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**GENETIC CATALOGUING OF  
HOT CHILLI (*Capsicum chinense* Jacq.)**



BY

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**THESIS  
submitted in partial fulfilment of the  
requirement for the degree  
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**Department of Olericulture  
COLLEGE OF AGRICULTURE  
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**2001**

*Dedicated*

*to*

*Amma and Achan*

## DECLARATION

I hereby declare that this thesis entitled **Genetic cataloguing of hot chilli (*Capsicum chinense* Jacq)** is a bonafide record of research work done by me during the course of research and that the thesis has not previously formed the basis for the award of any degree, diploma, associateship, fellowship or other similar title of any other university or society.

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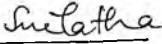


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## CERTIFICATE

Certified that this thesis entitled **Genetic cataloguing of hot chilli (*Capsicum chinense* Jacq)** is a record of research work done independently by Ms Manju P R (99 12 09) under my guidance and supervision and that she has not previously formed the basis for the award of any degree fellowship or associateship to her

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# *Introduction*

## 1 INTRODUCTION

Chilli or red pepper is an important vegetable cum spice crop grown throughout India for domestic as well as export markets. They are consumed in green, red as well as sun-dried conditions. It is unique among spice crops being a natural source of capsaicin, the pungent principle in chilli and colour.

Capsaicin is an effective counter-irritant and hence chilli extracts are used in pharmaceutical and cosmetic preparations. Since many countries including India are applying more and more restriction on the use of artificial colours, chilli colour may have a good demand as a substitute in food industry. Also, oleoresin obtained by solvent extraction finds application in food, pharmaceutical and cosmetic industries. The green chilli fruits are valuable on account of the richness in ascorbic acid.

Chilli belongs to the family Solanaceae and comes under the genus *Capsicum*. Recognizing the extent of variability, modern taxonomists have consolidated the cultivated *Capsicum* into the following five species: *Capsicum annum*, *C. frutescens*, *C. chinense*, *C. baccatum* and *C. pubescens*.

Most of the chillies available in the market belong to *C. annum* which is the most widely cultivated species. The present breeding works in chilli is mainly concentrated on the utilization of a relatively narrow gene base within the various cultivar groups, even though a wide genetic diversity exists both intraspecifically and interspecifically. To evolve varieties suited for specific purpose, widening of genetic base is inevitable.

*C chinense* one of the species grown in the homesteads of Kerala can be utilized for the improvement of the present day cultivars of *C annuum*. Characterized by its typical flavour and aroma the species is noted for its high oleoresin content and pungency. The plant is perennial in habit and bears two to five flowers per node as against solitary bearing in *C annuum* cultivars. Moreover the species is reported to be resistant to several pathogens including *Ralstonia solanacearum* and tomato spotted wilt virus which attack *C annuum* in the humid tropics.

Recently this crop is being exported to Maldives from Thiruvananthapuram airport where it has got much demand because of its flavour. It is known that this hot chilli is being largely used for the preparation of tuna fish which is one of the most favourite dishes of the people of Maldives.

Kerala is blessed with diverse climatic and soil conditions which have helped in the development of different landraces of *C chinense* having a wide range of variability. But so far there is no commercial variety available in this species in India. A critical estimate of genetic variability in the available germplasm is a pre requisite in a breeding programme for effective selection of superior genotypes and no systematic effort has been taken in this species so far.

Taking into consideration of all these aspects the present study was undertaken with the following objectives:

1. To genetically catalogue the available landraces in *C chinense*.

- 2 To identify superior genotypes based on yield quality and pest and disease resistance
- 3 To estimate the extent of available variability for important characters in *C. chinensis*
- 4 To study the extent of genetic divergence among the landraces and to group them into clusters based on genetic distance
- 5 To estimate the role of genetic contribution in the expression of each character and
- 6 To measure the degree and pattern of association between the characters

*Review of  
Literature*



## 2 REVIEW OF LITERATURE

*Capsicum chinense* is an economically important species of vegetable chili originated in the New World (Smith and Heiser 1957). The species which is noted for its biennial or perennial habit and for the highly pungent deep red coloured fruits are grown in the western hemisphere tropical South America Caribbean and South Central America (Loewenfeld and Back 1985). Scotch Bonnet and Habanero peppers the highly pungent cultivar classes of this crop are extremely popular in the United States (Fery and Thes 1997).

Eventhough considerable efforts have been made for the improvement of *C. annuum* the rich diversity in *C. chinense* has not received much attention.

The available literature on the variability in *Capsicum* spp are reviewed under the following subheads

- 2.1 Genetic cataloging
- 2.2 Variability
- 2.3 Heritability and genetic advance
- 2.4 Correlation studies
- 2.5 Path coefficient analysis
- 2.6 Selection index
- 2.7 Genetic divergence
- 2.8 Chemical constituents
- 2.9 Reaction to various noscoid disease

## 2.1 Genetic cataloguing

The genus *Capsicum* possess a wide variation in several morphological characters (Murthy and Bavajee 1982)

Ortiz and de la Flor (1990) evaluated five cultivated species of *Capsicum* and identified the principal characters differentiating them. Similarly Pradeepkumar (1990) grouped the accessions under *C. chinense* into three morphologically distinct groups based on corolla colour and calyx annular constriction.

Indra (1994) subjected eighty-two chilli genotypes to modern taxonomic treatments as suggested by IBPGR and assigned them into four *Capsicum* species namely *C. annum*, *C. frutescens*, *C. chinense* and *C. baccatum*.

In *C. frutescens* high variability for morphological characters was observed by Sheela (1998).

Based on plant growth habit, flower and fruit characters, Sin and Ahmad (1998) classified 160 accessions of chilli (*Capsicum* spp.) into four cultivated species.

A wide range of variation has been reported for various qualitative and quantitative characters in chilli (Verma *et al.* 1998 and Fatma 1999). Twenty-eight accessions of *C. chinense* were catalogued for morphological characters using the IBPGR descriptor list for *Capsicum* (Cheran 2000).

## 2.2 Variability

Genetic variability in the base population is a prerequisite for effective crop improvement. Considerable variation for several characters in

chilli was reported by Hiremath and Mathapatı (1977) Gopalakrishnan *et al* (1987) Kumar *et al* (1993) and Cherian (2000)

### 2 2 1 Morphological characters

High values for phenotypic and genotypic variances were observed for plant height by Arya and Sa n (1977) Ramalingam and Murugarajendran (1977) and Gopalakrishnan *et al* (1987) Ahmed *et al* (1990) and Sahoo *et al* (1990) revealed a low range of variation in plant height In *C. chinensis* Cherian (2000) obtained a phenotypic and genotypic coefficients of variation of 14.68 and 13.10 per cent respectively

In *C. annuum* Ramalingam and Murugarajendran (1977) noticed high values of phenotypic and genotypic variances for branches per plant whereas Ba *et al* (1987) observed low values Low phenotypic and genotypic variances for primary branches per plant was reported by Sahoo *et al* (1990) Similarly Varalakshmi and Har bab (1991) obtained low phenotypic and genotypic coefficients of variation for the character

In the case of plant spread high phenotypic and genotypic coefficients of variation was observed by Vijayalakshmi *et al* (1989) whereas Ahmed *et al* (1990) obtained low values Wide range of variation for plant spread was noticed by Sahoo *et al* (1990)

Low phenotypic and genotypic coefficients of variation was reported for days to first flowering (Arya and Sa n 1977 Varalakshmi and Har babu 1991 Cherian 2000)

Pollen fertility was observed to have a range of 95.6 – 96.0 per cent in sweet pepper (*C. annuum* L.) on the day of anthesis (Vijay *et al* 1979)

Pradeepkumar (1990) obtained a range of 37.24 per cent in *C. chacoense* to 94.11 per cent in *C. annuum*.

Arya and Sanil (1976) reported a moderate genotypic coefficient of variation for days to maturity in *C. annuum*. A high genetic variation for the character was observed by Kumar *et al.* (1993) and low variation by Cherian (2000).

## 2.2.2 Economic characters

### 2.2.2.1 Fruit and fruit yield

A wide range of variation in fruit and fruit yield per plant was reported by Arya and Sanil (1977), Hremath and Mathapat (1977), Ranalingan and Murugarajendran (1977), Gopalakrishnan *et al.* (1985), Ahmed *et al.* (1990), Sahoo *et al.* (1990) and Kataria (1997).

High phenotypic and genotypic coefficients of variation were observed by Singh and Brar (1979), Rajput *et al.* (1981), Nair *et al.* (1984), Jabeen *et al.* (1999) and Cherian (2000).

In *C. frutescens*, Sheela (1998) obtained high genotypic coefficient of variation for fruit yield.

### 2.2.2.2 Fruit characters

Low phenotypic and genotypic coefficients of variation were observed for fruit length, fruit girth and fruit weight by Vijayalakshmi *et al.* (1989), whereas Sheela (1998), Das and Choudhary (1999b), Fatima (1999) and Sreelathakumary (2000) reported high values.

Rajput *et al* (1981) and Na r *et al* (1984) observed high genotypic coefficients of variation in fruit weight. In the case of fruit growth a narrow genetic variability was reported by Gopalakrishnan *et al* (1987) and Munsh and Behera (2000)

Ped cell length exhibited a low range of variability in *C. n* (Choudhary *et al* 1985). On the other hand Ran (1996a) observed considerable variation for the character. In *C. chinense* a ped cell length range of 1.35 to 4.25 cm with 25.93 and 25.39 per cent of phenotypic and genotypic coefficients of variation respectively was reported by Cherian (2000)

### 2.2.2.3 Seed characters

Wide variability in seeds per fruit was observed by Arya and Sin (1976). V. Jayalakshmi *et al* (1989), Varalakshmi and Haribabu (1991) and Rani (1996b)

Sahoo *et al* (1990) reported a low range of variation in 100 seed weight whereas Rani (1996b) noticed a four fold variability in fruit seed weight. Rani *et al* (1996) revealed high genotypic coefficient of variation in 100 seed weight.

### 2.3 Heritability and genetic advance

Effectiveness of selection depends upon the heritability and genetic advance of the character studied.

Singh and Singh (1970) reported low values of heritability and genetic advance for most of the traits studied while Jabeen *et al* (1998) obtained high values for several characters.

High heritability has been reported for plant height in *C. annuum* (Singh and Brar 1979 Ghai and Thakur 1987) On the other hand low heritability was observed by Vijayalakshmi *et al* (1989)

In *C. annuum* branches per plant exhibited high heritability and genetic advance (Ramalingam and Murugarajendran 1977 Nar *et al* 1984) Low heritability and genetic advance for primary branches per plant was reported by Ran *et al* (1996)

Plant spread was reported to have low heritability and genetic advance by Ramakumar *et al* (1981) while Vijayalakshmi *et al* (1989) and Sahoo *et al* (1990) observed high values

Nar *et al* (1984) and Vijayalakshmi *et al* (1989) reported high heritability and low genetic advance for days to first flowering while moderate heritability along with low genetic advance for the character was observed by Rani *et al* (1996) Singh and Singh (1998) reported low heritability for days to first flowering

High heritability was observed for days to maturity in *C. annuum* (Ghai and Thakur 1987) Vallejo and Costa (1987) obtained a narrow sense heritability of 47 per cent for the character in *C. chinense*

Fruit and fruit yield per plant was reported to have high heritability and genetic advance (Arya and San 1977 Rajput *et al* 1981 Ramakumar *et al* 1981 Sahoo *et al* 1990 and Kumar *et al* 1993) Rani *et al* (1996) obtained heritability estimates of 99.3 per cent for fruits and 99.5 per cent for fruit yield Similar results were also reported by Das and Choudhary (1996b) and Jabeen *et al* (1999) in *C. annuum* and Cherian (2000) in *C. chinense*

Singh and Brar (1979) and Gopalakrishnan *et al* (1985) noticed moderate heritability for fruit yield

High heritability and genetic advance for several fruit characters was reported by Singh *et al* (1994) Pitchaimuthu and Pappiah (1995) Bhatt and Shah (1996) Ghildyal *et al* (1996) and Cherian (2000)

Choudhary *et al* (1985) revealed low heritability and genetic advance for fruit girth whereas Rani *et al* (1996) observed the same for fruit length Munshi and Behera (2000) reported moderate heritability and high genetic advance for fruit girth and fruit weight

In the case of seeds per fruit low heritability coupled with low genetic advance was reported by Ramakumar *et al* (1981)

Singh and Singh (1970) observed low heritability coupled with low genetic advance for 1000 seed weight whereas Mehra (1978) noticed high heritability for seed weight of 25 pods

#### 2.4 Correlation studies

A thorough knowledge of the relationship between yield and its component characters make crop improvement more effective

Pawade *et al* (1995) reported a strong positive correlation of fruits per plant with yield Sundaram and Irulappan (1998) concluded that fruits per plant is the primary yield component High positive association between yield and fruits per plant was also observed by Das and Choudhary (1999a) Aliyu *et al* (2000) and Cherian (2000)

Flowers produced was found to have a high positive correlation with yield (Suthanthirapandan and Sivasubramanian 1978) A high negative

correlation was observed between yield and days to first flowering (Warade *et al* 1996 Sreelathakumary 2000)

Arya and Sain (1976) found a negative association between yield and plant height whereas a positive correlation was observed by Dev and Arumugam (1999) and Ali *et al* (2000)

Ramakumar *et al* (1981) reported a strong correlation between plant height and plant spread. In *C. annuum* branches and plant spread were positively correlated with yield (Ghai and Thakur 1987 Rani 1995 Subashr and Natarajan 1999)

Fruit yield was observed to have a positive correlation with fruit weight, fruit length and fruit girth (Warade *et al* 1996). A strong association of yield with fruit weight and fruit circumference was also reported by Mishra *et al* (1998). Munshi *et al* (2000) noticed a non significant positive correlation of fruit length and fruit breadth with fruit and fruit yield.

Rani (1996d) observed high positive correlation of fruit weight with fruit diameter and seeds per fruit. Seeds per fruit was reported to have high correlation with yield (Chouvey *et al* 1986 Das *et al* 1989 Rani 1996b)

A positive correlation was observed between yield and seed weight of 25 pods by Mehra (1978) whereas Rani (1997) reported that yield has no correlation with 1000 seed weight.

In *C. frutescens* Sheela (1998) obtained a high positive correlation between harvests and fruit yield.



## 2.5 Path coefficient analysis

Rao *et al* (1974) concluded that in chilli the principal traits influencing green pod yield directly or indirectly were days to first flowering, days to pod maturity and pods per plant.

Seed weight of 25 pods was observed to be the major contributor to yield followed by days to first fruit set, locules per fruit and primary branches per plant (Mehra 1978).

Sundaram and Ranganathan (1978) reported that fruits per plant exerted the maximum positive direct effect on yield. The indirect effects of other characters through fruits per plant were consistently high.

Fruits, secondary branches, fruit weight, fruit circumference and duration exhibited positive direct effects on yield (Nair *et al* 1984).

Fruits per plant and fruit weight were reported to be the primary yield determinants in chilli (Rao and Chhonkar 1981, Depestre *et al* 1989, Pawade *et al* 1995, Das and Choudhary 1999a, Cherian 2000, Munshi *et al* 2000).

Chouvey *et al* (1986) obtained direct positive effects of seeds per fruit, fruit circumference and peduncle length on fruit yield.

In sweet pepper (*C. annuum* L.) fruits per plant and fruit size exhibited positive indirect effects, while branches per plant (Deka and Shadeque 1997).

Deshmukh *et al* (1997) reported that weight of 50 red ripe fruits exerted the highest direct positive effect on yield, followed by plant diameter and weight of 50 dried fruits.

Plant spread and fruits per plant recorded the highest positive direct effects on yield (Mishra *et al* 1998)

Dev and Arumugam (1999) noticed a negative direct effect of plant height on yield but influenced indirectly through fruits per plant, fruit shape index, secondary branches, capsaicin content and seeds per fruit.

Fatima (1999) found that fruit weight exhibited the highest positive direct effect on yield whereas Legesse *et al* (1999) observed that canopy width, leaf area, fruits per plant and pericarp thickness had positive direct effects on fruit yield.

In pepper (*C. annuum* L.) total dry weight exerted the maximum direct effect on yield followed by leaf area index, fruits and seeds per fruit (Aliyu *et al* 2000).

## 2.6 Selection index

Singh and Singh (1977) suggested that selection based on plant height, branches, days to first flowering, days to maturity, fruit length, fruit size and fruits per plant may be done to evolve high yielding lines in chilli. On the other hand, Mehra (1978) found that genetic advance through straight selection for yield per plant *per se* was higher than that calculated by discriminant function considering all combinations of component characters.

Fruits per plant and branches are the important characters that should be taken for selection in improvement programmes in chilli (Sundaram *et al* 1979).

Ramakumar *et al* (1981) opined that discriminant function based on fruit girth, fruits per plant and plant spread may be more efficient than straight selection.

Selection index involving yield per plant, fruit weight and fruits per plant was suitable to identify superior genotypes in *C. chinense* (Cherian 2000)

## 2.7 Genetic divergence

Singh and Singh (1976) grouped forty five strains of chilli into ten clusters based on the divergence pattern. The clustering pattern revealed that strains belonging to a particular geographical location generally tended to be in the same cluster.

Mehra (1978) studied genetic divergence in chilli using Mahalanobis  $D^2$  statistic and grouped 27 chilli varieties into nine gene constellations. Yield per plant was observed in contributing maximum towards divergence.

Sundaram *et al* (1980) could not observe any relation between genetic and genotypic diversity when they subjected 35 Indian and 15 exotic lines of *C. frutescens* to  $D^2$  analysis. Branches per plant and fruits per plant were the important characters contributing to genetic divergence.

In sweet pepper, six parents and the fifteen  $F_1$  hybrids formed seven clusters irrespective of the number of genotypes in each cluster (Gill *et al* 1982).

Varalakshmi and Harbabu (1991) grouped 32 genotypes of *C. annuum* based on ten characters into eleven gene constellations.

Indira (1994) assessed genetic diversity among chilli genotypes including *C. chinense* considering eight quantitative characters. Based on genetic distances, 71 genotypes were grouped into nine clusters during the first season and 72 genotypes into six clusters during the second season.

Roy and Sharma (1996) grouped 20 genotypes of chilli into seven clusters and observed that yield per plant primary branches per plant fruit girth and plant height were the highest contributors to total divergence

## 2.8 Chemical constituents of chilli

The important chemical constituents of chilli fruits include capsaicin oleoresin and ascorbic acid contents. But systematic studies on the physico-chemical qualities of *C. chinense* is very little.

### 2.8.1 Capsaicin

The pungent principle in chilli capsaicin ( $C_{18}H_{27}O_3N$ ) is a substituted benzylamine derivative (Narayanan *et al.* 1999).

About 90 per cent of capsaicin are concentrated in the placenta which has a capsaicin content of about 7 per cent (Sumathykutty and Mathew 1984). Teotia and Rana (1987) determined capsaicin by thin layer chromatography method and observed a range of 98.7 to 199.9 mg percentage. Tewari (1990) reported that capsaicin content can be improved by delaying the harvest to withering stage.

Indira (1994) classified genotypes with capsaicin in the range 1.0 to 1.5 per cent as highly pungent, 0.25 to 0.75 per cent as medium pungent and 0.11 to 0.25 per cent as less pungent.

Mani (1997) evaluated different *Capsicum* species and reported that *C. chinense* accessions CA 640 and CA 645 were having the highest pungency.

Mathur *et al.* (2000) conducted a survey to find out chilli with high capsaicin and dihydrocapsaicin. The Tezpur cult var of *C. frutescens* (Nagahar)

was found to contain the highest amounts of capsaicin and dihydrocapsaicin (4.28 and 1.42 per cent respectively) which is the hottest chilli known to date

Singh *et al* (2001) reported capsaicin contents of 2.15 per cent and 2.06 per cent with acetone and ethyl alcohol as solvents respectively

Nair *et al* (1984) observed high heritability and genetic advance for capsaicin. A negative correlation of capsaicin with fruit weight was reported by Jiang *et al* (1987) and Rani (1995)

The wide range of variation in capsaicin reported by several workers are summarized below

Reported by	Range of capsaicin (%)	Species
Pradeepkumar (1990)	0.42 - 2.54	<i>Capsicum</i> spp
Rani (1994)	0.056 - 1.81	<i>C. annuum</i> L.
Rani (1996c)	0.11 - 1.81	<i>C. annuum</i> L.
Mini (1997)	1.10 - 2.20	<i>Capsicum</i> spp
Sheela (1998)	0.43 - 1.70	<i>C. frutescens</i>
Cherian (2000)	0.82 - 1.85	<i>C. chinense</i>
Sreelathakumary (2000)	0.65 - 1.06	<i>Capsicum</i> spp

## 2.8.2 Oleoresin

Oleoresin consists of fixed oil, capsaicin, pigments, sugars and resin (Bajaj *et al* 1980). Pradeepkumar (1990) reported an oleoresin range from

18.70 per cent in *C. annuum* to 31.70 per cent in *C. chinense*. In *C. annuum* Papalkar *et al.* (1991) observed a range of 6.27 to 8.67 per cent.

Min (1997) reported a positive correlation of oleoresin with fruits per plant and negative correlation with earliness. Days to first flowering had the maximum direct effect on oleoresin yield. Oleoresin recovery was maximum at full ripe stage which was on par with turning stage.

Mandal *et al.* (1998) observed a higher oleoresin yield from the pulp of the fruit than the whole fruit. Mini *et al.* (1999) evaluated nine chilli genotypes under three different seasons and identified variety Arka Lohit as the highest oleoresin yielder (30.4 per cent). Genotypes were higher in oleoresin content during winter. Cheran (2000) observed a range of 9.0 to 25.75 per cent in *C. chinense*. Sreelathakumary (2000) obtained an oleoresin range of 12.40 per cent in *C. annuum* to 23.35 per cent in *C. chinense*. Singh *et al.* (2001) reported acetone as the best solvent for oleoresin extraction.

### 2.8.3 Ascorbic acid

The fruits of most *Capsicum* species contain high amounts of vitamin C upto 340 mg per 100 g when in fresh state (Anu and Peter 2000).

Ascorbic acid content were reported to have high heritability and genetic advance by Bhagyalakshmi *et al.* (1990), Kumar *et al.* (1993) and Rani *et al.* (1996).

Bajaj *et al.* (1980) observed an ascorbic acid range of 53.77 to 221.86 mg per 100 g and reported that the varietal variation also depended upon the maturity of the fruits. Maurya *et al.* (1984) reported a range of 158 to 171 mg per 100 g fresh fruit weight for ascorbic acid in *C. annuum*. A wide variation in ascorbic acid (58.73 to 193.1 mg per 100 g) was reported by

Rani (1994) Todorova *et al* (1997) observed a range of 147.1 to 343.8 mg per 100 g fresh fruit weight for ascorbic acid and reported a positive correlation with dry matter content and thickness of pericarp. Sreelathakumary (2000) reported a range of 92.74 to 97.01 mg per 100 g in *Chilli*.

## 2.9 Reaction towards mosaic disease of chilli

Mosaic is a serious viral disease found in chilli all over the country. It causes stunting of the chilli plant and reduction of leaves and fruits (Tewari 1983). Bidari and Reddy (1991) reported that mosaic incidence of *Capsicum* was widespread in commercially cultivated fields in Karnataka with disease incidence from 11.8 to 94.8 per cent.

### 2.9.1 Causal organism

A number of viruses have been associated with mosaic disease of chilli in India. Chief among them are cucumber mosaic virus (CMV), potato virus X (PVX), potato virus Y (PVY), chilli mosaic virus, tobacco mosaic virus (TMV) and tobacco leaf curl virus (TLCV).

The viruses are transmitted by aphid vectors, namely *Aphis gossypii*, *A. evonymi* and *Myzus persicae* and is not seed transmitted (Chattopadhyay and Satyabrata 1990).

### 2.9.2 Physiological effects

Viruses cause a lot of abnormalities and physiological imbalances leading to a drastic reduction in the yield (Jayaraj and Ramakrishnan 1961). Abnormal shedding of flower buds, failure of anthers to mature which are partially filled with non-functional pollen are reported by Jayaraj and Ramakrishnan (1961) and

Awasthi and Singh (1974) Chauhan (1980) observed 80 to 90 per cent sterility in virus infected chilli plants. In egg plant Gupta *et al* (1988) observed 55.68 to 87.45 per cent pollen sterility due to egg plant mild mosaic virus.

### 2.9.3 Source of resistance

A good number of sources of resistance to different viruses have been located by various scientists.

Singh (1973) evaluated 105 varieties of chilli involving five species and reported that the varieties Pur Red, Puri Orange G2, Kondverum and Suryamukhi were resistant to mosaic. Konai and Nariani (1980) reported that varieties Pant C1 and Pant C2 were tolerant to CMV, TMV, PVX and TLCV.

A local perennial chilli (IC 31339) has been reported to be immune against both CMV and PVX and tolerant to TMV and TLCV. It is also resistant to chilli mosaic virus (Tewari, 1983).

Thakur *et al* (1985) screened 51 varieties of pepper against PVY and found two resistant ones. Four varieties were moderately resistant and the rest were either susceptible or highly susceptible.

Sixty one genotypes of chilli were screened for resistance to PVY and identified one culture (Acc. No. 426) as moderately resistant (Anandam, 1992).

Kaloo (1994) reported a *C. chinense* accession PI 159236 resistant to TMV. Singh and Singh (1998) and Acharyya (1999) reported that variety Punjab Lal is resistant to CMV.

In a screening trial involving 53 accessions of chilli (*C. annuum* L.) Fatma (1999) observed that nine accessions were found to be resistant, twelve were moderately resistant and the remaining were susceptible.



*Materials and  
Methods*

### 3 MATERIALS AND METHODS

The present investigation was carried out in the Department of Olericulture College of Agriculture Vellayani during September 2000 May 2001. The area is situated at 8 5° N latitude 76 9° E longitude at an altitude of 29 0 m above mean sea level. Experimental site has a lateritic red loam soil with a pH of 5 2. The area enjoys a warm humid tropical climate.

The study consisted of the following experiments:

#### 3 1 Genetic cataloguing of *Capsicum chinense*

#### 3 2 Variability in *C chinense*

#### 3 1 Genetic cataloguing of *C chinense*

The basic material for the study consisted of 32 diverse accessions of *C chinense* collected through survey and correspondence. The details of the accessions and their sources are presented in Table 1. The descriptor developed by IBPGR (1995) for *Capsicum* was used for cataloguing (Table 2).

#### 3 2 Variability in *C chinense*

#### 3 2 1 Experimental materials and methods

Thirty two accessions of *C chinense* were grown during September 2000 to May 2001 to identify superior genotypes with yield, quality and reaction towards the incidence of pests and diseases.

**Table 1** Particulars of accessions of *C chinense* used in the study and their sources

Sl No	Accession number	Source
1	CC 1	Vellayani Thiruvananthapuram
2	CC 2	Anchal Kollam
3	CC 3	Neyyattinkara Thiruvananthapuram
4	CC 4	Veliyam Kollam
5	CC 5	Venganoor Thiruvananthapuram
6	CC 6	Nemom Th ruvananthapuram
7	CC 7	Vithura Th ruvananthapuram
8	CC 8	Vithura Th ruvananthapuram
9	CC 9	Indian Cardamom Research Institute Saklespur Karnataka
10	CC 10	Paudikkonam Th ruvananthapuram
11	CC 11	V thura Thiruvananthapuram
12	CC 12	Kumarapuram Thiruvananthapuram
13	CC 13	Vithura Thiruvananthapuram
14	CC 14	Neyyattinkara Thiruvananthapuram
15	CC 15	V lavoorkal Thiruvananthapuram
16	CC 16	Venganoor Th ruvananthapuram
17	CC 17	Paudikkonam Th ruvananthapuram
18	CC 18	Neyyatt nkara Thiruvananthapuram
19	CC 19	Venganoor Th ruvananthapuram
20	CC 20	Sreekaryam Th ruvananthapuram
21	CC 21	Sreekaryam Thiruvananthapuram
22	CC 22	Vithura Thiruvananthapuram
23	CC 23	Nemom Thiruvananthapuram
24	CC 24	V thura Th ruvananthapuram
25	CC 25	Vithura Thiruvananthapuram
26	CC 26	V thura Thiruvananthapuram
27	CC 27	V thura Thiruvananthapuram
28	CC 28	Pothankode Thiruvananthapuram
29	CC 29	Pothankode Th ruvananthapuram
30	CC 30	Nemom Th ruvananthapuram
31	CC 31	Nemom Th ruvananthapuram
32	CC 32	Neyyatt nkara Th ru vananthapuram

Table 2 Genetic cataloguing of *C chinense*

<b>1</b>	<b>Vegetative characters</b>	
1 1	Hypocotyl colour	White / green / purple
2	Stem pubescence	Sparse / intermediate / dense
3	Leaf colour	Light green / green / dark green / light purple / purple
4	Leaf shape	Deltoid / ovate / lanceolate / elong deltoid
5	Stem colour	Green / green with purple stripes / purple
6	Nodal anthocyanin	Green / light purple / purple / dark purple
7	Plant growth habit	Prostrate / intermediate / erect
8	Stem length (cm)	Light to first bifurcation
<b>2</b>	<b>Inflorescence characters</b>	
2 1	Number of flowers per axil	– One / two / three or more
2	Flower position	– Pendant / intermediate / erect
3	Corolla colour	– White / light yellow / yellow green / purple
4	Corolla spot	Absent / present
5	Corolla shape	Rotate / campanulate
6	Anther colour	White / yellow / pale blue / blue / purple
7	Filament colour	White / yellow / green / blue / light purple / purple
8	Stigma exertion	Inserted / same level / exerted
9	Calyx pigmentation	– Absent / present
10	Calyx margin	Entire / intermediate / dentate
11	Calyx annular constriction	Absent / present

Table 2 Contd

3 Fruit and seed characters		
3.1	Anthocyanin spots or stripes	Absent / present
2	Fruit colour at immature stage	White / yellow / green / orange / purple / deep purple
3	Fruit colour at mature stage	White / lemon yellow / orange yellow / orange / light red / red / dark red / purple
4	Fruit shape	- Elongate / almost round / triangular / campanulate / blocky
5	Fruit shape at peduncle attachment	- Acute / obtuse / truncate / cordate / lobate
6	Neck at base of fruit	Absent / present
7	Fruit shape at blossom end	- Pointed / blunt / sunken / sunken and pointed
8	Fruit cross sectional corrugation	Slightly corrugated / intermediate / corrugated
9	Number of locules	Two / three (as percentage of each category)
10	Seed colour	- Straw / brown / black

Statistical details were as furnished below

Design	RBD
Replications	3
Treatments	32 accessions
Plot size	6.75 m <sup>2</sup>
Spacing	75 x 60 cm
Number of plants per plot	15

The crops were raised as per package of practices recommendations of Kerala Agricultural University (KAU 1996). No insecticide or fungicide was applied on the plants during the course of experimentation to observe the reaction of the accessions towards pests and diseases.

### 3.2.2 Observations

Five plants were randomly selected per accession per replication for taking observations and the mean worked out. For recording observations on fruit characters, five fruits at fully mature green stage were selected at random from each accession in each replication. Observations on the following characters were recorded.

#### 3.2.2.1 Plant characters

##### (a) Plant height (cm)

Measured from the ground level to the top of the plant at the time of final harvest.

**(b) Primary branches per plant**

Branches arising from the main stem were counted

**(c) Plant spread / Plant circumference (cm)**

Expressed as  $\pi d$  where  $d$  is the average of North South and East West spread of the plant taken across the bush in North South and East West direction respectively

**3 2 2 2 Flowering characters****(a) Days to first flowering**

Number of days from transplanting until 50 per cent of plants in each accession have at least one open flower

**(b) Pollen viability**

Pollen viability was determined by the acetocarmine staining technique. Anthers about to dehisce were collected separately from each accession and the pollen grains were mounted in a drop of acetocarmine glycerine mixture (1:1). The slides were kept for about 30 minutes to allow pollen grains to take stain properly before examining under a microscope. Pollen viability was studied by counting the well filled and stained pollen grains. An average of 100 pollen were counted in different microscopic fields in each accession. Unfilled and unstained pollen grains were considered as sterile. Pollen viability was calculated as follows

$$\% \text{ of pollen viability} = \frac{\text{Number of well filled and stained pollen grains}}{\text{Total number of pollen grains counted}} \times 100$$

**(c) Days to maturity**

Number of days from fruit set until the fruit shows signs of ripening was observed

**3.2.2.3 Fruit and yield characters****(a) Fruits per plant**

Total number of fruits per plant was observed

**(b) Fruit length (cm)**

Distance between pedicel attachment and fruit apex

**(c) Pedicel length (cm)**

Distance between the point of attachment of pedicel with the stem and the fruit

**(d) Fruit girth (cm)**

Measured using twine and scale at the maximum width of the fruit

**(e) Fruit weight (g)**

Average of five fruits weight

**(f) Seeds per fruit**

Seeds per fruit were counted in five fruits and average was taken

**(g) 1000 seed weight**

The dry weight of randomly selected 1000 seeds were taken

**(h) Yield per plant (g)**

Weight of fruits harvested from each plant was recorded



**(i) Yield per harvest (g)**

Weight of fruits harvested from each plant was recorded for each harvest and the average taken

**(j) Number of harvests**

Total number of harvests from each plant was recorded

**3 2 2 4 Quality characters****(a) Capsaicin (%)**

Capsaicin content of different accessions were determined by Folin-Dennis method. The pungent principle reacts with Folin-Dennis reagent to give a blue coloured complex which is estimated colorimetrically (Mathew *et al* 1971)

**Reagents****(i) Folin-Dennis reagent**

Refluxed 750 ml distilled water, 100 g sodium tungstate, 20 g phosphomolybdic acid and 50 ml phosphoric acid for two hours. Cooled and diluted to 1000 ml with distilled water.

**(ii) 25 % aqueous sodium carbonate solution****(iii) Acetone****Procedure**

The fruits harvested at red ripe stage were dried in a hot air oven at 50°C and powdered finely in a mixer grinder. Five hundred mg each of the sample was weighed into test tubes. Added 10 ml acetone to it and kept

overnight Aliquots of 1 ml were pipetted into 100 ml conical flasks added 25 ml of Folin Dennis reagent and allowed to stand for 30 minutes Added 25 ml of freshly prepared sodium carbonate solution and shook vigorously The volume was made up to 100 ml with distilled water and the optical density was determined after 30 minutes at 725 nm against reagent blank (1 ml acetone + 25 ml Folin Dennis reagent + 25 ml aqueous sodium carbonate solution) using a UV spectrophotometer

To determine the EI per cent value for pure capsaicin a stock solution of standard capsaicin (200  $\mu\text{g ml}^{-1}$ ) was prepared by dissolving 20 mg in 100 ml acetone From this a series of solutions of different concentrations were prepared and their optical density measured at 725 nm Standard graph was prepared and calculated the content of capsaicin in the samples

#### (b) Oleoresin (%)

Oleoresin in chilli was extracted in a Soxhlet's apparatus using solvent acetone (Sadasvam and Manikam 1992)

#### Procedure

Chilli fruits harvested at red ripe stage were dried in a hot air oven at 50°C powdered finely in a mixer grinder Two g of chilli powder was weighed and packed in filter paper and placed in Soxhlet's apparatus Two hundred ml of acetone was taken in the round bottom flask of the apparatus and heated in a water bath The temperature was maintained at the boiling point of solvent After complete extraction (7 to 8 h) the solvent was evaporated to dryness under vacuum

Yield of oleoresin on dry weight basis was calculated using the formula

$$\text{Oleoresin (\%)} = \frac{\text{Weight of oleoresin}}{\text{Weight of sample}} \times 100$$

**(c) Ascorbic acid (mg per 100 g fresh fruit weight)**

Ascorbic acid content of fruits at red ripe stage was estimated by 2,6-dichlorophenol indophenol dye method (Sadasivam and Manickam 1992)

**Reagents**

(i) Oxalic acid (4%)

(ii) Ascorbic acid standard

Prepared a stock solution by dissolving 100 mg of ascorbic acid in 100 ml of four per cent oxalic acid. Diluted 10 ml of the stock solution to 100 ml with four per cent oxalic acid to get working standard solution

(iii) 2,6-dichlorophenol indophenol dye

Weighed 42 mg sodium bicarbonate into a small volume of distilled water. Dissolved 52 mg 2,6-dichlorophenol indophenol in it and made up to 200 ml with distilled water

**Procedure**

Pipetted out 5 ml of the working standard solution into a 100 ml conical flask and added 10 ml of four per cent oxalic acid. Titrated it against the dye (V ml). Endpoint is the appearance of pink colour which persisted for at least five seconds

Five g of fresh fruit was extracted in an acid medium (4 % oxal c acid) and titrated as above aga nst the dye solut on to a p nk colour ( $V_2$  ml) Ascorbic ac d content of the sample was calculated us ng the formula

$$\text{Amount of ascorb c acid /100 g sample} = \frac{0.5 \times V_2 \times 100}{V \times 5 \times \text{we ght of sample}} \times 100$$

### 3 2 2 5 Reaction towards major pests and diseases

No scor ng was done for pests since there was no major pest nc de ce in the crop Mosaic was the only disease observed and hence scor ng based on visual observations was done for mosaic incidence

#### Scoring for chilli mosaic incidence

The rat ng scale g ven by Rajamony *et al* (1990) n melon was used for scoring with minor modif cations Th s was done according to the character stic symptom of each observat onal plant (Table 3 and Plates 1 to 4)

**Table 3 Scoring for chilli mosaic disease**

Rating scale	Symptom	Category
0	No symptom	H ghly res stant
1	Very l ght mottl ng of green colour	Resistant
2	Prom nent mottl ng of leaves	Moderately res stant
3	Mottl ng covers the entire leaf lamina	Susceptible
4	Stunt ng of the plant Reduct on in leaf size with characterist c rat tailing and fork ng at the apex	Highly suscept ble

The individual plant score was utl ized to work out the Severity Index or Vulnerab lity Index (V I) so as to measure the res stance

**Plate 1**



**Plate 3**



**Plate 2**



**Plate 4**



The index was calculated using an equation adopted by Silbernagel and Jafri (1974) for measuring resistance in snap bean (*Phaseolus vulgaris*) to beet curly top virus and modified later by Bos (1982)

$$\text{Vulnerability Index (VI)} = \frac{(0n_0 + 1n_1 + 2n_2 + 3n_3 + 4n_4)}{n(n_c - 1)} \times 100$$

where

$n_0$  -  $n_4$  = number of plants in category 0 - 4 respectively

$n$  = total number of plants

$n_c$  = total number of categories = 5

The vulnerability index was used to classify the genotypes into different categories

Sl No	Vulnerability Index	Category
1	0 - 20	Highly resistant
2	21 - 40	Resistant
3	41 - 60	Moderately resistant
4	61 - 80	Susceptible
5	81 - 100	Highly susceptible

### 3.2.3 Statistical analysis

3.2.3.1 Analysis of variance (ANOVA) and covariance (ANCOVA) for Randomized Block Design (RBD) in respect of the various characters was done (Panse and Sukhatme 1967)

3.2.3.2 Mean The mean of the character  $\bar{X}$  was worked out

3.2.3.3 Variability components (phenotypic and genotypic) for different characters was estimated as suggested by Kempthorne (1977)

(a) The variance and covariance components were calculated as per the following formulae

For the character X

$$\begin{aligned} \text{Environmental variance } \sigma_e^2 &= \text{MSE} \\ \text{Genotypic variance } \sigma_g^2 &= \frac{\text{MST} - \text{MSE}}{r} \\ \text{Phenotypic variance } \sigma_p^2 &= \sigma_g^2 + \sigma_e^2 \end{aligned}$$

where MST and MSE are respectively the mean sum of squares for treatment and error from ANOVA and r the number of replications

For two characters X and X

$$\begin{aligned} \text{Environmental covariance } \sigma_e &= \text{MSPE} \\ \text{Genotypic covariance } \sigma_g &= \frac{\text{MSPT} - \text{MSPE}}{r} \\ \text{Phenotypic variance } \sigma_p &= \sigma_g + \sigma_e \end{aligned}$$

where MSPT and MSPE are respectively the mean sum of products between the  $i^{\text{th}}$  and  $j^{\text{th}}$  characters for genotype and environment respectively from Analysis of Covariance (ANCOVA)

The genotypes were classified into poor, medium and better categories with respect to each character as follows

Definition	Category
Less than mean $- SE_m$	Poor
Between mean $\pm SE_m$	Medium
More than mean $+ SE_m$	Better

where  $SE_m$  is the standard error of the mean for each character

$$SE_m = \sqrt{\frac{MSE}{r}}$$

For the character days to first flowering the reverse order is being followed for categorization

#### (b) Coefficient of variation

Variability that existed in the population for various characters were apportioned using the estimates of coefficient of variation (Singh and Chaudhary 1985)

For the character X

$$\text{Phenotypic coefficient of variation PCV} = \frac{\sigma_p}{\bar{X}} \times 100$$

$$\text{Genotypic coefficient of variation GCV} = \frac{\sigma_g}{\bar{X}} \times 100$$

$$\text{Environmental coefficient of variation ECV} = \frac{\sigma_e}{\bar{X}} \times 100$$

where  $\sigma_p$ ,  $\sigma_g$  and  $\sigma_e$  are respectively the phenotypic, genotypic and environmental standard deviations with respect to each character



### 3 2 3 4 Heritability

Hanson *et al* (1956) proposed the mathematical relationship of various estimates on computation of heritability which is usually expressed as percentage

$$\text{Heritability (broad sense) } H^2 = \frac{\sigma_g^2}{\sigma_p^2} \times 100$$

The range of heritability was categorized as suggested by Robinson *et al* (1949) as follows

Definition	Category
0 - 30 per cent	Low
31 - 60 per cent	Medium
61 per cent and above	High

### 3 2 3 5 Genetic advance

Genetic advance as percentage over mean was calculated as per the formula given by Lush (1949) and Johnson *et al* (1955)

$$\text{Genetic advance } GA = \frac{kH^2 \sigma_p}{\bar{X}} \times 100$$

where  $H^2$  heritability in broad sense

$\sigma_p$  phenotypic standard deviation

$k$  selection differential which is 2.06 in case of 5% selection in large samples (Miller *et al* 1958 and Allard 1960)

Genetic advance was categorized according to Robinson *et al* (1949) as follows

Definition	Category
Less than 20 per cent	Low
Greater than 20 per cent	High

### 3 2 3 6 Correlation analysis

Phenotypic, genotypic and environmental correlation coefficients were worked out according to the procedure suggested by Singh and Choudhary (1985)

### 3 2 3 7 Path analysis

The direct and indirect effects of yield contributing factors were estimated through path analysis technique (Wright 1954)

### 3 2 3 8 Mahalanobis D<sup>2</sup> analysis

Genetic divergence was studied based on ten characters taken together using Mahalanobis D<sup>2</sup> statistic as described by Rao (1952). The genotypes were clustered by Tocher's method.

### 3 2 3 9 Selection index

The various genotypes were discriminated based on ten characters using the selection index developed by Smith (1936) using the discriminant function of Fisher (1936). The selection index is described by the function

$$I = b_0 X_0 + b_1 X_1 + b_2 X_2 + \dots + b_k X_k$$

The function  $H = a_1 G_1 + a_2 G_2 + \dots + a_k G_k$  describes the merit of a plant where  $X_1, X_2, \dots, X_k$  are the phenotypic values and  $G_1, G_2, \dots, G_k$  are the genotypic values of the plant with respect to the characters  $X_1, X_2, \dots, X_k$ .  $H$  denotes the genetic worth of the plant. The economic worth assigned to each character is assumed to be equal to unity i.e.  $a_1 = a_2 = \dots = a_k = 1$ . The regression coefficients  $b_1, b_2, \dots, b_k$  are estimated in such a way that the correlation between  $H$  and  $I$  is maximum.

i.e.  $b = P^{-1} G a$  where  $P$  and  $G$  are the phenotypic and genotypic variance covariance matrices respectively. Based on the  $b$  estimates and the mean values for the ten characters with respect to each accession scores were calculated and the accessions were ranked.

*Results*

## 4 RESULTS

Experimental data recorded during the course of investigation were subjected to statistical analysis and are presented under the following heads

### 4.1 Genetic cataloguing in *Capsicum chinense*

#### 4.2 Variability in *C. chinense*

#### 4.1 Genetic cataloguing in *C. chinense*

Thirty two accessions of *C. chinense* were genetically catalogued based on the descriptor (IBPGR 1995). Morphological characters like vegetative (Table 4), inflorescence (Table 5) and fruit and seed characters (Table 6) were recorded and accessions were catalogued.

All the accessions had green to purple hypocotyl and stem colour. The leaf colour varied between light green to light purple with deltoid leaf shape. Most of the accessions had sparse stem pubescence with green to dark purple nodal anthocyanin. Plant growth habit were either erect or compact with a stem length of 21.00 to 55.75 cm.

Flowers per axil were either two or three with erect to pendant position. Corolla colour ranged from light yellow to purple with pale blue to purple anther and white to light purple filament. All the accessions had rotate corolla without corolla spot. All the accessions were identical with a prominent annular constriction at calyx. Calyx pigmentation was absent in all the

Table 4 Vegetative characters in *C. chinense* accessions

Accession number	Hypocotyl colour	Stem pubescence	Leaf colour	Stem colour	Nodal anthocyanin	Plant growth habit	Stem length (cm)
CC 1	Green	Sparse	Lght green	Green	Green	Erect	46 00
CC 2	Green	Sparse	Lgh green	Green v th purple str pes	Lght purple	Compact	26 60
CC 3	Green	Spa se	Lght g een	Green v th purple s r pes	Green	Erect	55 75
CC 4	Green	Spa se	Lght green	G een	Green	Erect	44 00
CC 5	Purple	Sparse	Lght purple	Purple	Dark purple	Erect	40 50
CC 6	Green	Spa se	Lght green	Green	Lght purple	Erct	23 33
CC 7	Green	Spa se	Lght green	G een v th purple str pes	L gl pu ple	Compac	30 50
CC 8	Green	Spa se	G een	G een	Lght purp e	E ect	25 17
CC 9	Green	Sparse	Lght g en	G een	Lght purp e	Erect	28 00
CC 10	Green	In e ned a e	Lght g een	G een	L gh pu ple	Er c	24 67
CC 11	Green	Sparse	Lgh g een	G ce l purple str pes	L gh purple	Compac	30 00
CC 12	Green	Sparse	Lght green	G een v h pu ple s r pes	Purple	Erect	32 25
CC 13	Green	Spa se	Lght green	G een	Green	Er ct	35 00
CC 14	Green	Spa se	Lght green	G een w h pu ple str pes	L gh pu ple	Erect	34 00
CC 15	Green	l te ed a c	L gl t green	G een v th purple s r pes	Green	Erect	27 00
CC 16	Green	Sparse	Lght g een	Green	Lght purple	Compact	34 67
CC 17	Green	Spa se	Lgh green	G een h purple str pes	Green	Erect	24 00
CC 18	Green	Sparse	lght g een	G een w th purp e str pes	Lght pu p e	Erect	28 00
CC 19	Green	Spa se	Lght green	G een v th purple str pes	Green	Compact	33 33
CC 20	Green	Spa se	Lght g een	G een w th purple s r pes	Lght pu ple	Erect	28 50
CC 21	Green	Spa se	Lght green	Green th purple str pes	Purple	E ect	26 50
CC 22	Green	Spa se	Lgh green	G een	Green	Compac	27 00
CC 23	Green	Sparse	Lght green	Green	Green	Compact	35 50
CC 24	Green	Sparse	Lgh green	G een v h pu ple st pes	Lght pu p e	E ect	23 25
CC 25	Purple	Sparse	Lght purple	Purple	Dark purple	Erect	26 00
CC 26	Green	nte n cd ate	Lght g een	G een v h purple str pes	Green	Erec	33 25
CC 27	Green	Sparse	Lght green	G een	Lght purp e	Erect	31 00
CC 28	G een	Spa se	Lght g een	G een w tl purple str pes	Green	Compact	25 67
CC 29	Green	Intermed a e	Lgh green	G een	G een	Compact	21 00
CC 30	Green	Spa se	Lght g een	G een	G een	Compact	35 00
CC 31	Green	Spa se	Lght g een	G e n v h pu ple str pes	G een	Compac	33 75
CC 32	Green	Spa se	l gh green	G e n	Green	E cc	28 00

Table 5 Inflorescence characters in *C chinense* accessions

Accession Number	Number of florets per a l	Flower position	Corolla colour	Anther colour	Filament colour	Stigma exsertion (out of 10)			Calyx pigmentation	Calyx margin
						Exserted	Same level	Inserted		
CC 1	Tavo	Erect	Light yellow	Pale blue	White	6	4		Absent	Intermedate
CC 2	To	Erect	Light yellow	Light purple	White	8	2		Absent	Intermedate
CC 3	Three	Erect	Light yellow	Pale blue	White	6		1	Absent	Entire
CC 4	Three	Erect	Light yellow	Light purple	White	10			Absent	Intermedate
CC 5	Three	Intermediate	Purple	Purple	Light purple	5	5		Present	Intermedate
CC 6	To	Intermediate	Light yellow	Pale blue	White	4	6		Absent	Intermedate
CC 7	Three	Intermediate	Light yellow	Light purple	White	7	3		Absent	Intermedate
CC 8	To	Intermediate	Light yellow	Light purple	White	6			Absent	Intermedate
CC 9	Three	Intermediate	Light yellow	Light purple	White	4	5	1	Absent	Intermedate
CC 10	To	Erect	Light yellow	Light purple	White	9		1	Absent	Intermedate
CC 11	Three	Erect	Light yellow	Light purple	White	10			Absent	Intermedate
CC 12	Tavo	Pendant	Light yellow	Pale blue	White	9	1		Absent	Intermedate
CC 13	To	Erect	Light yellow	Pale blue	White	10			Absent	Intermedate
CC 14	To	Intermediate	Light yellow	Light purple	White	8	2		Absent	Intermedate
CC 15	To	Pendant	Light yellow	Pale blue	White	9	1		Absent	Intermedate
CC 16	To	Erect	Light yellow	Light purple	White	5	4	3	Absent	Intermedate
CC 17	To	Intermediate	Light yellow	Light purple	White	7	2	1	Absent	Intermedate
CC 18	Three	Erect	Light yellow	Light purple	White	6	4		Absent	Intermedate
CC 19	Three	Intermediate	Light yellow	Pale blue	White	5	4	1	Absent	Intermedate
CC 20	Three	Intermedate	Light yellow	Light purple	White	6	5	1	Absent	Intermedate
CC 21	Three	Erect	Light yellow	Light purple	White	7			Absent	Intermedate
CC 22	Three	Intermediate	Light yellow	Light purple	White	10			Absent	Intermedate
CC 23	Three	Intermediate	Light yellow	Light purple	White	7	2	1	Absent	Intermedate
CC 24	To	Pendant	Light yellow	Light purple	White	8	1	1	Absent	Intermedate
CC 25	Three	Intermediate	Purple	Purple	Light purple	7	5	3	Present	Intermedate
CC 26	To	Intermediate	Light yellow	Pale blue	White	8		1	Absent	Intermedate
CC 27	Three	Pendant	Light yellow	Light purple	White	5	6	1	Absent	Intermedate
CC 28	To	Erect	Light yellow	Light purple	White	9	1		Absent	Intermedate
CC 29	Three	Intermediate	Light yellow	Light purple	White	8	2		Absent	Intermedate
CC 30	To	Erect	Light yellow	Pale blue	White	9	1		Absent	Intermedate
CC 31	Three	Erect	Light yellow	Light purple	White	10			Absent	Entire
CC 32	Three	Intermediate	Light yellow	Light purple	White	6		1	Absent	Intermedate

Table 6 Fruit and seed characters in *C chinense* accessions

Accession Number	Anthocyanin spots or spots on fruit	Fruit colour at immature stage	Fruit colour at mature stage	Fruit shape	Fruit shape at pedicel attachment	Neck at base of fruit	Fruit shape at blossom end	Fruit cross sectional corrugation	Seed colour
CC 1	Absent	Green	Red	Elongate	Acute	Absent	Po nted	Corrugated	Straw
CC 2	Absent	Green	Red	Campanulate	Truncate	Absent	Po nted	In c med ate	Straw
CC 3	Absent	Green	Red	Elongate	Acute	Present	Po nted	Corrugated	Straw
CC 4	Absent	Green	Dark red	Elongate	Acute	Present	Po nted	I n e n d a e	Straw
CC 5	Present	Green	Dark red	Campanulate	Acute	Absent	Po nted	I n e n d a e	Straw
CC 6	Absent	Green	Dark red	Angular	Truncate	Absent	Po nted	Co rugated	Straw
CC 7	Absent	Green	Dark red	Campanulate	Truncate	Present	Po nted	Co rugated	Straw
CC 8	Absent	Green	Red	Campanulate	Truncate	Absent	Blunt	Co rugated	Straw
CC 9	Absent	Green	Le non yel ow	Campanulate	Cordate	Absent	Blunt	Co rugated	Straw
CC 10	Absent	Green	Red	Angular	Cordate	Absent	Blunt	Corrugated	Straw
CC 11	Absent	Green	Dark red	Elongate	Acute	Present	Po nted	Corrugated	Straw
CC 12	Absent	Green	Red	Angular	Truncate	Absent	Po nted	Co rugated	Straw
CC 13	Absent	Green	Dark red	Elongate	Acute	Absent	Po nted	Corrugated	Straw
CC 14	Absent	Green	Red	Campanulate	Truncate	Absent	Blunt	Co rugated	Straw
CC 15	Absent	Green	Red	Blocky	Truncate	Present	Blunt	Corrugated	Straw
CC 16	Absent	Green	Red	Angular	Truncate	Absent	Po nted	Corrugated	Straw
CC 17	Absent	Green	Red	Angular	Truncate	Absent	Po nted	Corrugated	Straw
CC 18	Absent	Green	Red	Campanulate	Truncate	Absent	Blunt	Corrugated	Straw
CC 19	Absent	Green	Red	Elongate	Acute	Absent	Blunt	In ermed ate	Straw
CC 20	Absent	Green	Red	Campanulate	Truncate	Absent	Po nted	Corrugated	Straw
CC 21	Absent	Green	Red	Campanulate	Truncate	Absent	Po nted	Co rugated	Straw
CC 22	Absent	Green	Red	Blocky	Truncate	Absent	Blunt	Corrugated	Straw
CC 23	Absent	Green	Red	Almost round	Truncate	Absent	Blunt	Sl gh y corrugated	Straw
CC 24	Absent	Green	Red	Blocky	Truncate	Absent	Po nted	Corrugated	Straw
CC 25	Absent	Deep purple	Red	Campanulate	Acute	Absent	Po nted	Corrugated	Straw
CC 26	Absent	Green	Red	Campanulate	Truncate	Absent	Po nted	Co rugated	Straw
CC 27	Absent	Green	Lemon ye ow	Blocky	Cordate	Absent	Po nted	Co rugated	Straw
CC 28	Absent	Green	Red	Angular	Truncate	Absent	Blunt	Corrugated	Straw
CC 29	Absent	Green	Red	Blocky	Truncate	Absent	Blunt	Co rugated	Straw
CC 30	Absent	Green	Dark red	Angular	Truncate	Absent	Blunt	Corrugated	Straw
CC 31	Absent	Green	Red	Angular	Truncate	Absent	Blunt	Co rugated	Straw
CC 32	Absent	Green	Dark Red	Campanulate	Cordate	Absent	Po nted	Co rugated	Straw



accessions except CC 5 and CC 25 while calyx margin was intermediate in most of the accessions

Fruit colour at immature stage remained green except in CC 25 which had deep purple colour. Except for CC 5 all the accessions had no anthocyanin spots or stripes on fruit. At mature stage CC 9 and CC 27 recorded lemon yellow colour while the rest of the accessions had red to dark red colour.

Fruit shape varied from elongate to blocky with acute truncate or cordate shape at pedicel attachment. Neck at base of fruit was absent in most of the accessions with either pointed or blunt blossom end. Slightly corrugated to corrugated fruit cross section was noticed with two to three locules per fruit. Seed colour was found to be straw in all the accessions.

## 4.2 Variability in *C. chinense*

### 4.2.1 Mean performance

Analysis of variance showed significant differences among the accessions for all the characters studied (Table 7). The mean values of 32 accessions for different characters are presented in Table 8.

#### **Plant height**

There was significant difference among the accessions for plant height. It ranged from 61.33 to 133.33 cm with an overall mean of 98.69 cm. CC 27 was the tallest with a height of 133.33 cm which was on par with CC 7.

**Table 7 Analysis of variance for 20 characters in 32 accessions of *C. chinense* (Mean squares)**

Source	df	Plant height	Primary branches per plant	Plant spread	Days to first flowering	Pollen viability	Days to maturity	Fruits per plant
Replication	2	61.19	0.01	1005.50	22.20	5.41	13.20**	687.38
Genotype	31	1083.10**	4.95**	5945.94**	101.42**	839.88**	42.17**	68853.04**
Error	62	48.07	1.68	686.32	7.09	2.63	1.29	272.96

Source	df	Fruit length	Ped cell length	Fruit g r t l	Fruit weight	Seeds per fruit	1000 seed weight	Yield per plant
Replication	2	0.002	0.03	0.01	0.02	7.95	0.01	3149.00*
Genotype	31	3.11**	1.52**	6.71**	8.31**	438.77**	1.06**	474820.90**
Error	62	0.07	0.02	0.03	0.06	4.41	0.03	987.52

Source	df	Yield per harvest	Number of harvests	Capsaicin	Oleoresin	Ascorbic acid	Mosaic incidence
Replication	2	140.34	0.03	0.02**	0.24	14.84**	5.00
Genotype	31	12885.22**	5.49**	1.41**	59.04**	838.77**	97.21**
Error	62	67.89	0.02	0.002	0.67	0.81	15.83

\*Significant at 5 per cent level

\*\* Significant at 1 per cent level

**Table 8 Mean value of biometric characters**

Accession No	Plant height (cm)	Primary branches per plant	Plant spread (cm)	Days to fruits flowering	Pollen viability (%)	Days to maturity	Fruits per plant	Fruit length (cm)	Ped cell length (cm)	Fruit girth (cm)	Fruit weight (g)
CC 1	115 00	4 67	271 22	68 00	65 37	36 33	196 45	6 53	4 00	5 77	4 93
CC 2	100 00	3 67	270 44	65 33	75 70	31 00	216 33	5 60	3 80	10 23	8 63
CC 3	118 33	8 67	324 63	69 00	65 47	28 33	274 00	5 57	3 07	6 90	4 08
CC 4	125 67	6 67	304 47	64 00	77 90	37 33	233 33	5 97	3 53	5 27	1 22
CC 5	116 00	6 33	354 58	77 00	4 80	29 67	201 00	5 93	5 20	7 93	6 50
CC 6	94 67	4 33	315 21	72 33	40 10	29 00	37 67	6 17	2 95	9 47	5 82
CC 7	128 00	6 33	454 48	65 33	85 10	31 00	508 89	6 75	3 95	9 93	6 38
CC 8	103 67	4 90	286 33	68 33	33 77	36 67	87 60	5 67	3 2	10 37	6 55
CC 9	77 00	4 33	269 97	70 00	47 30	31 00	27 50	5 53	4 40	10 10	6 55
CC 10	93 00	5 67	294 76	70 00	44 47	29 33	49 00	5 93	2 97	7 23	2 07
CC 11	107 67	5 33	355 69	71 33	80 40	31 33	171 00	8 33	4 10	7 67	5 36
CC 12	122 67	3 33	79 87	65 33	47 3	35 67	188 50	5 60	3 8	10 10	6 72
CC 13	105 00	3 67	268 61	54 67	79 93	37 33	620 00	6 63	2 87	7 07	5 18
CC 14	102 00	4 67	297 17	66 33	38 23	29 35	56 67	4 57	2 60	10 07	5 97
CC 15	106 67	4 00	350 39	67 67	80 80	29 33	326 67	5 90	3 07	10 00	6 87
CC 16	106	8 33	308 66	69 67	69 93	31 67	201 87	6 63	3 10	8 17	3 32
CC 17	66 00	4 33	329 08	71 67	70 73	27 67	79 28	6 00	3 20	9 77	4 73
CC 18	72 67	6 00	265 73	73 33	70 83	35 67	20 37	4 83	2 50	9 73	4 51
CC 19	86 00	7 00	248 71	67 00	47 70	34 67	111 36	4 57	4 43	6 87	4 15
CC 20	114 67	6 00	382 75	79 33	61 57	31 00	66 00	3 80	2 87	7 77	4 05
CC 21	80 67	5 67	370 70	76 67	45 43	32 67	48 33	6 47	3 17	7 73	3 39
CC 22	81 33	4 33	320 44	71 67	72 50	32 67	136 50	5 57	3 97	9 70	4 00
CC 23	102 00	6 00	299 50	60 67	79 0	30 33	637 44	4 10	70	8 97	5 88
CC 24	86 00	4 67	324 89	66 67	60 00	28 00	18 99	6 00	3 23	7 50	5 28
CC 25	83 00	5 00	517 56	76 00	46 60	23 00	86 64	3 60	5 50	6 43	3 17
CC 26	99 67	7 00	316 25	76 33	40 27	30 00	35 33	5 87	3 67	6 80	4 27
CC 27	153 5	5 00	581 97	76 00	81 1	31 33	198 43	7 53	4 07	9 70	7 44
CC 28	97 67	5 33	577 75	74 67	86 40	34 67	213 90	6 47	60	9 20	7 05
CC 29	61	6 33	249 23	83 00	56 9	26 00	99 89	4 63	2 77	9 77	5 13
CC 30	118 00	4 33	568 61	63 00	77 60	22 00	212 33	6 93	3 90	9 10	3 64
CC 31	86 00	6 33	281 96	70 67	86 93	50 67	205 00	5 60	2 90	8 53	5 33
CC 32	68 00	6 00	92 95	74 67	62 43	28 67	19 00	4 93	2 87	6 43	2 42
CD (5%)	11 3719	2 1160	47 7808	4 3482	2 6488	1 8580	26 9797	0 4 75	0 202	0 2815	0 4104

Table 8 Contd

Accession No	Seeds per fruit	1000 seed weight (g)	Yield per plant (g)	Yield per harvest (g)	Number of harvests	Capsaicin (%)	Oleoresin (%)	Ascorbic acid (mg/100g)	Mosaic incidence (V.I.)
CC 1	13 33	3 79	319 49	106 50	3 00	1 20	13 09	119 60	55 42
CC 2	49 00	5 02	919 90	171 66	5 40	2 41	17 86	94 20	51 19
CC 3	25 00	3 73	408 03	108 55	3 77	3 59	10 91	120 37	51 39
CC 4	5 67	4 11	255 71	79 64	3 22	2 85	16 38	106 20	61 61
CC 5	78 33	3 72	479 80	74 43	6 45	2 36	27 18	94 53	51 67
CC 6	45 67	4 15	130 18	38 50	3 35	1 72	9 17	91 07	54 79
CC 7	42 67	3 46	1059 37	263 17	4 03	2 73	16 25	136 33	52 58
CC 8	30 00	3 15	15 05	83 69	3 75	1 4	9 53	104 10	56 55
CC 9	7 67	3 42	12 20	3 42	3 69	1 75	7 26	85 93	57 08
CC 10	45 00	3 04	102 80	2 90	3 2	2 69	10 79	85 60	61 79
CC 11	40 33	3 30	590 54	74 91	5 22	3 03	13 05	105 53	51 19
CC 12	40 33	4 36	807 62	157 87	5 12	2 73	12 64	61 83	54 86
CC 13	41 67	4 10	1455 60	206 97	6 94	3 03	13 94	102 70	40 6
CC 14	2 00	4 60	140 4	65 03	2 16	1 43	8 66	109 60	56 11
CC 15	44 67	4 21	743 45	1 7 38	6 34	1 98	10 38	102 70	51 39
CC 16	25 33	3 95	414 40	40 43	2 96	3 74	24 25	68 50	67 90
CC 17	41 00	4 63	197 38	45 98	4 29	2 48	11 07	102 70	56 35
CC 18	25 33	4 26	85 67	24 54	3 49	2 48	11 66	89 00	51 67
CC 19	27 33	2 83	748 31	59 81	4 15	2 83	7 16	82 20	58 77
CC 20	29 00	4 92	750 81	71 44	3 51	3 15	8 19	75 30	52 43
CC 21	20 00	5 21	154 53	66 59	2 33	2 17	4 92	89 00	53 62
CC 22	33 67	4 11	331 62	83 19	3 99	3 10	16 24	99 30	46 88
CC 23	47 67	4 29	1649 72	778 31	5 93	1 59	9 25	102 70	50 15
CC 24	18 67	4 58	55 00	2 82	2 42	2 97	13 35	98 60	64 58
CC 25	26 67	4 35	143 07	78 49	5 02	1 25	8 33	82 20	48 61
CC 26	21 00	3 74	73 57	6 40	2 03	2 97	13 96	102 70	64 40
CC 7	49 67	4 42	764 36	39 15	5 49	2 57	10 68	130 10	57 92
CC 28	14 67	4 55	667 71	25 93	5 1	2 53	20 30	123 30	54 25
CC 29	26 3	5 11	775 91	7 43	3 76	3 41	11 66	71 90	63 06
CC 0	26	4 09	696 23	16 06	6 00	2 2	7 89	102 70	60 02
CC 1	57 67	4 57	581 87	116 37	5 00	2 53	12 24	109 60	59 08
CC 2	32 67	4 15	51	19 78	7 0	3 43	15 01	106 20	5 27
CD (s)	3 4 94	0 2593	51 3 64	1 4553	0 234	0 0693	1 3412	1 4729	6 4976

(128.00 cm) CC 4 (125.67 cm) and CC 12 (122.67 cm). The accession CC 29 was the shortest (61.33 cm).

### **Primary branches per plant**

Primary branches per plant was found to vary from 3.33 to 8.67. The accessions on an average had 5.44 primary branches per plant. Maximum primary branches was observed in CC 3 (8.67) and minimum in CC 12 (3.33).

### **Plant spread**

Mean plant spread varied from 248.71 to 454.48 cm with a general mean of 315.39 cm. Plant spread was maximum in CC 7 (454.48 cm) and minimum in CC 19 (248.71 cm).

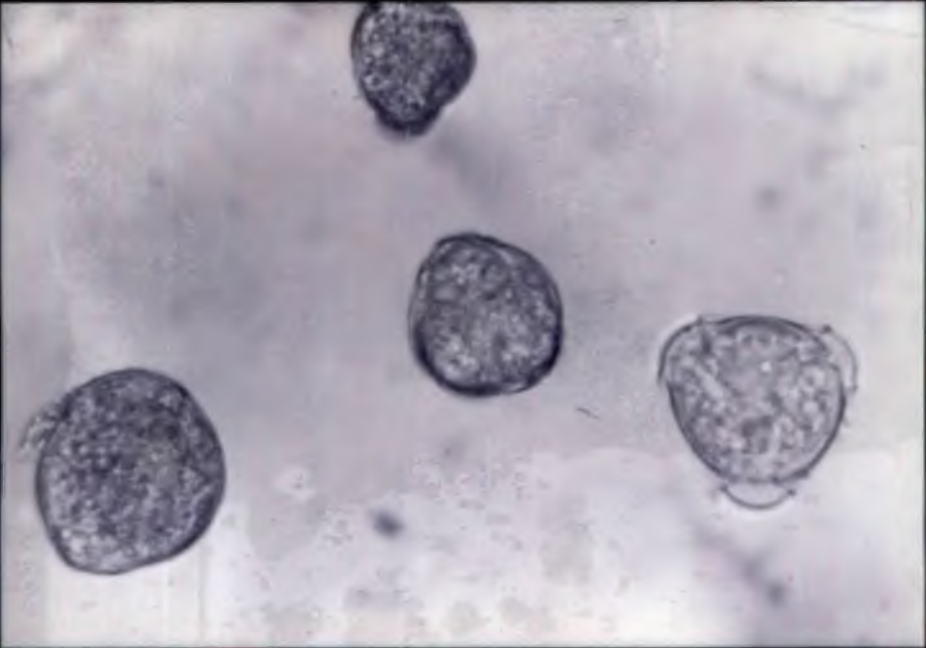
### **Days to first flowering**

Days to first flowering exhibited a range of 54.67 to 83.00. CC 13 was the earliest to flower (54.67) and was significantly different from all other accessions. CC 29 (83.00) was the latest which was on par with CC 20 (79.33).

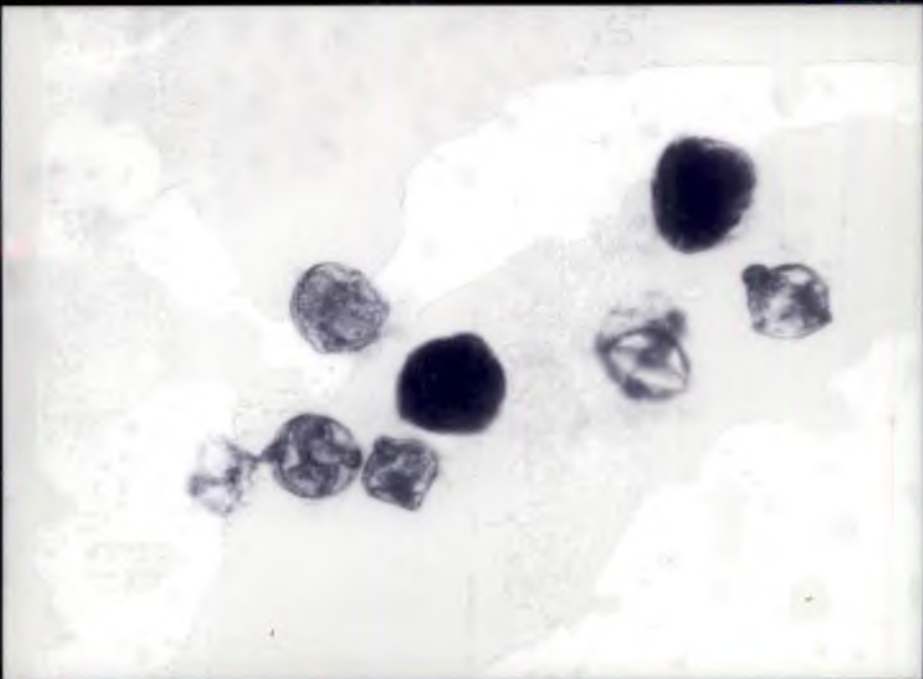
### **Pollen viability**

Wide variation among the accessions was observed for pollen viability (Plates 5 and 6). It ranged from 33.27 per cent in CC 8 to 86.93 per cent in CC 31. The accessions CC 28 (86.40 per cent) and CC 7 (85.10 per cent) were on par with CC 31.

**Plate 5**



**Plate 6**



### **Days to maturity**

The accessions varied significantly for days taken from fruit set to harvest. CC 30 was the earliest to mature, which took 22 days, while CC 4 was the latest (37.33).

### **Fruits per plant**

A wide range of variation was noticed for fruits per plant. Maximum fruits were obtained from CC 23 (637.44) which was on par with CC 13 (620.00). CC 24 had the minimum fruits (18.99) which was on par with CC 32 (19.00), CC 18 (20.37), CC 9 (22.50), CC 26 (35.33), CC 14 (36.67) and CC 6 (37.67).

### **Fruit length**

Fruit length varied considerably from 3.60 cm in CC 25 to 8.33 cm in CC 11 with an overall mean of 5.75 cm (Plate 7).

### **Pedicel length**

Pedicel length among the accessions was found to range from 2.50 (CC 18) to 5.50 cm (CC 25).

### **Fruit girth**

Girth of the fruits varied significantly among the accessions from 5.27 to 10.37 cm. Maximum fruit girth was recorded in CC 8 (10.37 cm) which was on par with CC 2 (10.23 cm) and CC 9 (10.10 cm). Accession CC 4 (5.27 cm) had the minimum fruit girth.

**Plate 7**



**1**



**2**



**3**



**4**



**5**



**6**



**7**



**8**



**9**



**10**



**11**



**12**



**13**



**14**



**15**



**16**



**17**



**18**



**19**



**20**



**21**



**22**



**23**



**24**



**25**



**26**



**27**



**28**



**29**



**30**



**31**



**32**



### **Fruit weight**

Range in fruit weight among the accessions was from 1.22 to 8.63 g highest in CC 2 (8.63 g) and lowest in CC 4 (1.22 g)

### **Seeds per fruit**

Seeds per fruit observed a range from 5.67 in CC 4 to 57.67 in CC 31 with an overall mean of 32.14

### **1000 seed weight**

A narrow range of variation was observed for 1000 seed weight from 2.83 to 5.21 g. CC 21 had the highest 1000 seed weight of 5.21 g and the lowest in CC 19 (2.83 g)

### **Yield per plant**

A wide range of variation was noticed for yield per plant. CC 23 had the highest yield (1649.72 g) which was significantly different from all other accessions. The lowest yield was obtained from CC 32 (51.31 g) which was on par with CC 24 (55.00 g), CC 26 (73.57 g) and CC 18 (85.67 g)

### **Yield per harvest**

Yield per harvest had a range from 19.78 g in CC 32 to 278.31 g in CC 23 with an overall mean of 95.78 g

### Number of harvests

Among the accessions the number of harvests was found to vary from 2.03 in CC 26 to 6.94 in CC 13

### Capsaicin

Capsaicin content observed a range of 1.20 to 3.74 per cent. Among the 32 accessions CC 16 contained the maximum capsaicin (3.74 per cent) whereas CC 1 had the minimum (1.20 per cent)

### Oleoresin

Among the accessions oleoresin content ranged from 4.92 per cent in CC 21 to 24.25 per cent in CC 16 with an overall mean of 12.44 per cent

### Ascorbic acid

Wide variation in ascorbic acid was observed among the accessions with CC 7 containing the maximum of 136.33 mg per 100 g and CC 12 with the minimum (61.83 mg per 100 g)

### Mosaic incidence

The vulnerability index for mosaic incidence ranged from 40.63 to 67.90. Maximum mosaic incidence was observed in CC 16 (67.90) whereas CC 13 (40.63) was the least affected. CC 13 was on par with CC 22 (46.88). The reaction of accessions towards mosaic incidence (Table 9) indicated that 26 accessions were moderately resistant and the remaining six (CC 4, CC 10, CC 16, CC 24, CC 26 and CC 29) were susceptible to the disease.

The accessions were classified into three based on normal distribution property and presented in Table 10. CC 2, CC 7, CC 12, CC 13 and CC 23 were found to be performing better for most of the characters whereas CC 18, CC 24, CC 26 and CC 32 were poor in most of the characters.

#### 4.2.2 Genetic parameters

The population mean, range, phenotypic, genotypic and environmental variances, phenotypic and genotypic coefficients of variation are given in Table 11.

High phenotypic and genotypic variances were observed for several characters including yield per plant, fruits per plant and plant spread. Wide variation was observed in phenotypic and genotypic variances among the characters. A close association between phenotypic and genotypic variances was noticed for yield per plant, fruits per plant, fruit growth and fruit weight. For most of the characters, genotypic variance makes up the major portion of phenotypic variance with very little effect of environment.

Phenotypic and genotypic coefficients of variation (PCV and GCV respectively) observed were high for most of the characters (Fig. 1). Fruits per plant had the highest PCV (90.08) and GCV (89.54) followed by yield per plant (89.39 and 89.12 respectively) and the lowest PCV and GCV were exhibited by days to first flowering (8.85 and 7.99 respectively).

Table 10 Classification of accessions

S No	Character	Poo	Medium	Be
1	Plant height	CC 6 CC 9 CC 10 CC 17 CC 18 CC 9 CC 21 CC 22 CC 24 CC 25 CC 29 CC 31 CC 32	CC 2 CC 14 CC 23 CC 26 CC 28	CC 1 CC 3 CC 4 CC 5 CC 7 CC 8 CC 1 CC 2 CC 3 CC 15 CC 6 CC 2 CC 27 CC 30
2	Primary branches per plant	CC 1 CC 2 CC 6 CC 9 CC 12 CC 13 CC 14 CC 15 CC 17 CC 22 CC 24 CC 30	CC 8 CC 10 CC 11 CC 18 CC 20 CC 21 CC 23 CC 25 CC 27 CC 28 CC 32	CC 3 CC 4 CC 5 CC 7 CC 6 CC 19 CC 26 CC 27 CC 31
3	Plant spread	CC 1 CC 2 CC 8 CC 9 CC 10 CC 13 CC 14 CC 18 CC 19 CC 23 CC 29 CC 31 CC 32	CC 3 CC 4 CC 6 CC 5 CC 16 CC 17 CC 21 CC 22 CC 24 CC 25 CC 26	CC 5 CC 7 CC 11 CC 12 CC 20 CC 27 CC 28 CC 30
4	Days to first flowering	CC 5 CC 6 CC 18 CC 20 CC 21 CC 25 CC 26 CC 27 CC 28 CC 29 CC 32	CC 3 CC 9 CC 10 CC 11 CC 16 CC 17 CC 22 CC 31	CC 1 CC 2 CC 4 CC 7 CC 8 CC 12 CC 13 CC 14 CC 15 CC 19 CC 23 CC 24 CC 30
5	Pollen viability	CC 5 CC 6 CC 8 CC 9 CC 10 CC 2 CC 14 CC 9 CC 20 CC 21 CC 24 CC 25 CC 26 CC 29	CC 32	CC 1 CC 2 CC 3 CC 4 CC 7 CC 11 CC 13 CC 5 CC 16 CC 7 CC 18 CC 22 CC 23 CC 27 CC 28 CC 30 CC 31
6	Days of maturity	CC 3 CC 5 CC 6 CC 0 CC 4 CC 15 CC 17 CC 23 CC 24 CC 25 CC 26 CC 29 CC 30 CC 32	CC 2 CC 7 CC 9 CC 1 CC 16 CC 20 CC 27 CC 3	CC CC 4 CC 8 CC 12 CC 13 CC 18 CC 19 CC 21 CC 22 CC 28
7	Fruits per plant	CC 6 CC 8 CC 9 CC 10 CC 14 CC 17 CC 18 CC 19 CC 20 CC 21 CC 22 CC 24 CC 25 CC 26 CC 29 CC 32	CC 11	CC 1 CC 2 CC 3 CC 4 CC 5 CC 7 CC 2 CC 13 CC 5 CC 16 CC 23 CC 27 CC 28 CC 30 CC



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Table 10 Contd

Sl No	Charac e	Poo	Med um	Bette
8	Fru t length	CC 2 CC 3 CC 9 CC 12 CC 4 CC 18 CC 19 CC 20 CC 22 CC 23 CC 25 CC 29 CC 31 CC 32	CC 8 CC 26	CC 1 CC 4 CC 5 CC 6 CC 7 CC 10 CC 11 CC 13 CC 15 CC 16 CC 17 CC 21 CC 24 CC 27 CC 28 CC 30
9	Ped ce length	CC 3 CC 6 CC 8 CC 10 CC 13 CC 14 CC 5 CC 16 CC 17 CC 18 CC 20 CC 21 CC 24 CC 29 CC 31 CC 32	CC 4 CC 28	CC 1 CC 2 CC 5 CC 7 CC 9 CC 11 CC 2 CC 19 CC 22 CC 23 CC 25 CC 26 CC 27 CC 30
10	Fru t g h	CC 1 CC 3 CC 4 CC 5 CC 10 CC 1 CC 13 CC 16 CC 19 CC 20 CC 2 CC 24 CC 25 CC 26 CC 32	CC 31	CC 2 CC 6 CC 7 CC 8 CC 9 CC 12 CC 14 CC 15 CC 17 CC 18 CC 27 CC 2 CC 27 CC 28 CC 29 CC 30
11	F u we gh	CC 3 CC 4 CC 0 CC 6 CC 17 CC 18 CC 9 CC 20 CC 2 CC 22 CC 25 CC 26 CC 30 CC 32	CC 1 CC 29	CC 2 CC 5 CC 6 CC 7 CC 8 CC 9 CC 1 CC 1 CC 3 CC 4 CC 15 CC 23 CC 24 CC 27 CC 28 CC 31
12	Seeds per fru t	CC 1 CC 3 CC 4 CC 5 CC 8 CC 9 CC 16 CC 8 CC 19 CC 20 CC 21 CC 24 CC 25 CC 26 CC 28 CC 29 CC 30	CC 14 CC 32	CC 2 CC 6 CC 7 CC 10 CC 1 CC 12 CC 13 CC 15 CC 7 CC 22 CC 23 CC 7 CC 3
13	1000 seed ve gh	CC 1 CC 3 CC 5 CC 7 CC 8 CC 9 CC 10 CC 1 CC 16 CC 19 CC 26	CC 4 CC 6 CC 13 CC 5 CC 22 CC 30 CC 32	CC 2 CC 12 CC 14 CC 17 CC 8 CC 20 CC 21 CC 23 CC 24 CC 25 CC 27 CC 28 CC 29 CC 31

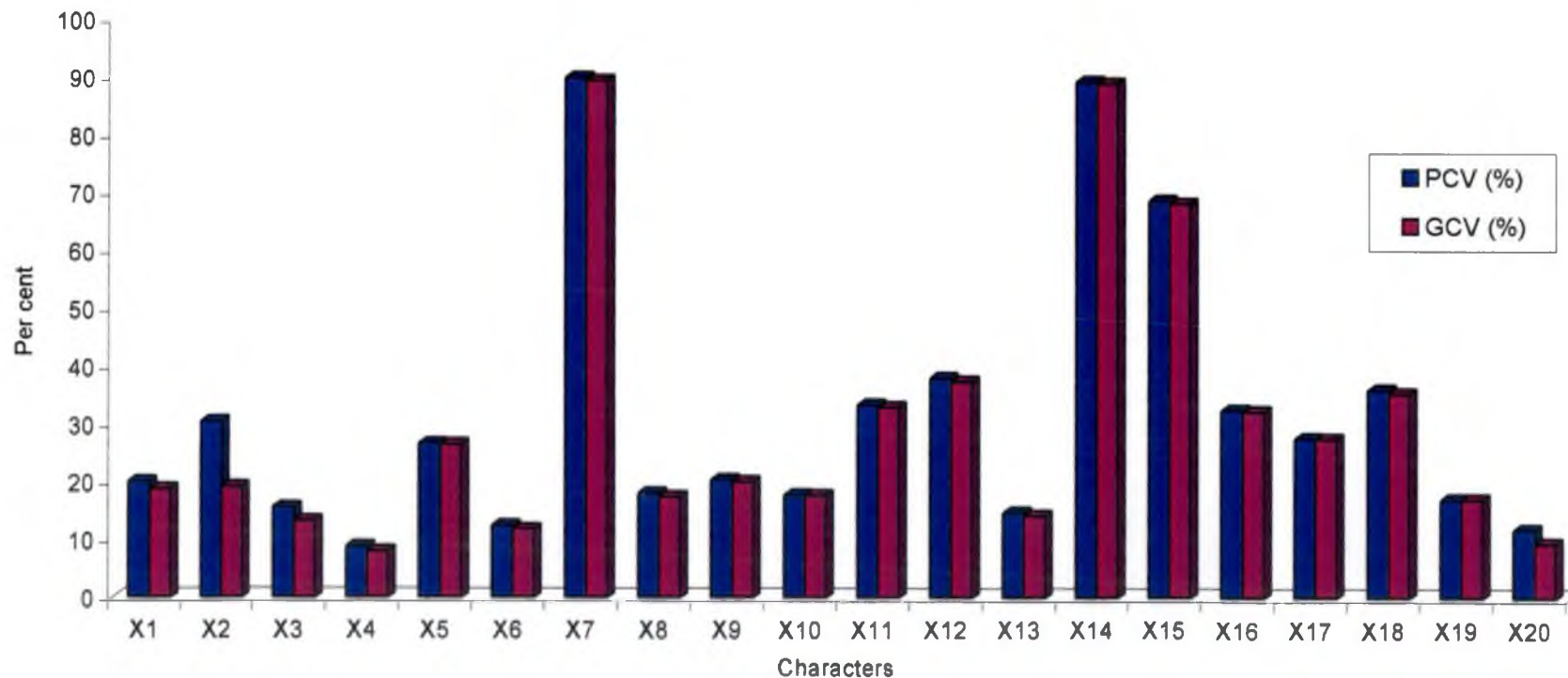
Table 10 Contd

S No	Character	Poor	Med um	Bette
14	Y eld per p an	CC 1 CC 3 CC 4 CC 6 CC 8 CC 9 CC 10 CC 11 CC 14 CC 16 CC 17 CC 18 CC 19 CC 20 CC 21 CC 22 CC 24 CC 25 CC 26 CC 29 CC 32		CC 2 CC 5 CC 7 CC 7 CC 13 CC 5 CC 23 CC 27 CC 28 CC 30 CC 31
15	Y eld per harvest	CC 4 CC 5 CC 6 CC 8 CC 9 CC 10 CC 11 CC 14 CC 17 CC 18 CC 19 CC 20 CC 21 CC 22 CC 24 CC 25 CC 26 CC 29 CC 32		CC 1 CC 2 CC 3 CC 7 CC 2 CC CC 15 CC 16 CC 23 CC 27 CC 28 CC 30 CC 31
16	Number of harves s	CC 1 CC 3 CC 4 CC 6 CC 7 CC 8 CC 9 CC 10 CC 14 CC 16 CC 18 CC 20 CC 21 CC 22 CC 24 CC 26 CC 29 CC 32	CC 19	CC 2 CC 5 CC 11 CC 17 CC 13 CC 5 CC 7 CC CC 25 CC 27 CC 28 CC 30 CC 31
17	Capsa c n	CC 1 CC 2 CC 5 CC 6 CC 8 CC 9 CC 12 CC 14 CC 15 CC 21 CC 23 CC 25 CC 27 CC 30	CC 17 CC 18	CC 3 CC 4 CC 7 CC 10 CC 11 CC 13 CC 16 CC 9 CC 20 CC 22 CC 24 CC 26 CC 28 CC 29 CC 31 CC 32
18	Oleores n	CC 3 CC 6 CC 8 CC 9 CC 10 CC 14 CC 15 CC 17 CC 18 CC 19 CC 20 CC 21 CC 23 CC 25 CC 27 CC 29 CC 30	CC 12 CC 31	CC 1 CC 2 CC 4 CC 5 CC 7 CC 11 CC 13 CC 16 CC 22 CC 24 CC 26 CC 28 CC 32
19	Ascorb c ac d	CC 2 CC 5 CC 6 CC 9 CC 10 CC 12 CC 6 CC 18 CC 9 CC 20 CC 21 CC 25 CC 29		CC 1 CC CC 4 CC 7 CC 8 CC 1 CC 13 CC 14 CC 15 CC 17 CC 22 CC 23 CC 24 CC 26 CC 27 CC 28 CC 30 CC 3 CC 32

**Table 11** Range mean phenotypic genotypic and environmental variances phenotypic and genotypic coefficients of variation for different characters in *C chinense*

Sl No	Character	Range	Mean + SE <sub>m</sub>	$\sigma_p^2$	$\sigma_g$	$\sigma_e^2$	PCV (%)	GCV (%)
1	Plant height (cm)	61.33-135.3	98.69+4.00	393.08	345.01	48.07	20.09	18.82
2	Primary branches per pan	3.33-8.67	5.44+0.75	2.77	1.09	1.68	30.50	19.17
3	Plant spread (cm)	74.8-145.48	115.39+15.1	2439.53	175.20	686.32	15.60	1.78
4	Days to first flowering	54.67-85.00	70.18+1.54	38.53	1.44	7.09	8.85	7.99
5	Pollen viability (%)	3.27-86.9	62.8+0.94	281.71	279.08	2.63	26.7	6.60
6	Days to maturity	22.00-37.33	31.04+0.66	14.92	13.62	1.30	12.44	1.89
7	Fruits per plant	18.99-637.44	168.85+9.54	23132.99	22860.03	272.96	90.08	89.54
8	Fruit length (cm)	3.60-8.5	5.75+0.5	1.08	1.01	0.07	18.07	17.51
9	Ped cell length (cm)	2.50-5.50	3.53+0.07	0.52	0.50	0.02	20.37	20.07
10	Fruit girth (cm)	5.27-10.37	8.45+0.099	2.26	2.23	0.03	17.79	17.67
11	Fruit weight (g)	1.22-8.65	5.02+0.15	2.81	2.75	0.06	33.41	5.03
12	Seeds per fruit	5.67-57.67	32.14+1.21	149.20	144.79	4.41	38.01	7.44
13	1000 seed weight (g)	2.83-5.71	4.12+0.09	0.37	0.34	0.03	14.74	14.23
14	Yield per plant (g)	51.31-1649.72	445.96+18.14	158932.00	157944.40	987.60	89.39	89.12
15	Yield per harvest (g)	19.78-278.51	95.78+4.76	4540.33	4272.44	67.89	68.87	68.28
16	Number of harvests	7.0-6.94	4.18+0.08	1.84	1.82	0.02	32.48	37.30
17	Capsaicin (%)	20.5-74	2.49+0.02	0.47	0.46	0.01	27.57	27.52
18	Oleocanin (%)	4.92-24.25	12.44±0.47	20.13	19.45	0.68	36.07	5.46
19	Ascorbic acid (mg per 100g)	61.83-156.3	98.07+0.52	280.15	279.32	0.81	17.07	17.04
20	Mosaic incidence (Vulnerability Index)	40.63-67.90	55.18+2.30	42.96	27.13	15.83	11.88	9.44

**Fig. 1 Phenotypic and genotypic coefficient of variation for 20 characters in *C. chinense***



X1-Plant height  
 X2-Primary branches per plant  
 X3-Plant spread  
 X4-Days to first flowering  
 X5-Pollen viability

X6-Days to maturity  
 X7-Fruits per plant  
 X8-Fruit length  
 X9-Pedicel length  
 X10-Fruit girth

X11-Fruit weight  
 X12-Seeds per plant  
 X13-1000 seed weight  
 X14-Yield per plant  
 X15-Yield per harvest

X16-Number of harvests  
 X17-Capsaicin  
 X18-Oleoresin  
 X19-Ascorbic acid  
 X20-Mosaic incidence



### 4.2.3 Heritability and genetic advance

Heritability and genetic advance for different characters are presented in Table 12 (Fig. 2).

High heritability coupled with high genetic advance was observed for most of the characters, except primary branches per plant, days to first flowering and mosaic incidence.

Heritability estimates were high for most of the characters studied *viz.*, ascorbic acid (99.71), capsaicin (99.62), yield per plant (99.38) and pollen viability (99.07). Primary branches per plant recorded the lowest but a moderate heritability (39.35).

Genetic advance was highest for fruits per plant (183.37), followed by yield per plant (183.01) and the lowest for days to first flowering (14.87) and mosaic incidence (15.45). High heritability combined with high genetic advance was observed for fruits per plant and yield.

### 4.2.4 Correlation analysis

The phenotypic, genotypic and environmental correlation coefficients were estimated for 20 characters (Tables 13, 14 and 15).

#### (A) Phenotypic correlation

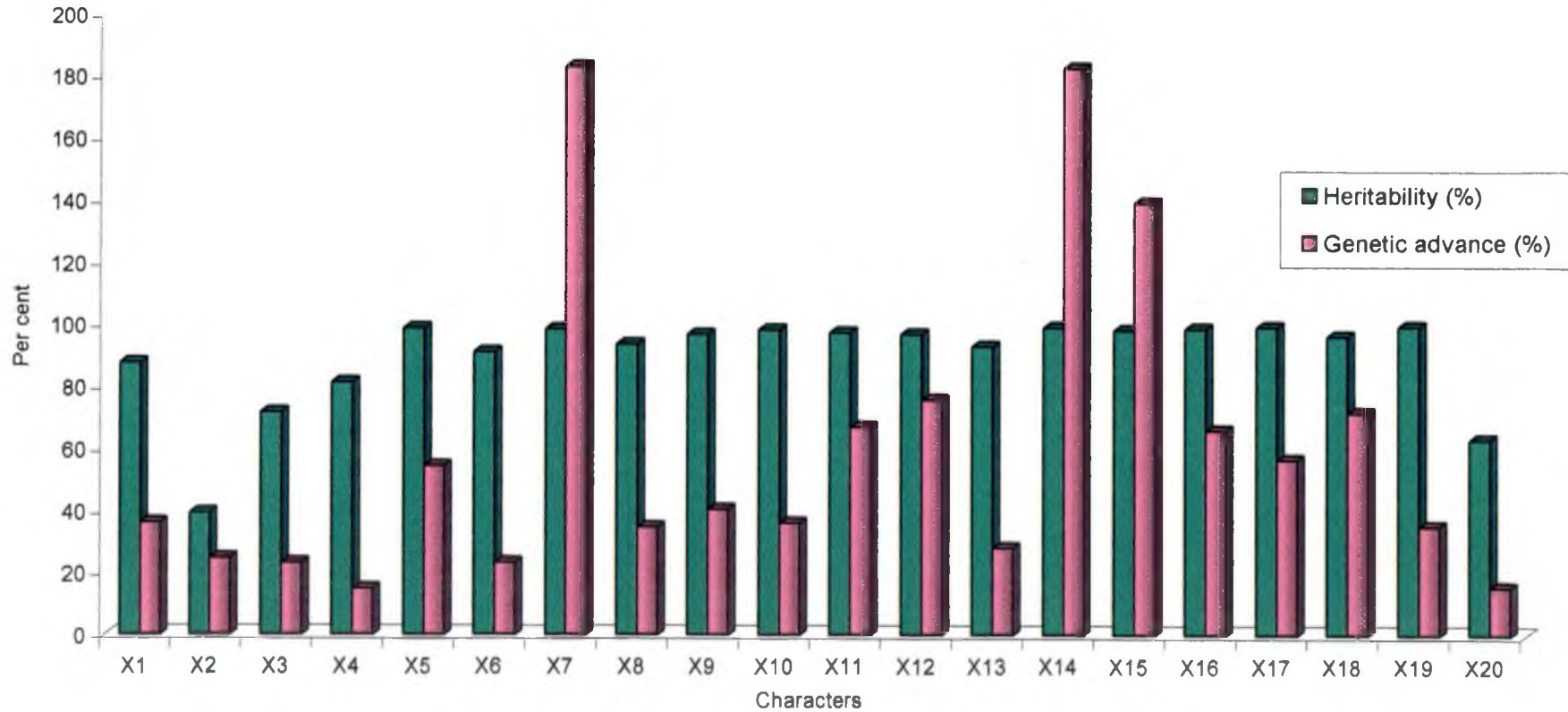
##### (i) Correlation between yield and other characters

Yield per plant recorded high positive correlation with plant height (0.4401), pollen viability (0.5759), fruits per plant (0.9300), fruit weight (0.4513),

**Table 12. Heritability and genetic advance for different characters in *C. chinense***

Sl. No.	Characters	Heritability (%)	Genetic advance (%)
1	Plant height	87.77	36.32
2	Primary branches per plant	39.35	24.77
3	Plant spread	71.87	23.18
4	Days to first flowering	81.60	14.87
5	Pollen viability	99.07	54.53
6	Days to maturity	91.32	23.41
7	Fruits per plant	98.82	183.37
8	Fruit length	93.95	34.97
9	Pedicle length	97.04	40.72
10	Fruit girth	98.68	36.17
11	Fruit weight	97.75	67.28
12	Seeds per plant	97.04	75.99
13	1000-seed weight	93.16	28.29
14	Yield per plant	99.38	183.01
15	Yield per harvest	98.44	139.55
16	Number of harvests	98.89	66.16
17	Capsaicin	99.62	56.58
18	Oleoresin	96.65	71.81
19	Ascorbic acid	99.71	35.05
20	Mosaic incidence	63.15	15.45

**Fig. 2 Heritability and genetic advance for 20 characters in *C. chinense***



X1-Plant height  
 X2-Primary branches per plant  
 X3-Plant spread  
 X4-Days to first flowering  
 X5-Pollen viability

X6-Days to maturity  
 X7-Fruits per plant  
 X8-Fruit length  
 X9-Pedicel length  
 X10-Fruit girth

X11- Fruit weight  
 X12-Seeds per plant  
 X13-1000 seed weight  
 X14-Yield per plant  
 X15-Yield per harvest

X16-Number of harvests  
 X17-Capsaicin  
 X18-Oleoresin  
 X19-Ascorbic acid  
 X20-Mosaic incidence

seeds per fruit (0.4798), number of harvests (0.7304) and ascorbic acid (0.2756). Days to first flowering and mosaic incidence were negatively correlated with yield (-0.5488 and -0.3531 respectively).

### **(ii) Correlation among the yield component characters**

Plant height was positively correlated with plant spread (0.5282), fruits per plant (0.4256), fruit length (0.3862), number of harvests (0.2833) and ascorbic acid content (0.3204) and negatively correlated with days to first flowering (-0.3255).

Fruit girth and fruit weight were negatively correlated with primary branches per plant (-0.3391 and -0.3531 respectively), while capsaicin (0.4058) and mosaic incidence (0.3072) were positively correlated with primary branches per plant.

Fruit length observed high positive correlation with plant spread (0.3080). Days to first flowering recorded negative correlation with most of the characters, the highest being with fruits per plant (-0.5963).

Pollen viability exhibited high positive correlation with several characters like fruits per plant (0.5800), fruit length (0.3460), seeds per fruit (0.2691), number of harvests (0.4827), capsaicin (0.3051), oleoresin (0.3307) and ascorbic acid (0.4774).

Days to maturity was negatively correlated with days to first flowering (-0.2636). Fruits per plant was positively correlated with seeds per fruit (0.3240) and negatively correlated with mosaic incidence (-0.3667).

**Table 13. Phenotypic correlation coefficients among yield and its components**

Character	X1	X2	X3	X4	X5	X6	X7	X8	X9	X10	X11	X12	X13	X14	X15	X16	X17	X18	X19	X20
Plant height (X1)	1.0000																			
Primary branches per plant (X2)	0.0494	1.0000																		
Plant spread (X3)	0.5282**	-0.0479	1.0000																	
Days to first flowering (X4)	-0.3255*	0.1937	0.0229	1.0000																
Pollen viability (X5)	0.2127	0.0029	0.2301	-0.2630	1.0000															
Days to maturity (X6)	0.2222	-0.0121	-0.1536	-0.2636	0.0674	1.0000														
Fruits per plant (X7)	0.4256**	-0.0212	0.0778	-0.5963**	0.5800**	0.2020	1.0000													
Fruit length (X8)	0.3862*	-0.0827	0.3080	-0.1765	0.3460*	0.1775	0.1414	1.0000												
Pedicle length (X9)	0.2150	-0.0758	0.1903	0.0384	-0.1068	-0.1280	0.0650	0.0446	1.0000											
Fruit girth (X10)	-0.0803	-0.3391*	0.1488	-0.0005	0.0637	-0.0767	-0.0103	0.0241	-0.1490	1.0000										
Fruit weight (X11)	0.1928	-0.3531*	0.1330	-0.0898	0.1252	0.1126	0.2430	0.1634	0.1466	0.6862**	1.0000									
Seeds per fruit (X12)	0.0382	-0.2300	0.1366	-0.1669	0.2691	-0.1513	0.3240*	0.0652	-0.1097	0.4553**	0.4151*	1.0000								
1000-seed weight (X13)	-0.2057	-0.1978	0.0381	0.2441	0.2137	-0.1961	-0.0045	-0.2203	-0.3117	0.2137	0.1374	0.0292	1.0000							
Yield per plant (X14)	0.4401**	-0.1410	0.2082	-0.5488**	0.5759**	0.1652	0.9300**	0.1592	0.0920	0.2440	0.4513**	0.4798**	0.0725	1.0000						
Yield per harvest (X15)	0.5176**	-0.0161	0.2900	-0.5145**	0.5611**	0.2138	0.8637**	0.1840	0.0528	0.2468	0.4076*	0.3985*	0.0503	0.9409**	1.0000					
Number of harvests (X16)	0.2833	-0.2675	0.1537	-0.2919	0.4827**	-0.0341	0.6804**	0.1665	0.3676*	0.2427	0.4517**	0.4608**	0.0054	0.7304**	0.5294**	1.0000				
Capsaicin (X17)	-0.0654	0.4058*	0.0781	0.1471	0.3051	-0.0152	0.0004	0.1551	-0.2990	-0.2485	-0.3551*	-0.0772	0.0139	-0.0684	-0.0257	-0.1345	1.0000			
Oleoresin (X18)	0.2029	0.1731	0.0745	-0.0085	0.3307*	0.2118	0.2042	0.2989	0.1056	-0.0770	0.0924	-0.0990	-0.0219	0.1741	0.2381	0.1250	0.4255**	1.0000		
Ascorbic acid (X19)	0.3204*	-0.0449	0.2651	-0.1914	0.4774**	0.1013	0.2670	0.4292**	0.0423	0.0059	0.2458	0.0965	-0.1111	0.2756	0.3101	0.1241	-0.1637	0.1238	1.0000	
Mosaic incidence (X20)	0.0032	0.3072	-0.0410	0.1558	-0.1717	-0.1644	-0.3667*	0.1123	-0.1621	-0.0939	-0.2076	-0.2430	-0.0229	-0.3531*	-0.2369	-0.4308**	0.2037	0.105	-0.1305	1.0000

\* Significant at 5 per cent level

\*\* Significant at 1 per cent level

Fruit length was positively correlated with oleoresin (0.2989) and ascorbic acid contents (0.4292). Pedicel length observed high negative correlation with 1000-seed weight (-0.3117) and capsaicin (-0.2990).

Fruit girth observed high positive correlation with fruit weight (0.6862), seeds per fruit (0.4553) and negatively correlated with capsaicin (-0.2485).

Seeds per fruit was positively correlated with fruit weight (0.4151), whereas capsaicin was negatively correlated with fruit weight (-0.3551).

Mosaic incidence observed high negative correlation with number of harvests (-0.4308).

A high positive correlation was observed between capsaicin and oleoresin contents (0.4255).

## **(B) Genotypic correlation**

### **(i) Correlation between yield and other characters**

High positive correlation was observed between yield per plant and plant height (0.4604), pollen viability (0.5808), fruits per plant (0.9323), fruit weight (0.4614), seeds per fruit (0.4876), yield per harvest (0.9425) and number of harvests (0.7356), whereas days to first flowering and mosaic incidence exhibited a high negative correlation (-0.6131 and -0.4581 respectively). Yield per harvest was positively correlated with plant spread (0.3426).

**Table 14. Genotypic correlation coefficients among yield and its components**

Character	X1	X2	X3	X4	X5	X6	X7	X8	X9	X10	X11	X12	X13	X14	X15	X16	X17	X18	X19	X20
Plant height (X1)	1.0000																			
Primary branches per plant (X2)	0.0594	1.0000																		
Plant spread (X3)	0.6222	-0.0307	1.0000																	
Days to first flowering (X4)	-0.3803	0.3746	0.0659	1.0000																
Pollen viability (X5)	0.2269	0.0027	0.2795	-0.2943	1.0000															
Days to maturity (X6)	0.2346	-0.0533	-0.2163	-0.3092	0.0717	1.0000														
Fruits per plant (X7)	0.4471	-0.0622	0.0830	-0.6651	0.5869	0.2144	1.0000													
Fruit length (X8)	0.4048	-0.1639	0.3772	-0.1925	0.3590	0.1797	0.1484	1.0000												
Pedicel length (X9)	0.2282	-0.1003	0.2217	0.0555	-0.1046	-0.1409	0.0679	0.0511	1.0000											
Fruit girth (X10)	-0.0839	-0.5248	0.1898	0.0091	0.0667	-0.0755	-0.0077	0.0194	-0.1541	1.0000										
Fruit weight (X11)	0.2192	-0.5483	0.1444	-0.1034	0.1324	0.1193	0.2503	0.1646	0.1452	0.6990	1.0000									
Seeds per fruit (X12)	0.0470	-0.3590	0.1757	-0.2050	0.2760	-0.1622	0.3313	0.0658	-0.1176	0.4637	0.4220	1.0000								
1000-seed weight (X13)	-0.2271	-0.2712	0.0199	0.2721	0.2199	-0.2163	-0.0082	-0.2293	-0.3289	0.2265	0.1382	0.0399	1.0000							
Yield per plant (X14)	0.4604	-0.2443	0.2459	-0.6131	0.5808	0.1726	0.9323	0.1626	0.0957	0.2479	0.4614	0.4876	0.0774	1.0000						
Yield per harvest (X15)	0.5391	-0.0655	0.3426	-0.5814	0.5690	0.2250	0.8667	0.1900	0.0575	0.2519	0.4206	0.4062	0.0565	0.9425	1.0000					
Number of harvests (X16)	0.3070	-0.4118	0.1819	-0.3233	0.4872	-0.0404	0.6881	0.1711	0.3743	0.2485	0.4594	0.4684	0.0060	0.7356	0.5395	1.0000				
Capsaicin (X17)	-0.0667	0.6510	0.0980	0.1637	0.3072	-0.0159	0.0003	0.1586	-0.3022	-0.2528	-0.3604	-0.0789	0.0123	-0.0692	-0.0269	-0.1353	1.0000			
Oleoresin (X18)	0.2136	0.2930	0.1084	-0.0153	0.3364	0.2222	0.2131	0.3136	0.1054	-0.0778	0.0941	-0.1062	-0.0316	0.1792	0.2463	0.1255	0.4341	1.0000		
Ascorbic acid (X19)	0.3369	-0.0718	0.3056	-0.2123	0.4803	0.1060	0.2688	0.4420	0.0436	0.0060	0.2498	0.1006	-0.1149	0.2764	0.3122	0.1252	-0.1639	0.1281	1.0000	
Mosaic incidence (X20)	-0.0260	0.4916	-0.0485	0.2672	-0.2044	-0.1788	-0.4701	0.1119	-0.2086	-0.1215	-0.2976	-0.3007	-0.0377	-0.4581	-0.3170	-0.5557	0.2558	0.1272	-0.1631	1.0000



## (ii) Correlation among the yield component characters

Plant height had high positive correlation with plant spread (0.6222), fruits per plant (0.4471), fruit length (0.4048) and ascorbic acid content (0.3369) and negatively correlated with days to first flowering (-0.3803).

Primary branches per plant exhibited high positive correlation with capsaicin (0.6510), mosaic incidence (0.4916) and days to first flowering (0.3746), whereas fruit girth (-0.5248), fruit weight (-0.5483) and seeds per fruit (-0.3590) were negatively correlated.

Fruit length and ascorbic acid were positively correlated with plant spread (0.3772 and 0.3056 respectively). Days to first flowering exhibited high negative correlation with fruits per plant (-0.6651), days to maturity (-0.3092) and pollen viability (-0.2943).

Pollen viability observed high positive correlation with several characters like fruits per plant (0.5869), fruit length (0.3590), seeds per fruit (0.2760), capsaicin (0.3072), oleoresin (0.3364) and ascorbic acid (0.4803) contents and negative correlation with mosaic incidence (-0.2044).

Fruits per plant had high positive correlation with seeds per fruit (0.3313) and negative correlation with mosaic incidence (-0.4701). Oleoresin and ascorbic acid exhibited high positive correlation with fruit length (0.3136 and 0.4420 respectively), whereas 1000-seed weight and capsaicin were negatively correlated with pedicel length (-0.3289 and -0.3022 respectively).



**Table 15. Environmental correlation coefficients among yield and its components**

Character	X1	X2	X3	X4	X5	X6	X7	X8	X9	X10	X11	X12	X13	X14	X15	X16	X17	X18	X19	X20
Plant height (X1)	1.0000																			
Primary branches per plant (X2)	0.0531	1.0000																		
Plant spread (X3)	0.1837	-0.0764	1.0000																	
Days to first flowering (X4)	-0.0248	-0.0555	-0.1210	1.0000																
Pollen viability (X5)	0.0316	0.0157	-0.1121	0.0394	1.0000															
Days to maturity (X6)	0.1183	0.0863	0.1384	0.0260	-0.0264	1.0000														
Fruits per plant (X7)	0.2433	0.2082	0.1349	0.0209	-0.0704	-0.0524	1.0000													
Fruit length (X8)	0.2165	0.0884	-0.0148	-0.0751	-0.0150	0.1520	-0.0607	1.0000												
Pedicle length (X9)	0.0732	-0.1028	0.0560	-0.1479	-0.2589	0.0921	-0.0774	-0.0994	1.0000											
Fruit girth (X10)	-0.0538	-0.1345	-0.1810	-0.1771	-0.2044	-0.1511	-0.2183	0.1897	0.0873	1.0000										
Fruit weight (X11)	-0.1959	-0.1115	0.1501	0.0399	-0.3558	-0.0024	-0.1868	0.1527	0.2025	-0.0199	1.0000									
Seeds per fruit (X12)	-0.0862	-0.0609	-0.1108	0.2105	-0.0914	0.0283	-0.0260	0.0562	0.1493	0.0781	0.1614	1.0000								
1000-seed weight (X13)	-0.0035	-0.1648	0.1569	0.0612	0.0975	0.0444	0.1198	-0.0898	0.0228	-0.1181	0.1407	-0.1959	1.0000							
Yield per plant (X14)	0.3653	0.1915	0.0100	0.0987	-0.0438	0.0356	0.7132	0.1097	-0.1494	-0.1616	-0.2926	0.0704	-0.0974	1.0000						
Yield per harvest (X15)	0.3770	0.2528	0.0267	0.1217	-0.0647	0.0146	0.6547	0.0401	-0.1618	-0.1015	-0.2657	0.0671	-0.1160	0.8840	1.0000					
Number of harvests (X16)	-0.0745	-0.1291	0.0067	-0.0336	0.0512	0.1382	0.0167	0.0610	0.0525	-0.2341	0.0012	0.1045	-0.0135	0.1352	-0.2197	1.0000				
Capsaicin (X17)	-0.1403	-0.0377	-0.1452	-0.0170	-0.0256	-0.0051	0.0119	0.1108	-0.1825	0.2947	0.0663	0.0376	0.1227	0.0800	0.1102	-0.0393	1.0000			
Oleoresin (X18)	0.0972	-0.0535	-0.1635	0.0653	0.0858	0.0554	-0.2053	0.0023	0.1097	-0.0448	0.0355	0.1224	0.1676	-0.1051	-0.0935	0.1194	0.0355	1.0000		
Ascorbic acid (X19)	0.2771	0.0036	0.2229	0.0067	0.0145	0.0142	0.0365	0.1066	-0.0728	-0.0032	-0.0956	-0.2664	-0.0242	0.0990	0.1134	-0.0380	-0.1054	-0.1941	1.0000	
Mosaic incidence (X20)	0.1062	0.1314	-0.0258	-0.1383	-0.1714	-0.1601	0.0709	0.1747	0.0113	0.0286	0.2885	-0.0731	0.0380	0.2034	0.1716	0.1305	0.0217	0.0509	-0.0313	1.0000

Fruit girth had high positive correlation with fruit weight (0.6990), seeds per fruit (0.4637) and negatively correlated with capsaicin content (-0.2528).

Fruit weight also had a negative correlation with capsaicin (-0.3604) along with mosaic incidence (-0.2976), but positively correlated with seeds per fruit (0.4220). Seeds per fruit exhibited high negative correlation with mosaic incidence (-0.3007).

Number of harvests had a more or less similar trend of correlation as that of yield per plant.

Capsaicin had high positive correlation with oleoresin (0.4341).

### **(C) Environmental correlation**

Environmental correlation coefficients were found to be negligible among yield and its component characters, except for the correlation between fruits per plant and yield per plant (0.7132).

#### **4.2.5 Path Analysis**

In path coefficient analysis, the genotypic correlation coefficients among yield and its component characters were partitioned into direct and indirect contribution of each character to fruit yield (Table 16 and Fig. 3). Plant height, days to first flowering, pollen viability, fruits per plant, fruit weight, seeds per fruit, number of harvests, ascorbic acid and mosaic incidence were selected for path coefficient analysis.

**Table 16. Direct and indirect effect of selected yield components on fruit yield in *C. chinense***

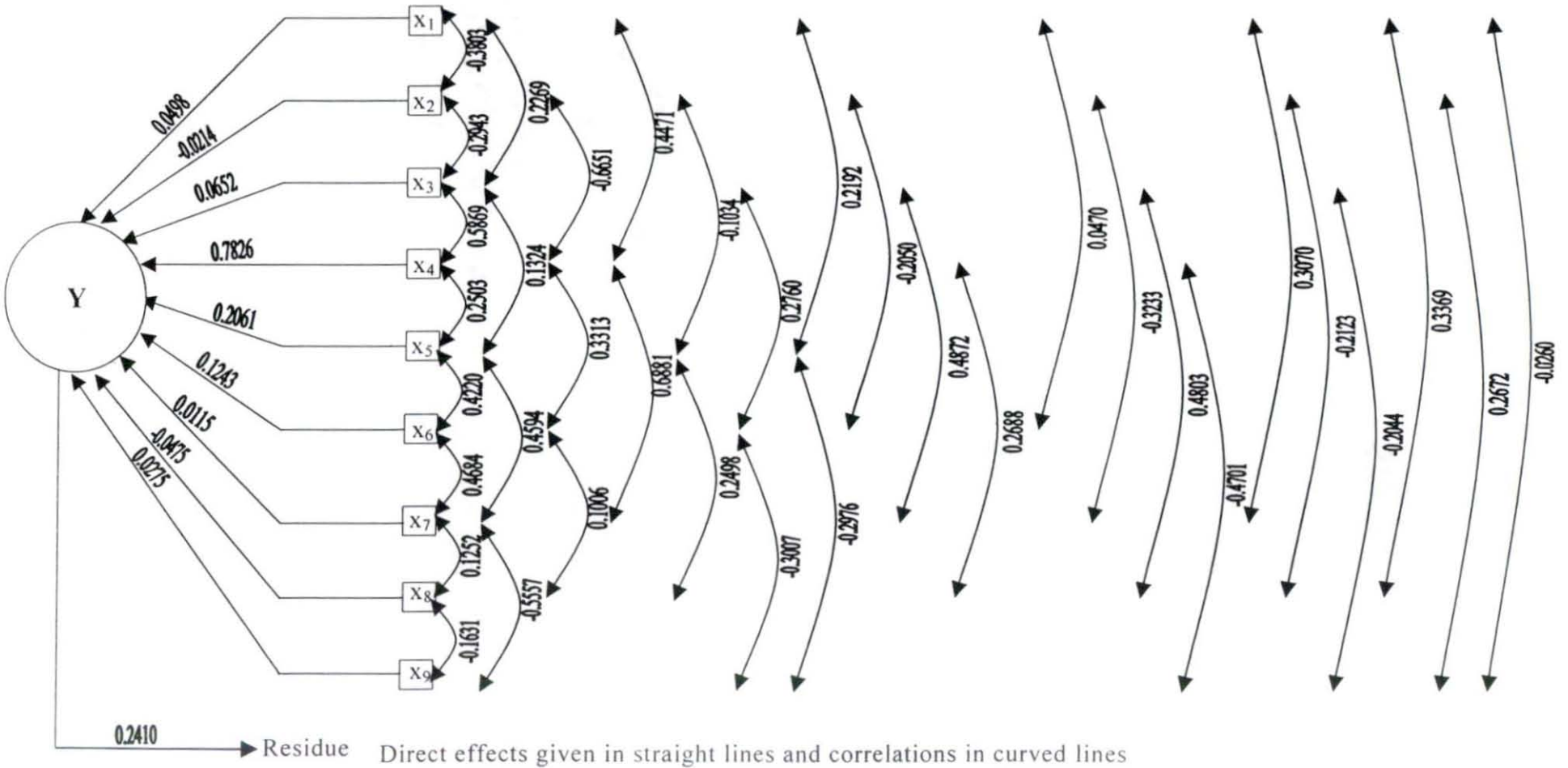
Character	Plant height	Days to first flowering	Pollen viability	Fruits per plant	Fruit weight	Seeds per fruit	Number of harvests	Ascorbic acid	Mosaic incidence	Correlation with yield
Plant height	<b>0.0498</b>	0.0081	0.0148	0.3499	0.0452	0.0058	0.0035	-0.0160	-0.0007	0.4604
Days to first flowering	-0.0189	<b>-0.0214</b>	-0.0192	-0.5205	-0.0213	-0.0255	-0.0037	0.0101	0.0073	-0.6131
Pollen viability	0.0113	0.0063	<b>0.0652</b>	0.4593	0.0273	0.0343	0.0056	-0.0228	-0.0056	0.5808
Fruits per plant	0.0222	0.0143	0.0382	<b>0.7826</b>	0.0516	0.0412	0.0079	-0.0128	-0.0129	0.9323
Fruit weight	0.0109	0.0022	0.0086	0.1959	<b>0.2061</b>	0.0525	0.0053	-0.0119	-0.0082	0.4614
Seeds per fruit	0.0023	0.0044	0.0180	0.2593	0.0870	<b>0.1243</b>	0.0054	-0.0048	-0.0083	0.4876
Number of harvests	0.0153	0.0069	0.0317	0.5385	0.0947	0.0582	<b>0.0115</b>	-0.0059	-0.0153	0.7356
Ascorbic acid	0.0168	0.0045	0.0313	0.2104	0.0515	0.0125	0.0014	<b>-0.0475</b>	-0.0045	0.2764
Mosaic incidence	-0.0013	-0.0057	-0.0133	-0.3679	-0.0613	-0.0374	-0.0064	0.0077	<b>0.0275</b>	-0.4581

Residue = 0.2410

Direct effects- diagonal elements

Indirect effects- off diagonal elements

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



Y – Yield per plant

X<sub>1</sub> – Plant height

X<sub>2</sub> – Days to first flowering

X<sub>3</sub> – Pollen viability

X<sub>4</sub> – Fruits per plant

X<sub>5</sub> – Fruit weight

X<sub>6</sub> – Seeds per fruit

X<sub>7</sub> – Number of harvests

X<sub>8</sub> – Ascorbic acid

X<sub>9</sub> – Mosaic incidence

Fruits per plant exhibited the highest positive direct effect on fruit yield (0.7826), followed by fruit weight (0.2061) and seeds per fruit (0.1243). The direct effects of plant height, pollen viability, number of harvests and mosaic incidence were negligible, whereas days to first flowering and ascorbic acid exerted small and negative direct effects on yield.

Indirect effects through fruits per plant were consistently high signifying the importance of that character. Thus in the case of plant height, pollen viability and number of harvests, high positive correlation with yield was mainly due to their positive indirect effects through fruits per plant (0.3499, 0.4593 and 0.5385 respectively). Similarly, high negative correlation of days to first flowering and mosaic incidence on yield was due to high negative indirect effects through fruits per plant (-0.5205 and -0.3679 respectively). In the case of fruit weight and seeds per fruit, the correlation was mainly built by the direct as well as indirect effect *via* fruits per plant.

#### 4.2.6 Selection Index

A discriminant function analysis was carried out for isolating superior genotypes. Selection index involving characters *viz.*, plant height ( $X_1$ ), days to first flowering ( $X_2$ ), pollen viability ( $X_3$ ), fruits per plant ( $X_4$ ), fruit weight ( $X_5$ ), seeds per fruit ( $X_6$ ), yield per plant ( $X_7$ ), number of harvests ( $X_8$ ), ascorbic acid ( $X_9$ ) and mosaic incidence ( $X_{10}$ ) were selected for the analysis.

The selection index worked out was as follows :

$$I = 0.6548872 X_1 - 0.8016408 X_2 + 1.396126 X_3 + 0.8978062 X_4 + 3.657391 X_5 \\ + 0.7279018 X_6 + 0.9976921 X_7 + 2.350527 X_8 + 0.9246957 X_9 - 0.4314854 X_{10}.$$

The scores obtained for the accession based on the selection index were given in Table 17.

Based on selection index, CC 23 (7471.67) ranked first, followed by CC 13 (6805.48) and CC 7 (4955.16) (Plates 8, 9 and 10). The minimum scores were obtained for CC 32 (762.16) and CC 24 (781.35).

#### 4.2.7 Mahalanobi's $D^2$ analysis

Following Mahalanobi's  $D^2$  statistic, the 32 accessions of *C. chinense* were subjected to cluster analysis, based on ten characters *viz.*, plant height, days to first flowering, pollen viability, fruits per plant, fruit weight, seeds per fruit, yield per plant, number of harvests, ascorbic acid and mosaic incidence.

The 32 accessions fell under six clusters. The clustering pattern is furnished in Table 18. Cluster I was the largest with 21 accessions, followed by cluster II with 6 accessions and cluster III with two accessions. Clusters IV, V and VI had one accession each.

The cluster means of the ten characters are presented in Table 19. Cluster IV consisted of taller accession (CC 5) which was late in flowering, whereas cluster V (CC 13) comprised of early flowering accession, which had more number of harvests and low mosaic incidence. Cluster III (CC 2 and

**Table 17. Selection indices arranged in descending order**

Rank	Accessions	Selection index
1	CC 23	7471.67
2	CC 13	6805.48
3	CC 7	4955.16
4	CC 2	4248.36
5	CC 15	3926.26
6	CC 27	3756.06
7	CC 12	3504.63
8	CC 30	3408.03
9	CC 28	3367.46
10	CC 31	3098.68
11	CC 3	2643.41
12	CC 5	2573.48
13	CC 11	2416.41
14	CC 16	2333.01
15	CC 1	2186.01
16	CC 4	2055.50
17	CC 22	2010.79
18	CC 8	1731.02
19	CC 19	1537.20
20	CC 29	1512.32
21	CC 20	1494.36
22	CC 17	1441.58
23	CC 25	1130.55
24	CC 14	1102.03
25	CC 6	1035.57
26	CC 21	1031.92
27	CC 10	941.39
28	CC 9	909.99
29	CC 18	883.80
30	CC 26	804.73
31	CC 24	781.35
32	CC 32	762.16



**Plate 8**



**Plate 9**



**Plate 10**





**Table 18. Clustering pattern of accessions**

Cluster No.	Number of accessions	Accessions
I	21	CC 14, CC 6, CC 9, CC 21, CC 18, CC 10, CC 25, CC 26, CC 17, CC 24, CC 32, CC 19, CC 20, CC 29, CC 4, CC 1, CC 8, CC 22, CC 11, CC 3, CC 16
II	6	CC 28, CC 30, CC 15, CC 31, CC 27, CC 12
III	2	CC 2, CC 7
IV	1	CC 5
V	1	CC 13
VI	1	CC 23

Table 18 Clustering pattern of accessions

Cluster No	Number of access ons	Accessions
I	21	CC 14 CC 6 CC 9 CC 21 CC 18 CC 10 CC 25 CC 26 CC 17 CC 24 CC 32 CC 19 CC 20 CC 29 CC 4 CC 1 CC 8 CC 22 CC 11 CC 3 CC 16
II	6	CC 28 CC 30 CC 15 CC 31 CC 27 CC 12
III	2	CC 2 CC 7
IV	1	CC 5
V	1	CC 13
VI	1	CC 23

**Table 19 Cluster means of ten biometric characters**

Cluster	Plant height (cm)	Days to first flowering	Pollen viability (%)	Fruits per plant	Fruit weight (g)	Seeds per fruit	Yield per plant (g)	Number of harvests	Ascorbic acid (no/100g)	Mosaic incidence (V I)
I	92.51	71.68	56.53	96.75	4.33	27.89	212.62	3.42	94.17	56.28
II	110.72	69.56	76.67	223.81	6.18	38.89	710.21	5.54	105.04	56.25
III	114.00	65.35	80.40	285.11	7.51	45.84	989.64	4.72	115.27	51.89
IV	116.00	77.00	42.80	201.00	6.50	28.33	479.80	6.45	94.53	51.67
V	105.00	54.67	79.93	620.00	5.18	41.67	1435.60	6.94	102.70	40.63
VI	102.00	60.67	79.30	637.44	5.88	47.67	1649.72	5.93	102.70	50.15

CC 7) had the highest pollen viability fruit weight and ascorbic acid content Yield per plant fruits per plant and seeds per fruit were highest in cluster VI (CC 23)

The average inter and intracluster distances are presented in Table 20 The cluster diagram is shown in Fig 4

The intracluster distance was on the increase with increasing cluster size Cluster I had the highest intracluster distance (229.93) followed by clusters II and III (217.55 and 188.74 respectively)

The highest intercluster distance was observed between clusters I and VI (1965.74) followed by clusters I and V (1640.10) and clusters IV and VI (1606.19) The genetic distance (D) between clusters I II and IV were largest with cluster VI The minimum intercluster distance was observed between clusters V and VI (339.74) indicating a close relationship among the accessions included

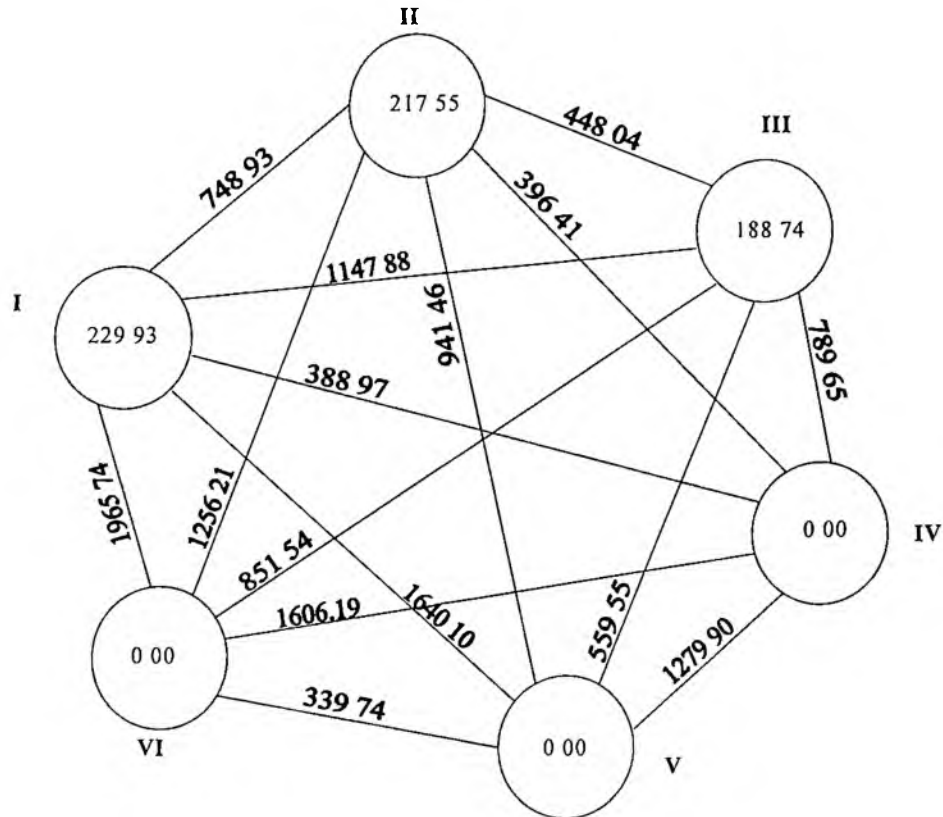
**Table 20** Average inter and intracluster distances

Cluster	I	II	III	IV	V	VI
I	229 93	748 93	1147 88	388 97	1640 10	1965 74
II		217 55	448 04	396 41	941 46	1256 21
III			188 74	789 65	559 55	851 54
IV				0 00	1279 90	1606 9
V					0 00	339 74
VI						0 00

Diagonal elements    ntracluster values

Off diagonal elements    ntercluster values

Fig 4 Cluster diagram



The values in circles indicate intracluster D values and the values on lines indicate intercluster D values

# *Discussion*

## 5 DISCUSSION

The genus *Capsicum* is noted for its richness in diversity. However the exploration of this diversity is mainly restricted to the most widely cultivated species *C. annuum*. *C. chinense* characterized by its perennial habit and highly pungent fruits cultivated in the homesteads of Kerala. The two recognized varieties of *C. chinense* viz. Scotch Bonnet and Habanero are known for their extreme pungency.

Kerala is blessed with diverse climatic and soil conditions which have helped in the development of different landraces of crops having high variability. These landraces are the products of natural selection maintain genetic heterogeneity in balance over time (Sharma 1994). The exploitation of this heterogeneity can help in the improvement of the crop.

The genetic improvement of any crop aims at increasing the production potential and quality by altering the genetic makeup of the existing varieties. To achieve this goal a plant breeder requires information on certain genetic parameters like variability, heritability, genetic advance and association between characters. For the development of superior varieties selection is the fundamental process which utilizes the available variability in a crop.

Hence a study was undertaken to collect and catalogue the available landraces of *C. chinense* for various morphological characters and to assess the magnitude of genetic variability for identifying superior genotypes based on yield, quality and pest and disease resistance.



## 5.1 Genetic cataloguing

Genetic cataloguing based on standard descriptors helps to easily describe the morphological features of a genotype and thus helps exchange of information about new accessions in a more clear way.

Thirty two accessions of *C. chinense* upon cataloguing showed distinct variations among each other with respect to vegetative inflorescence, fruit and seed characters. The accessions had either erect or compact growth habit with green to purple stem and leaves. Flowers per axil were either two or three with a prominent annular constriction at calyx. Fruit colour and fruit shape showed wide variation among the accessions. There are reports on high variability for morphological characters in *C. annuum* (Indra 1994), in *C. frutescens* (Sheela 1998) and in *C. chinense* (Cherian 2000).

## 5.2 Variability

An insight into the magnitude of variability present in a crop species is of utmost importance as it provides a basis for effective selection. The observed variability in the population is the sum total of the variations that arise due to genotypic and environmental effects. Hence a knowledge on the nature and magnitude of genetic variation contributing to gain under selection is essential.

In the present investigation analysis of variance revealed highly significant differences among the thirty two accessions for all the characters studied namely plant height, primary branches per plant, plant spread, days to first flowering, pollen viability, days to maturity, fruits per plant, fruit length, pedicel length, fruit girth, fruit weight, seeds per fruit, 1000 seed weight.

yield per plant yield per harvest number of harvests capsaicin oleoresin ascorbic acid and mosaic incidence. Such variation indicated the scope for improving the population for these characters as reported earlier by Hiremath and Mathapati (1977) Gopalakrishnan *et al* (1987) and Kumar *et al* (1993) in chilli.

In respect of vegetative characters ample variability was observed as evident from the wide range obtained for plant height and plant spread. Among the accessions evaluated CC 27 was the most vigorous registering the highest values for plant height and CC 7 for plant spread. Considerable variability was reported by Gopalakrishnan *et al* (1987) for plant height and Sahoo *et al* (1990) for plant spread. Primary branches per plant recorded a low range of variation compared to other characters as reported by Sahoo *et al* (1990) in chilli.

The accession CC 13 was the earliest to flower. It also had the maximum harvests and high fruit yields. In the present study days to first flowering varied from 54.67 to 83.00. Pollen viability showed a wide range of variation among the accessions. Similar results were also reported by Pradeepkumar (1990). Days to maturity recorded a narrow range of variation. Most of the accessions attained fruit maturity around 51 days after fruit set. Cherian (2000) reported a similar range for days to maturity in *C. chinense*.

Pedicle length in the present study ranged from 2.50 to 5.50 cm. Pedicle in chilli is non-edible and fruits with short pedicle is desirable. Considerable variation for the character was also reported by Rani (1996a) and Cherian (2000).

Among the accessions maximum fruit weight was observed in CC 2. Other accessions with better fruit weight were CC 27 and CC 28. Both fruit length and fruit girth contributed to better fruit weight in CC 2. In the present study fruit length ranged from 3.60 to 8.33 cm. Similarly fruit girth also varied from 5.27 to 10.37cm suggesting ample variability and scope for improvement of fruit size in *C. chinense*.

Fruits per plant and yield per plant exhibited high variability. Among the accessions evaluated fruits per plant and yield were maximum in CC 23 (Nemom Thiruvananthapuram) followed by CC 13 (Vithura Thiruvananthapuram) and CC 7 (Vithura Thiruvananthapuram). The high yield in CC 23 may be attributed to the high fruits per plant, pollen viability and more harvests. CC 13 was characterized by least mosaic incidence and earliness resulting in more number of harvests and finally yield. The accession CC 7 apart from being better for most of the fruit characters like fruit length, fruit girth and fruit weight, it also registered high values for plant height and plant spread indicating a vigorous nature. This confirms the fact that fruit yield is a complex trait and is the ultimate expression of many component characters.

Seeds per fruit exhibited a wide range of variation from 5.67 to 57.67 whereas 1000 seed weight had a narrow range as indicated by a low phenotypic and genotypic variances. Wide variability in seeds per fruit was observed by Arya and Sain (1976) and V. Jayalakshmi *et al* (1989). Sahoo *et al* (1990) reported a low range of variation in 100 seed weight. Varieties with high fruit seed weight and fruit seed number are preferred not only to increase crop production but also to meet the needs of the seed industry and farmers.

Capsaicin the pungent principle of chilli is considered to be one of the most important quality characters. In the present study wide variation was observed between the accessions for capsaicin content. This variation could probably be due to the presence of gene modifying factors for pungency and the ratio of placental tissue to seed and pericarp. Varietal variation in capsaicin contents in chilli was also reported by Pradeepkumar (1990) Sheela (1998) Cherian (2000) and Sreelathakumary (2000). All the accessions evaluated in the present study had high capsaicin content (>1 per cent) which are particularly valued for their pungency and for the manufacture of high capsaicin oleoresin. In this context the accessions CC 9 and CC 27 which produced yellow fruits at mature stage will be more economical for capsaicin extraction as there will not be any interference of red pigments.

Oleoresin represents the total flavour extract of ground spices and consists of fixed oil, capsaicin, pigments, sugars and resin. They are now being extensively used in processed foods and also pharmaceutical products (Bajaj *et al.* 1980). The results obtained in the current investigation revealed considerable variation among the accessions for oleoresin content. This is in agreement with the results obtained by Pradeepkumar (1990) Cherian (2000) and Sreelathakumary (2000). The high pungency oleoresin obtained from the yellow fruited accessions namely CC 9 and CC 27 can be of high value in the pharmaceutical and cosmetic industries where high pungency and low colour are desirable.

The nutritive value of chilli is largely determined by the content of ascorbic acid. Significant variation in ascorbic acid content between accessions was noted in the present study. Such wide variation was also

reported by Rani (1994) and Todorova *et al* (1997). Accessions with high ascorbic acid content are suitable for vegetable purposes.

Mosaic is a serious disease affecting chilli and is a major constraint in chilli cultivation in Kerala. Significant differences were observed among the accessions for mosaic incidence which clearly indicated that the level of resistance or susceptibility to the disease varied with the accession. Out of the 32 accessions evaluated, 26 were moderately resistant and the remaining six were found to be susceptible to the disease. Screening for chilli mosaic resistance was also done by Thakur *et al* (1985), Anandam (1992), Kalloo (1994) and Fatima (1999). The accessions which were found to be moderately resistant in the present study may be used as resistant donors for imparting disease resistance to otherwise desirable genotypes.

High coefficients of variation (phenotypic [PCV] and genotypic [GCV]) were observed for fruits per plant, yield per plant and fruit weight. Similar results were also reported by Nair *et al* (1984), Jabeen *et al* (1999) and Cherian (2000). The high PCV and GCV observed for these characters are evident from their high variability which in turn offers good scope for selection. The lowest PCV and GCV was exhibited by days to first flowering which was in conformity with the findings of Cherian (2000). The GCV was very near to PCV for most of the characters, indicating a highly significant effect of genotype on phenotypic expression with very little effect of environment.

### 5.3 Heritability and genetic advance

The total variability existing in a population is a sum of heritable and non heritable components and it is necessary to partition these components since the magnitude of heritable variability is an important aspect of genetic constitution of breeding material.

High values of heritability were observed for most of the characters studied. Higher magnitude of heritability (>90 %) was registered for yield per plant, fruits per plant, fruit length, fruit girth, fruit weight, seeds per fruit, 1000 seed weight, pollen viability, days to maturity, number of harvests, capsacin, oleoresin and ascorbic acid contents. Similar findings were also reported by Rajput *et al* (1981) for fruits per plant and fruit yield, Narci *et al* (1984) for capsacin, Bhagyalakshmi *et al* (1990) and Kumar *et al* (1993) for ascorbic acid and Singh *et al* (1994) for fruit characters. High heritability estimates indicate the presence of large number of fixable additive factors and hence these traits can be improved by selection.

High heritability estimates does not necessarily mean a high genetic advance for a particular character. The effectiveness of selection depends upon the heritability and genetic advance of the character selected. The present investigation revealed high heritability coupled with high genetic advance for several biometric characters including fruits per plant, yield per plant, fruit weight, fruit girth and fruit length. Jabeen *et al* (1998) also observed high heritability and genetic advance for several yield characters.

High heritability coupled with low genetic advance attributable to non additive gene action was noticed for days to first flowering. Similar results were reported by Nar *et al* (1984) and V. Jayalakshmi *et al* (1989).

On the basis of the present study it can be concluded that simultaneous selection based on multiple characters having high estimates of heritability and genetic advance might be of appreciable use in this crop.

#### 5.4 Correlation studies

Correlation provides information on the nature and extent of relationship between all pairs of characters. A study of correlation among yield and its components will be of great value in planning and evaluating breeding programmes.

In the present study both at phenotypic and genotypic levels the characters viz. plant height, pollen viability, fruits per plant, fruit weight, seeds per fruit and number of harvests showed strong positive association with yield per plant. Days to first flowering and mosaic incidence were negatively correlated with yield.

The very high positive association of fruits per plant with yield indicated that fruits per plant was the primary yield attribute in chili. Similar reports were also suggested by Sundaram and Irulappan (1998) and Cheran (2000).

The present investigation revealed that plant height was positively correlated with plant spread and yield. Ramakumar *et al* (1981) reported a strong correlation between plant spread and plant height. Aliyu *et al* (2000) supported the association between plant height and yield. Production of

increased vegetative growth like plant spread and plant height might lead to larger canopy of the plant resulting in increased fruits per plant and finally yield

Positive association of fruit weight and seeds per fruit with yield was in agreement with the findings of Ranı (1996b) and Mishra *et al* (1998) High correlation observed between number of harvests and yield was in conformity with that of Sheela (1998)

The study revealed a strong positive association between pollen viability and yield This can be explained as when pollen viability increases it increases the availability of viable pollen for pollination and fruit set thereby leading to an increased fruit set fruits per plant and finally yield This was supported by high correlation of pollen viability with fruits per plant which in turn was correlated with yield

The high negative correlation between days to first flowering and yield was supported by the findings of Warade *et al* (1996) and Sreelathakumary (2000) Hence any selection aimed for earliness will be useful for improving yield and yield associated characters

Capsaicin was found to be positively correlated with primary branches per plant pollen viability and oleoresin and negatively with pedicel length and fruit weight Hence selection based on these characters may be done to improve capsaicin and a simultaneous improvement in oleoresin contents A negative correlation of capsaicin with fruit weight was also reported by Jiang *et al* (1987) and Ranı (1995) Aiming at capsaicin improvement selection of small fruited accessions will reduce the fruit yield leading to decrease capsaicin output per unit area Hence medium weight fruited accessions with fairly better capsaicin content may be selected Similarly ascorbic acid was also correlated with yield and yield components



Mosaic incidence exhibited high negative correlation with yield. This might be due to its negative association with other yield contributing factors like pollen viability, fruits per plant, fruit weight, seeds per fruit and number of harvests. Moreover, late flowering accessions were more affected by the disease, which also showed reduced pollen viability. Mosaic induced pollen sterility leading to reduced fruit set and fruits per plant was reported by Jayarajan and Ramakrishnan (1961).

On the basis of the present study, it is evident that selection based on earliness increased plant height, pollen viability, fruit weight, fruits per plant, seeds per fruit and number of harvests along with mosaic resistance may be done for yield improvement.

### 5.5 Path coefficient analysis

Yield is a complex quantitative character governed by a large number of genes and is greatly influenced by environmental factors. The present investigation of path coefficient analysis provided information on the nature of association of several characters contributing to yield by means of untangling the direct and indirect contribution of various characters in building up a complex correlation. As evidenced from correlation studies, path coefficient analysis also signifies the importance of the character fruits per plant which exhibited the highest direct and indirect effects on fruit yield. Similar results were also reported by Cherian (2000) and Munshi *et al* (2000).

Fruit weight and seeds per fruit also exhibited direct positive effects on yield. The direct effects of plant height, pollen viability, number of harvests

nosa c n c d e n c e days to first flower ng and ascorbic acid content were small and negligible but their nd rect effects through fruits per plant were consistently high This was n conform ty with the f ndings of Sundaram and Ranganathan (1978)

## 5 6 Selection index

Select on ndex prov des information on y eld components and thus aids n indirect select on for the mprovement of y eld It involves discriminant function analysis which is meant for solating superior genotypes based on the phenotyp c and genotyp c correlations Ident fication of super or genotypes of *C ch nense* based on d scr minant function analys s was done by Cherian (2000) A model nvolving the same set of ten characters wh ch was used for path coefficient analysis was selected for rank ng the accessions On rank ng the scores obtained the accession CC 23 (Nemom Thiruvananthapuram) ranked frst followed by CC 13 (V thura Th ruvananthapuram) and CC 7 (V thura Th ruvananthapuram) These accessions w th high yield quality and mosaic res stance may be recommended as elite types after refinement and mult locational testing

## 5 7 Mahalanobis $D^2$ analysis

Breeding crop plants adopt ng hybrid zation as a tool is one of the most nportant crop improvement methods The success of hybrid zat on programme is ma nly dependent on the genetic d versity of the parents chosen for the purpose Crosses between genetically diverse parents are likely to produce high heterot c effects Mahalanob s  $D^2$  stat stic s one of the potent techniques for measuring genetic divergence at both ntra and ntercluster

levels and thus provides a basis for selection of genetically divergent parents in hybridization programmes. Genetic divergence in chili was assessed by Mehra (1978) Sundaram *et al* (1980) Gill *et al* (1982) Varalakshmi and Harbabu (1991) Indra (1994) and Roy and Sharma (1996).

In the present study based on Mahalanobis  $D^2$  statistic the 32 accessions were grouped into six gene constellations. The maximum number of accessions (21) were included in cluster I followed by cluster II (six) and cluster III (two). Clusters IV, V and VI had one accession each. The pattern of clustering closely followed the ranking obtained from selection index.

Considering the cluster means for the various characters studied, clusters III, V and VI were superior for most of the biometric characters, whereas clusters I and IV were generally poor. Cluster II was found to be intermediate. For crop improvement programmes, intercrossing among accessions with outstanding mean performance for these characters would be effective.

The study revealed maximum divergence between clusters I and VI followed by clusters I and V as shown by the high intercluster distances. Clusters V and VI with least divergence showed a close relationship between the accessions CC 13 and CC 23. The intracuster distance was maximum for cluster I which had the maximum number of accessions. On the basis of the present study, the accessions of clusters V and VI may be used as base materials for hybridization with selected accessions of cluster I in order to obtain desirable segregants with high yield potential.

The present investigation on 32 *C. chinense* accessions showed wide variation for almost all the characters studied. High heritability coupled with

high genetic advance was observed for most of the biometric characters which indicates the scope for effective selection. Correlation and path coefficient analysis revealed that fruits per plant is the primary yield component. The accessions CC 23 (Nemom Thiruvananthapuram), CC 13 (Vithura Thiruvananthapuram) and CC 7 (Vithura Thiruvananthapuram) were found to be promising with regard to yield, quality and mosaic resistance. The same may be used for further improvement programmes.

*Summary*

## 6 SUMMARY

The present investigation on Genetic cataloguing of hot chilli (*Capsicum chinense* Jacq.) was conducted at the vegetable research plot of the Department of Olericulture College of Agriculture Vellayani during September 2000 to May 2001.

The study envisaged genetic cataloguing of the available germplasm in *C. chinense* assessment of genetic variability divergence association among the characters including the direct and indirect effects of various characters on yield and formulation of a selection index for identifying suitable lines based on yield quality and pest and disease resistance.

The experimental material consisted of 32 accessions collected from different parts of Kerala. The experiment was laid out in a randomised block design with three replications. The accessions were genetically catalogued based on the descriptor list for *Capsicum* (IBPGR 1995). The results revealed distinct variations among the accessions with respect to vegetative inflorescence fruit and seed characters.

Significant differences were observed among the accessions for all the characters studied viz. plant height primary branches per plant plant spread days to first flowering pollen viability days to maturity fruits per plant fruit length pedicel length fruit girth fruit weight seeds per fruit 1000 seed weight yield per plant yield per harvest number of harvests capsaicin oleoresin ascorbic acid and mosaic incidence.

The highest yield was observed in CC 23 (Nemom Thiruvananthapuram 1649 72 g) which also recorded the maximum fruits per plant (637 44). CC 13 (V th ra Th ruva anthapura n) was the earliest to flower (54 67 days) with maximum number of harvests (6 94) and least vulnerability index for mosaic (40 63). Among the accessions CC 27 was the tallest (133 33 cm). The maximum plant spread was exhibited by CC 7 which also recorded the highest ascorbic acid content (136 33 mg per 100 g). CC 2 had the maximum fruit weight (8 63 g). The highest capsaicin (3 74 per cent) and oleoresin (24 25 per cent) contents were recorded by CC 16. Out of the 32 accessions evaluated for mosaic resistance 26 were moderately resistant and the remaining six were susceptible to the disease.

High coefficients of variation (phenotypic [PCV] and genotypic [GCV]) were recorded for fruits per plant, yield per plant and fruit weight. The lowest PCV and GCV were exhibited by days to first flowering.

High heritability coupled with high genetic advance was observed for yield per plant, fruit per plant, fruit weight, fruit girth and fruit length and catting scope for improvement of these characters through selection.

Correlation studies revealed that at both phenotypic and genotypic levels characters like plant height, pollen viability, fruits per plant, fruit weight, seeds per fruit, yield per harvest and number of harvests were positively correlated with yield. Days to first flowering and mosaic incidence were negatively correlated with yield.

Path coefficient analysis indicated that fruits per plant exerted the maximum positive direct effect (0 7826) on yield followed by fruit weight.

(0 2061) and seeds per fruit (0 1243) The indirect effects through fruits per plant were consistently high signifying the importance of that characters

A selection index was worked out using ten characters viz plant height days to first flowering pollen viability fruits per plant fruit weight seeds per fruit yield per plant number of harvests ascorbic acid content and mosaic incidence Based on the index scores obtained CC 23 (Nemom Thruvananthapuram) ranked first followed by CC 13 (Vthura Thruvananthapuram) and CC 7 (Vthura Thruvananthapuram)

The 32 accessions were grouped into six clusters based on Mahalanobis  $D^2$  statistic Cluster I was the largest which contained 21 accessions followed by cluster II with six and cluster III with two accessions Clusters IV V and VI had one accession each With regard to the cluster means clusters V (CC 13) and VI (CC 23) performed better for most of the characters taken The maximum intercluster distance was observed between clusters I and VI (1965 74) followed by clusters I and V (1640 10) Cluster I had the highest intracluster distance (229 93)

Comparison among the accessions for various biometric characters revealed that CC 23 CC 13 and CC 7 were found to be promising based on the superiority in yield quality and mosaic resistance and hence they may be utilized for further crop improvement

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**GENETIC CATALOGUING OF  
HOT CHILLI (*Capsicum chinense* Jacq.)**

BY

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**ABSTRACT OF THE THESIS  
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## ABSTRACT

The research project Genetic cataloguing of hot chilli (*Capsicum chinense* Jacq) was carried out in the vegetable research plot of the Department of Olericulture College of Agriculture Vellayani during September 2000 to May 2001. The objective of the study was to catalogue the accessions based on the IBPGR descriptor for *Capsicum* and to estimate the genetic parameters for different traits in the germplasm for identifying superior lines based on yield, quality and pest and disease resistance.

Thirty two accessions of *C. chinense* collected from various sources upon cataloguing pointed out wide variation for several morphological characters. Analysis of variance revealed significant difference among the accessions for all the characters studied namely plant height, primary branches per plant, plant spread, days to first flowering, pollen viability, days to maturity, fruits per plant, fruit length, pedicel length, fruit girth, fruit weight, seeds per fruit, 1000 seed weight, yield per plant, yield per harvest, number of harvests, capsaicin, oleoresin, ascorbic acid and mosaic incidence.

Among the accessions CC 23 recorded the maximum yield (1649.72 g) as well as fruits per plant (637.44). CC 13 was the earliest to flower (54.67 days) with the maximum number of harvests (6.94) and least vulnerability index for mosaic (40.63). Fruits per plant recorded the maximum phenotypic

and genotypic coefficients of variation followed by yield per plant and fruit weight

High heritability coupled with high genetic advance was observed for fruits per plant yield per plant fruit weight fruit girth and fruit length

Correlation studies and path coefficient analysis revealed that fruits per plant is the primary yield component as evidenced from its high positive correlation as well as high direct and indirect effects on yield

In the discriminant function analysis CC 23 (Nemom Thiruvananthapuram) ranked first followed by CC 13 (Vithura Thiruvananthapuram) and CC 7 (Vithura Thiruvananthapuram)

Based on the analysis for genetic divergence the 32 accessions were grouped into six clusters with the maximum intercluster distance observed between clusters I and VI

On the basis of the present study the accessions CC 23 CC 13 and CC 7 were found to be promising with regard to yield quality and mosaic resistance and the same may be utilized for further improvement