50% flowering (0.0568), fruits per cluster (0.0918), fruit width (cm) (0.2473), fruit length (cm) (0.0529), fruits per plant (0.0834) and seeds per fruit (0.0204), while it negative indirect effect on secondary branches per plant at maturity (-0.0077) and days to first fruit initiation (-0.1370).

Seeds per fruit (0.0816) revealed that high positive direct effect on fruit dry weight per plant (g) (0.0791) while, it expressed positive indirect correlation plant height (cm) at maturity (0.0139), primary branches per plant at maturity (0.0157), secondary branches per plant at maturity (0.0171), fruits per cluster (0.0225), fruit width (cm) (0.0141), fruit length (cm) (0.0224), fruits per plant (0.0169) and days to maturity (0.0019), while it negative indirect effect on days to 50% flowering (-0.0156) and days to first fruit initiation (-0.0055).

CHAPTER - VI

SUMMARY, CONCLUSION AND SUGGESTIONS FOR FUTURE WORK

6.1 Summary

The present investigation entitled “**Assessment of Morphological, Fruit Yield & Quality Characters in Hybrid Chilli (*Capsicum annum* L.)**” was undertaken on eighteen hybrid of chilli collected from different sources of Madhya Pradesh. The experiment was laid in Randomized Block Design replicated thrice in *kharif* season during the year 2014-15 at Horticulture Research Field Of RAK College of Agriculture Sehore.

Different observations were recorded on Plant height (cm) at an equal interval 30, 60, 90,120 DAT and at maturity, No. of primary branches per plant at 30, 60, 90,120 DAT and at maturity, No. of secondary branches per plant at 30, 60, 90,120 DAT and at maturity, Days to first flower initiation, Days of 50% flowering, Days to first picking of fruits. Days to first fruit initiation, Number of fruits per cluster, Fruit width (cm), Fruit length (cm), Number of fruits per plant, Days to maturity, Fresh weight of fruit per plant (g), Fresh weight of fruit per plot (kg), Fresh weight of fruit per ha. (q.), Dry weight of fruit per plant (g), Dry fruit weight per plot (kg), Dry fruit weight per ha. (q.), Number of seeds per fruit and 1000 seed weight (g)

6.1 Variability studies:

The highest GCV value was recorded for Fruits width cm, while moderate for fruit length, fruits per cluster, secondary branches per plant at 60 DAT, primary branches per plant 90 DAT, primary branches per plant maturity, fruits per plant, secondary branches per plant 90 DAT, fruit fresh weight plant (g), fruit fresh weight per plot (kg), fruit fresh weight per ha. (q.), fruit dry weight per plant (g), fruit dry weight per plot (kg), fruit dry weight per ha. (q.), Seeds per Fruit, 1000 seeds weight (g), primary branches per plant 60 DAT, secondary branches per plant 120 DAT, plant height (cm) 120 DAT, days to first flower initiation, plant height (cm) maturity, plant height (cm) 30 DAT, plant height (cm) 90 DAT, secondary branches per plant maturity, primary branches per plant 30 DAT, days to maturity, days of 50% flowering, days to first picking of fruits and days to first fruit initiation. The

highest PCV value was recorded for fruits width, while moderate for fruits per cluster, fruit fresh weight per plot (kg), fruit dry weight per plot (kg), fruit fresh weight plant (g), fruit fresh weight per ha. (q.), fruit dry weight per plant (g), fruit dry weight per ha. (q.), fruit length, while low value for PCV was recorded for Days to First Fruit Initiation, Days to maturity, Days to 50 % flowering and Days to first fruit picking.

6.2 Heritability

Heritability estimates in broad sense were high for Plant Height (cm) 30 DAT, plant height (cm) 60 DAT (97 %), plant height (cm) 90 DAT, plant height (cm) 120 DAT, plant height (cm) maturity, primary branches per plant 90 DAT, primary branches per plant 120 DAT, primary branches per plant maturity, secondary branches per plant 30 DAT, secondary branches per plant 60 DAT, secondary branches per plant 90 DAT, secondary branches per plant 120 DAT, days to first picking of fruits, fruit length (cm), fruits per plant, seeds per fruit, 1000 seed weight.

Heritability estimates in broad sense were medium for primary branches per plant 60 DAT, secondary branches per plant maturity, days to first flower initiation, fruits per cluster, days to maturity.

Heritability estimates in broad sense were low for primary branches per plant 30 DAT, fruit width, days to maturity, fruit fresh weight plant (g), fruit fresh weight per plot (kg), fruit fresh weight per ha. (q.), fruit dry weight per plant (g), fruit dry weight per plot (kg), fruit dry weight per ha. (q.).

6.3 Genetic advance

High value of genetic advance was recorded for fruit fresh weight plant (g) followed by fruit fresh weight per ha. (q), Fruit yield per plant, fruit dry weight per plant (g), plant height (cm) 120 DAT, plant height (cm) maturity, plant height (cm) 90 DAT and plant height (cm) 60 DAT.

Medium value of genetic advance as percentage of mean was recorded for plant height (cm) 60 DAT, plant height (cm) 120 DAT, primary branches per plant 90 DAT, primary branches per plant 120 DAT, primary branches per plant maturity, secondary branches per plant 30 DAT, secondary branches per plant 60 DAT, secondary branches per plant 90 DAT, secondary branches per plant 120 DAT, fruits per cluster, fruit width (cm), fruit length (cm), fruits per plant, seeds per fruit and 1000 Seed Weight (g).

6.4 Phenotypic Correlation Coefficient

Plant height expressed positive phenotypic correlation with primary branches per plant maturity, secondary branches per plant maturity, fruits per cluster, fruit width (cm), fruit length (cm), fruits per plant, days to maturity and seeds per fruit. Negative association of this character was recorded with days of 50% flowering and days to first fruit initiation.

Primary branches per plant at maturity was observed positive with number of secondary branches per plant at maturity, fruits per cluster, fruit width, fruit length, fruits per plant, days to maturity and seeds per fruit. Negative association of this character was recorded with days of 50% flowering and days to first fruit initiation.

Secondary branches per plant at maturity was observed positive correlation with fruits per cluster, fruit width, fruit length, fruits per plant and seeds per fruit. Negative association of this character was recorded with days of 50% flowering, days to first fruit initiation and days to maturity.

Days to 50% flowering were exhibited positive with days to 1st first fruit initiation and days to maturity. Negative association of this character was recorded with fruit per clusters, fruit width, fruit length, fruits per plant and seeds per fruit.

Correlation coefficient of negative association of this character was recorded with fruit per clusters, fruit width, fruit length, fruits per plant, days to maturity and seeds per fruit.

Fruit length expressed a significant and positive association with fruits per plant, days to maturity and seeds per fruit.

Fruits per plant expressed a significant and positive association with days to maturity and seeds per fruit. Days to maturity expressed a positive association with and seeds per fruit.

6.5 Path coefficients

6.5.1 Genotypic path coefficients

Plant height revealed that positive direct effect on fruit dry weight per plant (g) while, it expressed positive indirect correlation primary branches per plant at maturity, secondary branches per plant at maturity, fruits per cluster, fruit width (cm), fruit length, fruits per plant,

days to maturity and seeds per fruit, while it negative indirect effect on days of 50% flowering and days to first fruit initiation.

Primary branches per plant showed high negative direct effect on fruit dry weight per plant while, it expressed positive indirect correlation days to 50% flowering, days to first fruit initiation and days to maturity, while it negative indirect effect on plant height (cm) at maturity, primary branches per plant at maturity, secondary branches per plant at maturity, fruits per cluster, fruit width, fruit length, fruits yield plant and seeds per fruit.

Secondary branches per plant high positive direct effect on fruit dry weightperplant while, it expressed positive indirect correlation days to 50% flowering and days to first fruit initiation, while it negative indirect effect on plant height, primary branches per plant at maturity, secondary branches per plant at maturity, fruits per cluster, fruit width, fruit length, fruits per plant, days to maturity and seeds per fruit.

Days to 50% flowering high negative direct effect on fruit dry weight per plant while, it expressed positive indirect correlation plant height, primary branches per plant, secondary branches per plant, fruits per cluster, fruit width, fruit length, fruits per plant, days to maturity and seeds per fruit, while it negative indirect effect on days to first fruit initiation.

Days to first fruit initiation high positive direct effect on fruit dry weight per plant while, it expressed positive indirect correlation plant height, primary branches per plant at maturity, secondary branches per plant, fruits per cluster, fruit width, fruit length, fruits per plant and seeds per fruit, while it negative indirect effect on days to 50% flowering and days to maturity.

Fruits per cluster high negative direct effect on fruit dry weight per plant, it expressed positive indirect correlation plant height, primary branches per plant, secondary branches per plant, fruit width, fruit length, fruits per plant and seeds per fruit, while it negative indirect effect on days to 50% flowering and days to first fruit initiation and days to maturity.

Fruit width high negative direct effect on fruit dry weight per plant while, it expressed positive indirect correlation plant height, primary branches per plant, secondary branches per plant, fruits per cluster, fruit length, fruits per plant, days to maturity and seeds per fruit, while it negative indirect effect on days to 50% flowering and days to first fruit initiation.

6.7 Conclusion:

The following conclusions have been drawn from the present investigation as:

1. The high genetic variability were observed for major fruit yield traits namely plant height, branches per plant, days to fruit initiation, fruit length and fruit per plant.
2. High genotypic coefficient of variation were noticed for fruit width followed fruit length, fruit per cluster while low magnitudes were recorded for days to first fruit initiation, days to first fruit picking and days to 50% flowering.
3. High heritability estimated were exhibited for traits like fruits per plant, fruit length, plant height at 60 DAT, plant height at 90 DAT, plant height at maturity, secondary branches at 90 DAT, seeds per fruit, 1000 seed weight, However low estimated were noticed by days to first fruit initiation and fresh fruit weight.
4. High heritability coupled with high genetic advance were found for fruits per plant, plant height at 120 DAT and plant height at maturity, While low estimates were shown by days to first fruit initiation followed by fruit dry weight per plant.
5. Significant and positive association of fruit yield per plant were recorded with fruit length, plant height at maturity, primary branches per plant secondary branches per plant and fruits per cluster with it showed significant negative association with days to 50% flowering followed by days to first fruit initiation.
6. The maximum positive direct effect were observed by days to maturity, plant height, fruit per cluster on fruit yield per plant, However significant negative direct effect were observed by secondary branches per plant, days to 50% flowering and fruit length. Similarly maximum positive

indirect effect of plant height via secondary branches per plant, followed by fruits per cluster and fruit width on fruit yield were noticed secondary branches per plant though days to first fruit initiation and fruit per cluster via fruit width and fruit length were recorded.

7. Over all fruit yield contributing traits namely plant height, days to maturity, fruit length, primary branches per plant, fruit per cluster were found promising in chilli hybrids under present investigation.
8. Based on high fruit yield coupled with good quality traits. The hybrids namely Saniya, Jalsa and ARCH 930 were found promising for fresh marketable produce, while Tara, Amrita, Jwala garima-12 and TS0017 were found better for dry chilli purpose.

6.8 Suggestions for further work

1. Stability of distinct traits of chilli hybrid should be established over the years and seasons.
2. Traits identified for high heritability coupled with high or moderate genetic gain may be considered well in selection for the genetic improvement of crop.
3. Characters having desirable association with fresh and dry fruit yield should be given due consideration under selection for genetic improvement in chilli.
4. Some hybrids showing greater yield potential with other desirable qualities may be tested under different agro climatic conditions and those found suitable should be recommended for release for commercial cultivation.
5. The genetic diversity analysis be carried out to determine diverse parents for hybridization purpose to create further new variability

BIBLIOGRAPHY

- Abu N.E and Uguru M.I. (2006). Evaluation of genetic variations in growth and yield components of aromatic pepper lines in a derived savanna ecology of Nigeria. *Agronomic. Science*.**5** (1): 1-7.
- Adpawar, R. M.; Kale, P. B.; Kale, V. S.; Parlawar, N. D. and Yadgirwar, B. M. (2006). Heterosis for yield and yield components in chilli. *Annals of Plant Physiology*. **20** (1): 69-73.
- Ajjapplavara PS, Patil SS, Hosamani RM, Patil AA and Gangaprasad S. (2005) . Correlation and path coefficient analysis in chilli. *Karnataka Journal of Agricultural Sciences*.**18** (3): 748 –751.
- Amit, K, Ahad, I, Kumar V and Thakur S (2014) Genetic variability and correlation studies for growth and yield characters in chilli (*Capsicum annum* L.) *Journal of Spices and Aromatic Crops* **23** (2) : 170–177
- Anandanayaki, D. and Natarajan, S. (2000). Genetics of certain growth and yield parameters in chilli (*Capsicum annum* L.). *South Indian Horticulture*. **48** (1/6): 123-125.
- Ananthi, N., Jebaraj S. and Banu R. (2006). Variability studies in two-line rice (*Oryza sativa* L.)*Res. on Crops*, **7**(1):140–142.
- Ben Chaim A, Borovsky Y, Rao GU, Tanyolac B and Paran I. (2003) A major fruit shape QTL conserved in *Capsicum*. *Genome*. **46**: 1–9.
- Bharadwaj, D. N.; Harvinder Singh and Yadav, R. K. (2007a). Genetic variability and association of component characters for yield in chilli (*Capsicum annum* L.). *Progressive Agriculture*. **7** (1/2): 72-74.
- Bharadwaj, D. N.; Singh, S. K. and Singh, H. L. (2007). Genetic variability and association of component characters for yield in chilli. *International Journal of Plant Sciences* (Muzaffarnagar). **2** (2): 93-96.
- Burton, G.W. (1952). Quantitative inheritance in grasses. Proc. 6th Int. Grassland Cong. **1**: 227-285.
- Chattopadhyay, Arup; Sharangi, Amit Baran; Dai, Nuka and Dutta, Subrata. (2011) Diversity of genetic resources and genetic association analysis of green and dry chillies of Eastern India. *Chilean Journal of Agricultural Research*; Chillán **71**(3): 350-356.

- Chaudhary, Alok; Rajesh Kumar and Solankey, S. S. (2013). [Estimation of heterosis for yield and quality components in chilli \(*Capsicum annuum* L.\)](#). *African Journal of Biotechnology*. **12** (47): 6605-6610.
- Datta, S. and Jana, J. C. (2010). Genetic variability, heritability and correlation in chilli genotypes under Terai zone of West Bengal. *SAARC Journal of Agriculture*. **8** (1): 33-45.
- Diwaker Kumar; Vijay Bahadur; Rangare, S. B. and Devi Singh (2012). Genetic variability, heritability and correlation studies in chilli (*Capsicum annuum* L.). *Hort Flora Research Spectrum*. **1** (3): 248-252.
- Dutonde, S. N.; Bhalekar, M. N.; Warade, S. D.; Gupta, N. S. and Mahatale, P. V. (2006). Genetic variability and heritability in chilli (*Capsicum annuum* L.) with reference to heat tolerance. *Orissa Journal of Horticulture*. **34** (1): 107-109.
- Farhad M., Hasanuzzaman M., Biswas B. K., Azad A. K. and Arifuzzaman M. (2008). Reliability of Yield Contributing Characters for Improving Yield Potential in Chilli (*Capsicum annuum*). *Int. J. Sustain. Crop Prod.* **3** (3):30-38
- Gogoi, Dipendra and Gautam, B. P. (2002). Variability, heritability and genetic advance in chilli (*Capsicum* spp.). *Agricultural Science Digest*. **22** (2): 102-104.
- Gupta, A. M.; Singh Daljeet and Ajay Kumar (2009). Genetic variability, genetic advance and correlation in chilli (*Capsicum annuum* L.). *Indian Journal of Agricultural Sciences*. **79** (3): 221-223.
- Hanson, C.H.; Robinson, H.F. and Comstock, R.E. (1956). Biometrical studies of yield in segregating population of Korean Lespedeza. *Agron, J.*, **48** : 268-272.
- Hasanuzzaman, M.; Hakim, M. A.; Hanafi, M. M.; Shukor-Juraimi, A.; Islam, M. M. and Shamsuddin, A. K. M. (2013). Study of heterosis in Bangladeshi chilli (*Capsicum annuum* L.) landraces. *Agrociencia* (Montecillo). **47** (7): 683-690.
- Hosamani R M and Shivkumar (2008) Correlation and path coefficient analysis in chilli. *Indian J. Hort.* **65**: 349–352.
- Jabeen, N.; Mufti, S.; Khan, S. H.; Hussain, K.; Tasaduk Shafi and Sonam Spaldon (2012). Mean performance and genetic variability in chilli

- (*Capsicum annuum* L.). *International Journal of Plant Sciences* (Muzaffarnagar). **7** (1): 155-157.
- Jackson, M. L. (1967). Soil chemical analysis, Pub. Prentice Hall India, New Delhi, pp. 87-93.
- Johnson, H.W.; Robinson, H.F. and Comstock, R.E. (1955). Estimates of genetics and environment variability in soybean. *Agron. J.*, **47** : 314-318.
- Kaur, Balvir and Singh, Daljit (2007). Genetic variability in chilli (*Capsicum annuum* L.). *Haryana Journal of Horticultural Sciences*. **36** (3per4): 305-306.
- Krishna, C. Ukkund, M.P. Patil, M.B. Madalageri, Ravindra Mulage and Jagadeesh. R.C (2007). Character association and path analysis studies in green chilli for yield and yield attributes (*Capsicum annuum* L.) *Karnataka J. Agric. Sci.* **20(1)**: (99 -101).
- Krishnamurthy, S. L.; Rao, A. M.; Reddy, K. M.; Ramesh, S.; Shailaja Hittalmani and Rao, M. G. (2013). Limits of parental divergence for the occurrence of heterosis through morphological and AFLP marker in chilli (*Capsicum annuum* L.). *Current Science*. **104** (6): 738-746.
- Kumar, Rajesh and Lal, Gulshan (2001). Expression of heterosis in hot pepper (*Capsicum annuum* L.). *Capsicum and Eggplant Newsletter*. (20): 38-41.
- Kumar, S.; Kumar, R., and Singh, J. (2006). Cayenne per American pepper (*Capsicum* species). Peter K.V. (ed), Handbook of Herbs and spices, Vo1.3, Wood head Publishing, Cambridge, U.K., pp. 299-312.
- Manju, P. R. and Sreelathakumary, I. (2002 b). Genetic cataloguing of hot chilli (*Capsicum chinense* Jacq.) types of Kerala. *Journal of Tropical Agriculture*. **40** (1per2): 42-44.
- Manju, P. R. and Sreelathakumary, I. (2002 a). Genetic variability, heritability and genetic advance in hot chilli (*Capsicum chinense* Jacq.). *Journal of Tropical Agriculture*. **40** (1per2): 4-6.
- Miller, D.A.; Williams, J.C.; Robinson, H.F. and Comstock, K.B. (1958). Estimates of genotypic and environmental variances and covariances in upland cotton and their implication in selection. *Agron. J.* **50**: 126-131.

- Mini, S. and Khader, K. M. A. (2004). Variability, heritability and genetic advance in wax type chilli (*Capsicum annuum* L.). *Capsicum & Eggplant Newsletter*. (23): 49-52.
- Munshi A D and Behera T K (2000) Genetic variability, heritability and genetic advance for some traits in chilli. (*Capsicum annuum* L.) *Veg. Sci.* **27**: 39–41.
- Nandadevi and Hosamani, R. M. (2003). Variability, correlation and path analysis in *kharif*-grown chilli (*Capsicum annuum* L.) genotypes for different characters. *Capsicum & Eggplant Newsletter*. (22): 43-46.
- Nehru, S. D.; Manjunath, A. and Rangaiah, S. (2003). Genetic variability and stability for fruit yield and other metrical characters in chilli (*Capsicum annuum* L.). *Karnataka Journal of Agricultural Sciences*. **16** (1): 44-47.
- NHB (2013).www.nhb.gov.in
- Olsen, S. R.; C V. Cole and L. A. Dean (1954). Estimation of available phosphorus in soil by extraction with sodium carbonate. USDA Circular No.939, Washington. pp 15.
- Padhar, P. R. and Zaveri, P. P. (2010). Genetic studies in relation to selection criteria in chilli. *Research on Crops*. **11** (3): 722-727.
- Pandit M.K and Ahikary S. (2014). Variability and heritability estimates in some reproductive characters and yield in chilli (*Capsicum annuum* L.) *International Journal of Plant and Soil Science*.**3** (7): XX-XX.
- Pandit M.K, Muthukumar P and Mukhopadhyay T.P. (2009). Genetic variability, character association and path analysis in chilli (*Capsicum annuum* L.) genotypes. ICH pp 41.
- Panse, V. C. and P. V. Sukhatme (1985). Statistical methods for agricultural workers. ICAR Publications, New Delhi. pp 155.
- Patel, H. B.; Bhatt, M. M.; Patel, J. S. and Patel, J. A. (2008). Heterosis for green fruit yield and its quality attributes in chilli (*Capsicum annuum* L.). *Research on Crops*. **9** (2): 350-352.
- Patel, J. A.; Patel, M. J.; Acharya, R. R.; Bhanvadia, A. S. and Bhalala, M. K. (2004). Hybrid vigour, gene action and combining ability in chilli (*Capsicum annuum* L.) hybrids involving male sterile lines. *Indian Journal of Genetics and Plant Breeding*. **64** (1): 81-82.

- Patel, J. A.; Patel, M. J.; Patel, A. D.; Acharya, R. R. and Bhalala, M. K. (2001). Heterosis [studies over environments in chilli \(*Capsicum annuum* L.\)](#). *Vegetable Science*. **28** (2): 130-132.
- Patel, M. P.; Patel, A. R.; Patel, J. B. and Patel, J. A. (2010). Heterosis for green fruit yield and its components in chilli (*Capsicum annuum* var. longicum (D.G) Sendt) over environments. *Electronic Journal of Plant Breeding*. **1** (6): 1443-1453.
- Patel, P. N.; Fougat, R. S. and Sasidharan, N. (2009). Studies on genetic variability, correlation and path analysis in chillies (*Capsicum annuum* L.). *Research on Crops*. **10** (3): 626-631.
- Piper, C S. (1967). Soil and plant analysis. *Uni.of Adelaide, Australia*. Hans Publishers, Bombay.
- Prabhakaran, T. S.; Natarajan, S. and Veeraragavatham, D. (2004). Studies on genetic variability, heritability and genetic advance in chilli (*Capsicum annuum* L.). *South Indian Horticulture*. **52** (1/6): 70-72.
- Prajapati, D. B. and Agalodia, A. V. (2011). Heterosis and inbreeding depression in chilli (*Capsicum annuum* L.). *Journal of Spices and Aromatic Crops*. **20** (2): 72-76.
- Rathod, R. P.; Deshmukh, D. T.; Sable, N. H. and Rathod, N. G. (2002). Genetic variability studies in chilli (*Capsicum annuum* L.). *Journal of Soils and Crops*. **12** (2): 210-212.
- Saffarod, W. E. (1926). Our heritage from the American Indians. *Annual Report, Smithsonian Institute*, pp. 405-410.
- Samadia, D. K. (2007). Genetic variability studies in chilli germplasm under hot arid eco-system. *Indian Journal of Horticulture*. **64** (4): 477-479.
- Sarkar, S.; Murmu, D.; Chattopadhyay, A. and Hazra, P. (2009). Genetic variability, correlation and path analysis of some morphological characters in chilli. *Journal of Crop and Weed*. **5** (1): 162-166.
- Satish, R. G. and Lad, D. B. (2007). Heterosis studies in chilli. *Journal of Maharashtra Agricultural Universities*. **32** (1): 68-71.
- Shankarnag, B.; Madalageri, M. B. and Ravindra Mulge (2006). Manifestation of heterosis for growth, earliness and early green fruit yield in chilli. *Indian Journal of Horticulture*. **63** (4): 410-414.

- Sharma, Madhu; Singh, Yudhvir and Jamwal, R. S. (2009). Variability studies for various metric traits in chilli. *Haryana Journal of Horticultural Sciences*. **38** (3/4): 284-287.
- Sharma, V. K., Semwal, C. S. and Uniyal, S. P. (2010). Genetic variability and character association analysis in bell pepper (*Capsicum annuum* L.). *J. Horti forestry*. **2** (3): 058-065.
- Shirshat S.S, Giritammannavar V.A and Patil S.J. (2007.) Analysis of genetic variability for quantitative traits in chilli. *Karnataka Journal of Agricultural Sciences*. **20** (1): 29–32.
- Singh, D. K.; Pramod Tewari and Jain, S. K. (2012). Heterosis studies for growth, flowering, and yield of chilli (*Capsicum annuum* L.). *Pantnagar Journal of Research*. **10** (1): 61-65.
- Singh, R. K. and Singh, D. B. (2013). Genetic variability and characters association in chilli (*Capsicum annuum* L). *SAARC Journal of Agriculture*. **10** (1): 71-80.
- Singh, Yudhvir; Sharma, Madhu and Sharma, Akhilesh (2009). Genetic variation, association of characters and their direct and indirect contributions for improvement in chilli peppers. *International Journal of Vegetable Science*. **15** (4): 340-368.
- Smitha, R. P. and Basavaraja, N. (2006). Variability and correlation studies in chilli (*Capsicum annuum* L.). *Karnataka Journal of Agricultural Sciences*. **19** (4): 888-891.
- Snedecor, G.W. and Cochran, C.W. 1967. *Stastical methods*. The Iowa State University Press, Iowa, USA.
- Sreelathakumary, I. and Rajamony, L. (2004). Variability, heritability and genetic advance in chilli (*Capsicum annuum* L.). *Journal of Tropical Agriculture*. **42** (1per2): 35-37.
- Subbiah, B. V. and Asija, G. L. (1956). A rapid procedure for the determination of available N in soils. *Curr. Sc.* **25**: 256-260.
- Tembhurne, B. V. and Rao, S. K. (2012). Heterosis and combining ability in CMS based hybrid chilli (*Capsicum annuum* L.). *Journal of Agricultural Science (Toronto)*. **4** (10): 89-96.
- Tembhurne, B. V.; Revanappa, R. and Kuchanur, P. H. (2008). Varietal performance, genetic variability and correlation studies in chilli

- (*Capsicum annuum* L.). *Karnataka Journal of Agricultural Sciences*. **21** (4): 541-543.
- Ukkund, K. C.; Patil, M. P.; Madalageri, M. B.; Ravindra Mulge; Patil, B. R. and Jagadeesh, R. C. (2006). Studies on genetic variability, heritability and genetic advances in green chilli (*Capsicum annuum*) var. Longum. *Journal of Asian Horticulture*. **2** (1per2): 4-8.
- Ullah M.Z., M.J. Hasan, A.I. Saki, M.A. Rahman and Biswas P.L. (2011). Association of correlation and cause effect analysis among morphological traits in chilli (*Capsicum frutescens* L.). *Intl. J. Bio Res.* **10** (6): 19-24.
- Vaishnav, R. S.; Srivastava, K. and Singh, P. K. (2010). Heterosis and combining ability in chilli for yield and yield contributing traits (*Capsicum annuum* L.). *Asian Journal of Bio Science*. **4** (2): 244-248.
- Vani, S. K.; Sridevi, O. and Salimath, P. M. (2007). Studies on genetic variability, correlation and path analysis in chilli (*Capsicum annuum* L.). *Annals of Biology*. **23** (2): 117-121.
- Varkey, J.; Saiyed, M. P.; Patel, J. S. and Patel, D. B. (2005). Genetic variability and heritability in chilli. *Journal of Maharashtra Agricultural Universities*. **30** (3): 346-347.
- Vikram Amit , Warshamana, I.K and Gupta Meenu (2014) Genetic correlation and path coefficient studies on yield and biochemical traits in chilli (*Capsicum annuum* L) . *International Journal of Farm Sciences* **4**(2):70-75.
- Walkley, A. J. and Black, J. A. (1934). Estimation of organic carbon by chromic acid titration method. *Soil Science*, **37**: 29-38.
- Wilson D and Philip (2009). Genetic variability and genetic divergence in paprika (*Capsicum annuum* L.). ICH pp 16.
- Yadwad, Arati; Sridevi, O. and Salimath, P. M. (2008). Genetic variability in segregating progenies of chilli (*Capsicum annuum* L.). *International Journal of Plant Sciences* (Muzaffarnagar). **3** (1): 106-110.

APPENDIX

| ANOVA for Plant Height (cm) 30 DAT | | | | | | |
|------------------------------------|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 0.8411 | 0.4206 | 0.2278 | 0.7975 | |
| Treatments | 17 | 594.8467 | 34.9910 | 18.9546 | 0.0000 | *** |
| Error | 34 | 62.7656 | 1.8460 | | | |
| General Mean | | 28.3222 | 0.7844 | 95% | 99% | |
| S.E.Diff. | | 1.1094 | Critical Diff. | 2.2545 | 3.0268 | |
| S.E.Diff from Mean | | 0.7623 | Critical Diff. | 1.5493 | 2.08 | |
| Var Environmental | | 1.846 | ECV | 4.7973 | | |
| Var Genotypical | | 11.0483 | GCV | 11.736 | | |
| Var Phenotypical | | 12.8944 | PCV | 12.6786 | | |
| h ² (Broad Sense) | | 0.8568 | S.E. of h ² | 0.7547 | | |
| Genetic Advancement | | 6.3382 | 8.1227 | | | |
| Gen.Adv as % of Mean | | 22.3788 | 28.6796 | (selection intensity at 5% & 1%) | | |

| ANOVA for Plant Height (cm) 60 DAT | | | | | | |
|------------------------------------|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 1.3011 | 0.6506 | 0.3295 | 0.7215 | |
| Treatments | 17 | 3007.3284 | 176.9017 | 89.6120 | 0.0000 | *** |
| Error | 34 | 67.1189 | 1.9741 | | | |
| General Mean | | 49.6944 | 0.8112 | 95% | 99% | |
| S.E.Diff. | | 1.1472 | Critical Diff. | 2.3314 | 3.13 | |
| S.E.Diff from Mean | | 0.7883 | Critical Diff. | 1.6021 | 2.1509 | |
| Var Environmental | | 1.9741 | ECV | 2.8273 | | |
| Var Genotypical | | 58.3092 | GCV | 15.366 | | |
| Var Phenotypical | | 60.2833 | PCV | 15.6239 | | |
| h ² (Broad Sense) | | 0.9673 | S.E. of h ² | 0.7683 | | |
| Genetic Advancement | | 15.4706 | 19.8263 | | | |
| Gen.Adv as % of Mean | | 31.1314 | 39.8965 | (selection intensity at 5% & 1%) | | |

| ANOVA for Plant Height (cm) 90 DAT | | | | | | |
|------------------------------------|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 0.5233 | 0.2617 | 0.0791 | 0.9241 | |
| Treatments | 17 | 3228.2200 | 189.8953 | 57.3923 | 0.0000 | *** |
| Error | 34 | 112.4967 | 3.3087 | | | |
| General Mean | | 71.1333 | 1.0502 | 95% | 99% | |
| S.E.Diff. | | 1.4852 | Critical Diff. | 3.0183 | 4.0522 | |
| S.E.Diff from Mean | | 1.0206 | Critical Diff. | 2.0741 | 2.7846 | |
| Var Environmental | | 3.3087 | ECV | 2.5572 | | |
| Var Genotypical | | 62.1955 | GCV | 11.0868 | | |
| Var Phenotypical | | 65.5042 | PCV | 11.3779 | | |
| h ² (Broad Sense) | | 0.9495 | S.E. of h ² | 0.7662 | | |
| Genetic Advancement | | 15.8304 | 20.2875 | | | |
| Gen.Adv as % of Mean | | 22.2545 | 28.5204 | (selection intensity at 5% & 1%) | | |

| ANOVA for Plant Height (cm) 120 DAT | | | | | | |
|-------------------------------------|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 10.8811 | 5.4406 | 0.6977 | 0.5047 | |
| Treatments | 17 | 4974.0947 | 292.5938 | 37.5244 | 0.0000 | *** |
| Error | 34 | 265.1122 | 7.7974 | | | |
| General Mean | | 76.3389 | 1.6122 | 95% | 99% | |
| S.E.Diff. | | 2.28 | Critical Diff. | 4.6334 | 6.2206 | |
| S.E.Diff from Mean | | 1.5668 | Critical Diff. | 3.184 | 4.2747 | |
| Var Environmental | | 7.7974 | ECV | 3.6579 | | |
| Var Genotypical | | 94.9321 | GCV | 12.7632 | | |
| Var Phenotypical | | 102.7296 | PCV | 13.2771 | | |
| h ² (Broad Sense) | | 0.9241 | S.E. of h ² | 0.763 | | |
| Genetic Advancement | | 19.2945 | 24.7269 | | | |
| Gen.Adv as % of Mean | | 25.2748 | 32.3909 | (selection intensity at 5% & 1%) | | |

| ANOVA for Plant Height (cm) Maturity | | | | | | |
|--------------------------------------|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 5.3404 | 2.6702 | 0.6485 | 0.5292 | |
| Treatments | 17 | 4349.3540 | 255.8444 | 62.1397 | 0.0000 | *** |
| Error | 34 | 139.9863 | 4.1172 | | | |
| General Mean | | 76.987 | 1.1715 | 95% | 99% | |
| S.E.Diff. | | 1.6568 | Critical Diff. | 3.3669 | 4.5202 | |
| S.E.Diff from Mean | | 1.1385 | Critical Diff. | 2.3137 | 3.1062 | |
| Var Environmental | | 4.1172 | ECV | 2.6356 | | |
| Var Genotypical | | 83.909 | GCV | 11.8984 | | |
| Var Phenotypical | | 88.0263 | PCV | 12.1868 | | |
| h ² (Broad Sense) | | 0.9532 | S.E. of h ² | 0.7666 | | |
| Genetic Advancement | | 18.4234 | 23.6106 | | | |
| Gen.Adv as % of Mean | | 23.9305 | 30.6682 | (selection intensity at 5% & 1%) | | |

| ANOVA for Primary Branchesper Plant 30 DAT | | | | | | |
|--|----|----------------|------------------------|----------------------------------|-------------|----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 0.9537 | 0.4769 | 1.5134 | 0.2346 | |
| Treatments | 17 | 18.0953 | 1.0644 | 3.3782 | 0.0013 | ** |
| Error | 34 | 10.7130 | 0.3151 | | | |
| General Mean | | 4.7963 | 0.3241 | 95% | 99% | |
| S.E.Diff. | | 0.4583 | Critical Diff. | 0.9314 | 1.2505 | |
| S.E.Diff from Mean | | 0.315 | Critical Diff. | 0.6401 | 0.8593 | |
| Var Environmental | | 0.3151 | ECV | 11.7033 | | |
| Var Genotypical | | 0.2498 | GCV | 10.4201 | | |
| Var Phenotypical | | 0.5649 | PCV | 15.6699 | | |
| h ² (Broad Sense) | | 0.4422 | S.E. of h ² | 0.7012 | | |
| Genetic Advancement | | 0.6846 | 0.8774 | | | |
| Gen.Adv as % of Mean | | 14.274 | 18.2929 | (selection intensity at 5% & 1%) | | |

| ANOVA for Primary Branchesper Plant 60 DAT | | | | | | |
|--|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 5.4815 | 2.7407 | 1.4594 | 0.2466 | |
| Treatments | 17 | 129.9259 | 7.6427 | 4.0696 | 0.0003 | *** |
| Error | 34 | 63.8519 | 1.8780 | | | |
| General Mean | | 10.2963 | 0.7912 | 95% | 99% | |
| S.E.Diff. | | 1.1189 | Critical Diff. | 2.2739 | 3.0529 | |
| S.E.Diff from Mean | | 0.7689 | Critical Diff. | 1.5626 | 2.0979 | |
| Var Environmental | | 1.878 | ECV | 13.3096 | | |
| Var Genotypical | | 1.9216 | GCV | 13.4632 | | |
| Var Phenotypical | | 3.7996 | PCV | 18.9315 | | |
| h ² (Broad Sense) | | 0.5057 | S.E. of h ² | 0.7097 | | |
| Genetic Advancement | | 2.0307 | 2.6025 | | | |
| Gen.Adv as % of Mean | | 19.7231 | 25.2762 | (selection intensity at 5% & 1%) | | |

| ANOVA for Primary Branchesper Plant 90 DAT | | | | | | |
|--|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 5.7778 | 2.8889 | 3.1799 | 0.0542 | |
| Treatments | 17 | 328.6667 | 19.3333 | 21.2806 | 0.0000 | *** |
| Error | 34 | 30.8889 | 0.9085 | | | |
| General Mean | | 13.1111 | 0.5503 | 95% | 99% | |
| S.E.Diff. | | 0.7782 | Critical Diff. | 1.5816 | 2.1233 | |
| S.E.Diff from Mean | | 0.5348 | Critical Diff. | 1.0868 | 1.4591 | |
| Var Environmental | | 0.9085 | ECV | 7.2698 | | |
| Var Genotypical | | 6.1416 | GCV | 18.9017 | | |
| Var Phenotypical | | 7.0501 | PCV | 20.2516 | | |
| h ² (Broad Sense) | | 0.8711 | S.E. of h ² | 0.7565 | | |
| Genetic Advancement | | 4.7649 | 6.1064 | | | |
| Gen.Adv as % of Mean | | 36.3423 | 46.5746 | (selection intensity at 5% & 1%) | | |

| ANOVA for Primary Branchesper Plant 120 DAT | | | | | | |
|---|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 1.3333 | 0.6667 | 0.5075 | 0.6065 | |
| Treatments | 17 | 510.8333 | 30.0490 | 22.8731 | 0.0000 | *** |
| Error | 34 | 44.6667 | 1.3137 | | | |
| General Mean | | 18.3889 | 0.6617 | 95% | 99% | |
| S.E.Diff. | | 0.9359 | Critical Diff. | 1.9019 | 2.5534 | |
| S.E.Diff from Mean | | 0.6431 | Critical Diff. | 1.3069 | 1.7546 | |
| Var Environmental | | 1.3137 | ECV | 6.233 | | |
| Var Genotypical | | 9.5784 | GCV | 16.8303 | | |
| Var Phenotypical | | 10.8922 | PCV | 17.9474 | | |
| h ² (Broad Sense) | | 0.8794 | S.E. of h ² | 0.7575 | | |
| Genetic Advancement | | 5.9787 | 7.662 | | | |
| Gen.Adv as % of Mean | | 32.5124 | 41.6664 | (selection intensity at 5% & 1%) | | |

| ANOVA for Primary Branchesper Plant Maturity | | | | | | |
|--|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 1.3333 | 0.6667 | 0.5075 | 0.6065 | |
| Treatments | 17 | 510.8333 | 30.0490 | 22.8731 | 0.0000 | *** |
| Error | 34 | 44.6667 | 1.3137 | | | |
| General Mean | | 18.3889 | 0.6617 | 95% | 99% | |
| S.E.Diff. | | 0.9359 | Critical Diff. | 1.9019 | 2.5534 | |
| S.E.Diff from Mean | | 0.6431 | Critical Diff. | 1.3069 | 1.7546 | |
| Var Environmental | | 1.3137 | ECV | 6.233 | | |
| Var Genotypical | | 9.5784 | GCV | 16.8303 | | |
| Var Phenotypical | | 10.8922 | PCV | 17.9474 | | |
| h ² (Broad Sense) | | 0.8794 | S.E. of h ² | 0.7575 | | |
| Genetic Advancement | | 5.9787 | 7.662 | | | |
| Gen.Adv as % of Mean | | 32.5124 | 41.6664 | (selection intensity at 5% & 1%) | | |

| ANOVA for Secondary Branchesper Plant 30 DAT | | | | | | |
|--|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 2.7037 | 1.3519 | 1.0299 | 0.3679 | |
| Treatments | 17 | 403.6482 | 23.7440 | 18.0888 | 0.0000 | *** |
| Error | 34 | 44.6296 | 1.3126 | | | |
| General Mean | | 14.9815 | 0.6615 | 95% | 99% | |
| S.E.Diff. | | 0.9355 | Critical Diff. | 1.9011 | 2.5523 | |
| S.E.Diff from Mean | | 0.6428 | Critical Diff. | 1.3064 | 1.7539 | |
| Var Environmental | | 1.3126 | ECV | 7.6475 | | |
| Var Genotypical | | 7.4771 | GCV | 18.2521 | | |
| Var Phenotypical | | 8.7898 | PCV | 19.7895 | | |
| h ² (Broad Sense) | | 0.8507 | S.E. of h ² | 0.754 | | |
| Genetic Advancement | | 5.1953 | 6.6581 | | | |
| Gen.Adv as % of Mean | | 34.6784 | 44.4422 | (selection intensity at 5% & 1%) | | |

| ANOVA for Secondary Branchesper Plant 60 DAT | | | | | | |
|--|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 9.1481 | 4.5741 | 1.6512 | 0.2068 | |
| Treatments | 17 | 1351.2037 | 79.4826 | 28.6925 | 0.0000 | *** |
| Error | 34 | 94.1852 | 2.7702 | | | |
| General Mean | | 24.9074 | 0.9609 | 95% | 99% | |
| S.E.Diff. | | 1.359 | Critical Diff. | 2.7617 | 3.7078 | |
| S.E.Diff from Mean | | 0.9339 | Critical Diff. | 1.8978 | 2.5479 | |
| Var Environmental | | 2.7702 | ECV | 6.6823 | | |
| Var Genotypical | | 25.5708 | GCV | 20.3022 | | |
| Var Phenotypical | | 28.341 | PCV | 21.3737 | | |
| h ² (Broad Sense) | | 0.9023 | S.E. of h ² | 0.7604 | | |
| Genetic Advancement | | 9.8947 | 12.6806 | | | |
| Gen.Adv as % of Mean | | 39.7261 | 50.9111 | (selection intensity at 5% & 1%) | | |

| ANOVA for Secondary Branchesper Plant 90 DAT | | | | | | |
|--|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 2.1111 | 1.0556 | 0.7600 | 0.4754 | |
| Treatments | 17 | 1412.0000 | 83.0588 | 59.8024 | 0.0000 | *** |
| Error | 34 | 47.2222 | 1.3889 | | | |
| General Mean | | 34.8889 | 0.6804 | 95% | 99% | |
| S.E.Diff. | | 0.9623 | Critical Diff. | 1.9555 | 2.6254 | |
| S.E.Diff from Mean | | 0.6612 | Critical Diff. | 1.3438 | 1.8041 | |
| Var Environmental | | 1.3889 | ECV | 3.3779 | | |
| Var Genotypical | | 27.2233 | GCV | 14.9549 | | |
| Var Phenotypical | | 28.6122 | PCV | 15.3316 | | |
| h ² (Broad Sense) | | 0.9515 | S.E. of h ² | 0.7664 | | |
| Genetic Advancement | | 10.4841 | 13.436 | | | |
| Gen.Adv as % of Mean | | 30.0501 | 38.5108 | (selection intensity at 5% & 1%) | | |

| ANOVA for Secondary Branchesper Plant 120 DAT | | | | | | |
|---|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 0.3333 | 0.1667 | 0.0642 | 0.9380 | |
| Treatments | 17 | 1500.6666 | 88.2745 | 33.9774 | 0.0000 | *** |
| Error | 34 | 88.3333 | 2.5980 | | | |
| General Mean | | 40.7778 | 0.9306 | 95% | 99% | |
| S.E.Diff. | | 1.3161 | Critical Diff. | 2.6745 | 3.5907 | |
| S.E.Diff from Mean | | 0.9044 | Critical Diff. | 1.8379 | 2.4675 | |
| Var Environmental | | 2.598 | ECV | 3.9527 | | |
| Var Genotypical | | 28.5588 | GCV | 13.1053 | | |
| Var Phenotypical | | 31.1569 | PCV | 13.6884 | | |
| h ² (Broad Sense) | | 0.9166 | S.E. of h ² | 0.7621 | | |
| Genetic Advancement | | 10.5398 | 13.5073 | | | |
| Gen.Adv as % of Mean | | 25.8468 | 33.1241 | (selection intensity at 5% & 1%) | | |

| ANOVA for Secondary Branchesper Plant Maturity | | | | | | |
|---|-----------|-----------------------|------------------------|----------------------------------|--------------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 0.3333 | 0.1667 | 0.0276 | 0.9728 | |
| Treatments | 17 | 1282.1666 | 75.4216 | 12.5089 | 0.0000 | *** |
| Error | 34 | 205.0000 | 6.0294 | | | |
| General Mean | | 42.5 | 1.4177 | 95% | 99% | |
| S.E.Diff. | | 2.0049 | Critical Diff. | 4.0744 | 5.4701 | |
| S.E.Diff from Mean | | 1.3777 | Critical Diff. | 2.7999 | 3.759 | |
| Var Environmental | | 6.0294 | ECV | 5.7776 | | |
| Var Genotypical | | 23.1307 | GCV | 11.3163 | | |
| Var Phenotypical | | 29.1601 | PCV | 12.7059 | | |
| h ² (Broad Sense) | | 0.7932 | S.E. of h ² | 0.7468 | | |
| Genetic Advancement | | 8.8239 | 11.3083 | | | |
| Gen.Adv as % of Mean | | 20.7622 | 26.6078 | (selection intensity at 5% & 1%) | | |

| ANOVA for Days to First Flower Initiation. | | | | | | |
|---|-----------|-----------------------|------------------------|----------------------------------|--------------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 19.1111 | 9.5556 | 1.4980 | 0.2380 | |
| Treatments | 17 | 1287.3334 | 75.7255 | 11.8709 | 0.0000 | *** |
| Error | 34 | 216.8889 | 6.3791 | | | |
| General Mean | | 39.2222 | 1.4582 | 95% | 99% | |
| S.E.Diff. | | 2.0622 | Critical Diff. | 4.1909 | 5.6265 | |
| S.E.Diff from Mean | | 1.4171 | Critical Diff. | 2.8799 | 3.8664 | |
| Var Environmental | | 6.3791 | ECV | 6.4394 | | |
| Var Genotypical | | 23.1155 | GCV | 12.258 | | |
| Var Phenotypical | | 29.4946 | PCV | 13.8465 | | |
| h ² (Broad Sense) | | 0.7837 | S.E. of h ² | 0.7456 | | |
| Genetic Advancement | | 8.768 | 11.2366 | | | |
| Gen.Adv as % of Mean | | 22.3546 | 28.6486 | (selection intensity at 5% & 1%) | | |

| ANOVA for Days to First Picking of Fruits | | | | | | |
|---|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 0.3341 | 0.1671 | 0.1919 | 0.8263 | |
| Treatments | 17 | 237.9830 | 13.9990 | 16.0787 | 0.0000 | *** |
| Error | 34 | 29.6022 | 0.8707 | | | |
| General Mean | | 75.1413 | 0.5387 | 95% | 99% | |
| S.E.Diff. | | 0.7619 | Critical Diff. | 1.5483 | 2.0786 | |
| S.E.Diff from Mean | | 0.5235 | Critical Diff. | 1.064 | 1.4284 | |
| Var Environmental | | 0.8707 | ECV | 1.2418 | | |
| Var Genotypical | | 4.3761 | GCV | 2.784 | | |
| Var Phenotypical | | 5.2468 | PCV | 3.0484 | | |
| h ² (Broad Sense) | | 0.8341 | S.E. of h ² | 0.7519 | | |
| Genetic Advancement | | 3.9356 | 5.0437 | | | |
| Gen.Adv as % of Mean | | 5.2376 | 6.7122 | (selection intensity at 5% & 1%) | | |

| ANOVA for Days of 50% Flowering | | | | | | |
|---------------------------------|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 0.3341 | 0.1671 | 0.1919 | 0.8263 | |
| Treatments | 17 | 237.9830 | 13.9990 | 16.0787 | 0.0000 | *** |
| Error | 34 | 29.6022 | 0.8707 | | | |
| General Mean | | 66.1413 | 0.5387 | 95% | 99% | |
| S.E.Diff. | | 0.7619 | Critical Diff. | 1.5483 | 2.0786 | |
| S.E.Diff from Mean | | 0.5235 | Critical Diff. | 1.064 | 1.4284 | |
| Var Environmental | | 0.8707 | ECV | 1.4107 | | |
| Var Genotypical | | 4.3761 | GCV | 3.1628 | | |
| Var Phenotypical | | 5.2468 | PCV | 3.4632 | | |
| h ² (Broad Sense) | | 0.8341 | S.E. of h ² | 0.7519 | | |
| Genetic Advancement | | 3.9356 | 5.0437 | | | |
| Gen.Adv as % of Mean | | 5.9503 | 7.6256 | (selection intensity at 5% & 1%) | | |

| ANOVA for Days to First Fruit Initiation | | | | | | |
|---|-----------|-----------------------|------------------------|----------------------------------|--------------------|--|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 2.2553 | 1.1276 | 0.1141 | 0.8925 | |
| Treatments | 17 | 237.9830 | 13.9990 | 1.4160 | 0.1891 | |
| Error | 34 | 336.1343 | 9.8863 | | | |
| General Mean | | 71.4746 | 1.8153 | 95% | 99% | |
| S.E.Diff. | | 2.5673 | Critical Diff. | 5.2173 | 7.0045 | |
| S.E.Diff from Mean | | 1.7642 | Critical Diff. | 3.5852 | 4.8134 | |
| Var Environmental | | 9.8863 | ECV | 4.3991 | | |
| Var Genotypical | | 1.3709 | GCV | 1.6381 | | |
| Var Phenotypical | | 11.2572 | PCV | 4.6942 | | |
| h ² (Broad Sense) | | 0.1218 | S.E. of h ² | 0.6569 | | |
| Genetic Advancement | | 0.8417 | 1.0787 | | | |
| Gen.Adv as % of Mean | | 1.1776 | 1.5092 | (selection intensity at 5% & 1%) | | |

| ANOVA for Fruitsper Cluster | | | | | | |
|------------------------------------|-----------|-----------------------|------------------------|----------------------------------|--------------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 0.3371 | 0.1685 | 0.0697 | 0.9328 | |
| Treatments | 17 | 448.5953 | 26.3880 | 10.9077 | 0.0000 | *** |
| Error | 34 | 82.2529 | 2.4192 | | | |
| General Mean | | 13.1209 | 0.898 | 95% | 99% | |
| S.E.Diff. | | 1.27 | Critical Diff. | 2.5809 | 3.4649 | |
| S.E.Diff from Mean | | 0.8727 | Critical Diff. | 1.7735 | 2.381 | |
| Var Environmental | | 2.4192 | ECV | 11.8542 | | |
| Var Genotypical | | 7.9896 | GCV | 21.5426 | | |
| Var Phenotypical | | 10.4088 | PCV | 24.5887 | | |
| h ² (Broad Sense) | | 0.7676 | S.E. of h ² | 0.7435 | | |
| Genetic Advancement | | 5.1014 | 6.5377 | | | |
| Gen.Adv as % of Mean | | 38.8801 | 49.8269 | (selection intensity at 5% & 1%) | | |

| ANOVA for Fruit Width (cm) | | | | | | |
|------------------------------|----|----------------|------------------------|----------------------------------|-------------|----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 0.0010 | 0.0005 | 0.1449 | 0.8657 | |
| Treatments | 17 | 0.2000 | 0.0118 | 3.3952 | 0.0012 | ** |
| Error | 34 | 0.1178 | 0.0035 | | | |
| General Mean | | 0.2193 | 0.034 | 95% | 99% | |
| S.E.Diff. | | 0.0481 | Critical Diff. | 0.0977 | 0.1311 | |
| S.E.Diff from Mean | | 0.033 | Critical Diff. | 0.0671 | 0.0901 | |
| Var Environmental | | 0.0035 | ECV | 26.8453 | | |
| Var Genotypical | | 0.0028 | GCV | 23.9871 | | |
| Var Phenotypical | | 0.0062 | PCV | 36.0007 | | |
| h ² (Broad Sense) | | 0.4439 | S.E. of h ² | 0.7015 | | |
| Genetic Advancement | | 0.0722 | 0.0925 | | | |
| Gen.Adv as % of Mean | | 32.9239 | 42.1938 | (selection intensity at 5% & 1%) | | |

| ANOVA for Fruit Length (cm) | | | | | | |
|------------------------------|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 0.4288 | 0.2144 | 14.2375 | 0.0000 | *** |
| Treatments | 17 | 72.7945 | 4.2820 | 284.3532 | 0.0000 | *** |
| Error | 34 | 0.5120 | 0.0151 | | | |
| General Mean | | 5.1628 | 0.0708 | 95% | 99% | |
| S.E.Diff. | | 0.1002 | Critical Diff. | 0.2036 | 0.2734 | |
| S.E.Diff from Mean | | 0.0689 | Critical Diff. | 0.1399 | 0.1879 | |
| Var Environmental | | 0.0151 | ECV | 2.3769 | | |
| Var Genotypical | | 1.4223 | GCV | 23.1002 | | |
| Var Phenotypical | | 1.4374 | PCV | 23.2222 | | |
| h ² (Broad Sense) | | 0.9895 | S.E. of h ² | 0.7711 | | |
| Genetic Advancement | | 2.4439 | 3.132 | | | |
| Gen.Adv as % of Mean | | 47.3365 | 60.6642 | (selection intensity at 5% & 1%) | | |

| ANOVA for Fruitsper Plant | | | | | | |
|------------------------------|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 320.0370 | 160.0185 | 92.7966 | 0.0000 | *** |
| Treatments | 17 | 14338.3701 | 843.4335 | 489.1169 | 0.0000 | *** |
| Error | 34 | 58.6296 | 1.7244 | | | |
| General Mean | | 108.4074 | 0.7582 | 95% | 99% | |
| S.E.Diff. | | 1.0722 | Critical Diff. | 2.1789 | 2.9253 | |
| S.E.Diff from Mean | | 0.7368 | Critical Diff. | 1.4973 | 2.0103 | |
| Var Environmental | | 1.7244 | ECV | 1.2113 | | |
| Var Genotypical | | 280.5697 | GCV | 15.4512 | | |
| Var Phenotypical | | 282.2941 | PCV | 15.4986 | | |
| h ² (Broad Sense) | | 0.9939 | S.E. of h ² | 0.7716 | | |
| Genetic Advancement | | 34.3999 | 44.0853 | | | |
| Gen.Adv as % of Mean | | 31.7321 | 40.6663 | (selection intensity at 5% & 1%) | | |

| ANOVA for Days to Maturity | | | | | | |
|------------------------------|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 3.7378 | 1.8689 | 0.1524 | 0.8592 | |
| Treatments | 17 | 1858.9199 | 109.3482 | 8.9161 | 0.0000 | *** |
| Error | 34 | 416.9827 | 12.2642 | | | |
| General Mean | | 154.7767 | 2.0219 | 95% | 99% | |
| S.E.Diff. | | 2.8594 | Critical Diff. | 5.8109 | 7.8015 | |
| S.E.Diff from Mean | | 1.9649 | Critical Diff. | 3.9932 | 5.3611 | |
| Var Environmental | | 12.2642 | ECV | 2.2626 | | |
| Var Genotypical | | 32.3613 | GCV | 3.6754 | | |
| Var Phenotypical | | 44.6255 | PCV | 4.316 | | |
| h ² (Broad Sense) | | 0.7252 | S.E. of h ² | 0.7382 | | |
| Genetic Advancement | | 9.9793 | 12.7891 | | | |
| Gen.Adv as % of Mean | | 6.4476 | 8.2629 | (selection intensity at 5% & 1%) | | |

| ANOVA for Fruit Fresh Weight Plant (g) | | | | | | |
|--|----|----------------|------------------------|----------------------------------|-------------|----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 31038.9043 | 15519.4521 | 1.7627 | 0.1869 | |
| Treatments | 17 | 414423.6875 | 24377.8633 | 2.7689 | 0.0057 | ** |
| Error | 34 | 299347.1250 | 8804.3271 | | | |
| General Mean | | 501.4117 | 54.1736 | 95% | 99% | |
| S.E.Diff. | | 76.613 | Critical Diff. | 155.695 | 209.0291 | |
| S.E.Diff from Mean | | 52.6472 | Critical Diff. | 106.991 | 143.6415 | |
| Var Environmental | | 8804.3269 | ECV | 18.7134 | | |
| Var Genotypical | | 5191.1789 | GCV | 14.3694 | | |
| Var Phenotypical | | 13995.5059 | PCV | 23.5939 | | |
| h ² (Broad Sense) | | 0.3709 | S.E. of h ² | 0.6916 | | |
| Genetic Advancement | | 90.3939 | 115.8446 | | | |
| Gen.Adv as % of Mean | | 18.0279 | 23.1037 | (selection intensity at 5% & 1%) | | |

| ANOVA for Fruit Fresh Weightper Plot (kg) | | | | | | |
|---|----|----------------|------------------------|----------------------------------|-------------|----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 27.8947 | 13.9474 | 1.7594 | 0.1875 | |
| Treatments | 17 | 373.0112 | 21.9418 | 2.7678 | 0.0057 | ** |
| Error | 34 | 269.5368 | 7.9276 | | | |
| General Mean | | 15.0422 | 1.6256 | 95% | 99% | |
| S.E.Diff. | | 2.2989 | Critical Diff. | 4.6719 | 6.2723 | |
| S.E.Diff from Mean | | 1.5798 | Critical Diff. | 3.2105 | 4.3102 | |
| Var Environmental | | 7.9276 | ECV | 18.7179 | | |
| Var Genotypical | | 4.6714 | GCV | 14.3685 | | |
| Var Phenotypical | | 12.599 | PCV | 23.5969 | | |
| h ² (Broad Sense) | | 0.3708 | S.E. of h ² | 0.6916 | | |
| Genetic Advancement | | 2.7111 | 3.4744 | | | |
| Gen.Adv as % of Mean | | 18.0234 | 23.098 | (selection intensity at 5% & 1%) | | |

| ANOVA for Fruit Fresh Weightper ha. (q.) | | | | | | |
|---|-----------|-----------------------|------------------------|----------------------------------|--------------------|----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 4966.1411 | 2483.0706 | 1.7627 | 0.1869 | |
| Treatments | 17 | 66307.6250 | 3900.4482 | 2.7689 | 0.0057 | ** |
| Error | 34 | 47894.9688 | 1408.6755 | | | |
| General Mean | | 200.5648 | 21.6693 | 95% | 99% | |
| S.E.Diff. | | 30.645 | Critical Diff. | 62.2778 | 83.6111 | |
| S.E.Diff from Mean | | 21.0588 | Critical Diff. | 42.7963 | 57.4563 | |
| Var Environmental | | 1408.6755 | ECV | 18.7133 | | |
| Var Genotypical | | 830.591 | GCV | 14.3694 | | |
| Var Phenotypical | | 2239.2665 | PCV | 23.5938 | | |
| h ² (Broad Sense) | | 0.3709 | S.E. of h ² | 0.6916 | | |
| Genetic Advancement | | 36.1578 | 46.3381 | | | |
| Gen.Adv as % of Mean | | 18.028 | 23.1038 | (selection intensity at 5% & 1%) | | |

| ANOVA for Fruit Dry Weightper Plant (g) | | | | | | |
|--|-----------|-----------------------|------------------------|----------------------------------|--------------------|----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 3448.4624 | 1724.2312 | 1.7625 | 0.1869 | |
| Treatments | 17 | 46047.0977 | 2708.6528 | 2.7688 | 0.0057 | ** |
| Error | 34 | 33261.4219 | 978.2770 | | | |
| General Mean | | 167.1367 | 18.058 | 95% | 99% | |
| S.E.Diff. | | 25.5379 | Critical Diff. | 51.8989 | 69.677 | |
| S.E.Diff from Mean | | 17.5492 | Critical Diff. | 35.6641 | 47.881 | |
| Var Environmental | | 978.2771 | ECV | 18.7137 | | |
| Var Genotypical | | 576.7919 | GCV | 14.3694 | | |
| Var Phenotypical | | 1555.069 | PCV | 23.5941 | | |
| h ² (Broad Sense) | | 0.3709 | S.E. of h ² | 0.6916 | | |
| Genetic Advancement | | 30.1309 | 38.6143 | | | |
| Gen.Adv as % of Mean | | 18.0277 | 23.1034 | (selection intensity at 5% & 1%) | | |

| ANOVA for Fruit Dry Weightper Plot (kg) | | | | | | |
|---|----|----------------|------------------------|----------------------------------|-------------|----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 3.1014 | 1.5507 | 1.7600 | 0.1874 | |
| Treatments | 17 | 41.4468 | 2.4380 | 2.7671 | 0.0057 | ** |
| Error | 34 | 29.9565 | 0.8811 | | | |
| General Mean | | 5.0143 | 0.5419 | 95% | 99% | |
| S.E.Diff. | | 0.7664 | Critical Diff. | 1.5575 | 2.0911 | |
| S.E.Diff from Mean | | 0.5267 | Critical Diff. | 1.0703 | 1.4369 | |
| Var Environmental | | 0.8811 | ECV | 18.7197 | | |
| Var Genotypical | | 0.519 | GCV | 14.3672 | | |
| Var Phenotypical | | 1.4001 | PCV | 23.5976 | | |
| h ² (Broad Sense) | | 0.3707 | S.E. of h ² | 0.6916 | | |
| Genetic Advancement | | 0.9036 | 1.1579 | | | |
| Gen.Adv as % of Mean | | 18.0196 | 23.0931 | (selection intensity at 5% & 1%) | | |

| ANOVA for Fruit Dry Weightper ha. (q.) | | | | | | |
|--|----|----------------|------------------------|----------------------------------|-------------|----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 551.6238 | 275.8119 | 1.7622 | 0.1870 | |
| Treatments | 17 | 7367.1948 | 433.3644 | 2.7688 | 0.0057 | ** |
| Error | 34 | 5321.6216 | 156.5183 | | | |
| General Mean | | 66.8546 | 7.2231 | 95% | 99% | |
| S.E.Diff. | | 10.215 | Critical Diff. | 20.7592 | 27.8703 | |
| S.E.Diff from Mean | | 7.0196 | Critical Diff. | 14.2654 | 19.152 | |
| Var Environmental | | 156.5183 | ECV | 18.7133 | | |
| Var Genotypical | | 92.282 | GCV | 14.369 | | |
| Var Phenotypical | | 248.8003 | PCV | 23.5936 | | |
| h ² (Broad Sense) | | 0.3709 | S.E. of h ² | 0.6916 | | |
| Genetic Advancement | | 12.052 | 15.4453 | | | |
| Gen.Adv as % of Mean | | 18.0272 | 23.1028 | (selection intensity at 5% & 1%) | | |

| ANOVA for Seedspers Fruit | | | | | | |
|------------------------------|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 3.5973 | 1.7986 | 1.2641 | 0.2954 | |
| Treatments | 17 | 847.2592 | 49.8388 | 35.0278 | 0.0000 | *** |
| Error | 34 | 48.3764 | 1.4228 | | | |
| General Mean | | 28.3704 | 0.6887 | 95% | 99% | |
| S.E.Diff. | | 0.9739 | Critical Diff. | 1.9793 | 2.6573 | |
| S.E.Diff from Mean | | 0.6693 | Critical Diff. | 1.3601 | 1.826 | |
| Var Environmental | | 1.4228 | ECV | 4.2045 | | |
| Var Genotypical | | 16.1386 | GCV | 14.1602 | | |
| Var Phenotypical | | 17.5615 | PCV | 14.7712 | | |
| h ² (Broad Sense) | | 0.919 | S.E. of h ² | 0.7624 | | |
| Genetic Advancement | | 7.9333 | 10.1669 | | | |
| Gen.Adv as % of Mean | | 27.9633 | 35.8365 | (selection intensity at 5% & 1%) | | |

| ANOVA for 1000 Seed Weight (g) | | | | | | |
|--------------------------------|----|----------------|------------------------|----------------------------------|-------------|-----|
| Source of Variations | df | Sum of Squares | Mean Squares | F Ratio | Probability | |
| Replicate | 2 | 0.3972 | 0.1986 | 1.2571 | 0.2974 | |
| Treatments | 17 | 94.1760 | 5.5398 | 35.0617 | 0.0000 | *** |
| Error | 34 | 5.3720 | 0.1580 | | | |
| General Mean | | 9.4565 | 0.2295 | 95% | 99% | |
| S.E.Diff. | | 0.3246 | Critical Diff. | 0.6596 | 0.8855 | |
| S.E.Diff from Mean | | 0.223 | Critical Diff. | 0.4532 | 0.6085 | |
| Var Environmental | | 0.158 | ECV | 4.2034 | | |
| Var Genotypical | | 1.7939 | GCV | 14.1635 | | |
| Var Phenotypical | | 1.9519 | PCV | 14.7741 | | |
| h ² (Broad Sense) | | 0.9191 | S.E. of h ² | 0.7624 | | |
| Genetic Advancement | | 2.6451 | 3.3898 | | | |
| Gen.Adv as % of Mean | | 27.9711 | 35.8465 | (selection intensity at 5% & 1%) | | |

VITA

The author of this thesis Mr. Ravindra Patidar S/O Shri Premchandra Patidar was born on 20th January 1990 at Khargone, M.P. He completed his primary and middle school education from Khargone H.S.C. (10th) from Bal Siksha Niketan H.S.Khargone and H.S.S.C. (12th) from Govt. D.A.Boys H.S.School Khargone, District Khargone(M.P.) in 2007

He joined College of Agriculture, Indore (M.P.) in 2009-10 and completed B.Sc. (Ag) degree in 2013 affiliated to Rajmata Vijayaraje Scindia Krishi Vishwa Vidyalaya (RVSKVV), Gwalior with first division (7.2 OGPA). He further admitted in department of Horticulture, RVSKVV - R.A.K. College of Agriculture, Sehore and completed his M.Sc.(Hort.) in Vegetable science with first division in 2017.

Ravindra Patidar