BREEDING FOR CASSAVA MOSAIC DISEASE RESISTANCE

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INTRODUCTION

PREVALENCE OF CASSAVA MOSAIC DISEASE HAS RESULTED IN REDUCED YIELD IN TAMIL NADU: REDUCED STARCH TURN OVER IN KERALA: OBLITERATION OF VARIETAL CHARACTERS & VARIETIES BECOME EXTINCT

REDUCED VEGETATIVE GROWTH LEADING TO SHORTAGE OF PLANTING MATERIAL

DISEASE SEVERITY MORE IN RAINFED AREAS & DROUGHT PRONE AREAS. LESS SYMPTOM EXPRESSION SEEN IN SOME HILLY AREAS OF TAMIL NADU. IN KADAMBUR FOREST, ERODE Dist.

Severe symptom expression (CMD) in H220

Symptom free plants of H165 in Kadambur, Erode
MATERIALS AND METHODS

- Repeated field screening done in different seasons and locations.
- Graft testing done with susceptible symptom expressing stalk for confirmation of resistance.
- Field evaluation of MNga-1 for direct utilization: Repeated over different seasons & locations.
- Hybridization varieties with MNga-1

Varieties involved in crosses with MNga-1:
Sree Rekha, Sree Jaya, CI-848, CO-2, Sree Prabha, Sree Vijaya, CI-742, Ambakadan.

Graft testing of Mnga-1 & M. Caerulescens

MATERIALS AND METHODS

- EVALUATION OF F1 LINES AND SELECTION
- CLONAL EVALUATION & SELECTION
- CLONAL MULTIPLICATION & REPLICATED TRIALS
- FARMER'S PARTICIPATORY EVALUATION
- ALL INDIA MULTI LOCATION TRIALS
- GENETIC ENHANCEMENT OF CMD RESISTANT GENE
- BACK CROSS BREEDING, SELFING & OP. PROGENY
- STUDIES ON INHERITANCE OF CMD RESISTANCE
- INTERSPECIFIC HYBRIDIZATION WILD GENES
- TRUE SEED PROGRAMME - RAPID SELECTION
SELECTION CRITERIA USED
DISEASE RESISTENCE

RESISTANT – R
MODERATELY RESISTANT
WITH RECOVERING SYMPTOM \{ RV & RV+ \}

SUSCEPTIBLE S

DISEASE RECOVERY SYMPTOM

SELECTION CRITERIA

Tuber Yield, Starch Content, Tuber Shape (Top Priority)
Culinary Quality, Tuber Flesh Colour - (White & Yellow)
  rind pealability, purple/pink rind (Market preference).
  • Starch content measured instantly based on specific gravity method (factory method).
Later confirmed by chemical methods.
Evaluation trials : Design : RBD, Replications :3
Plot size : 25 plants

Measuring starch of potato using Specific Gravity Method (Factory Method) – FPR Programme.
Farmers participatory evaluation

FOR SELECTION OF LOCATION SPECIFIC VARIETIES.

RESULT

*Mnga -1 selected as resistant variety (germplasm introduction)

proposed for release in irrigated plains (Tamil Nadu) - Sree Padmanabha.


- Sree Sahya - 24.8% - (6-46.7)
- Sree Vijaya - 18.5% - (8-28.5)
- Sree Prakash - 35.1% - (20.6-50.0)
- Sree Visakham - 31.6% - (2.6-66.9)
- Sree Harsha - 18.6%
- H 226 - 27.7% - (15.6-39.8)
- H 165 - 28.5% - (2.5-48.3)
- H 97 - 31.3% - (1.4-45.9)
- Mnga1 - 0.3% - (0-1.0)

Mnga -1: yield & tuber characters.

- Yield : 23.1-34.2 t/ha
- Dry matter : 30.6 - 37.4
- Starch : 23.5-28.2 %
- Cyanogen : 36.5-40 /ug/100gm
- Culinary quality : well cooked, non bitter, tasty.
**MNga-1:(TMS 30001) Distinguishing Features**

- Resistant to Cassava Mosaic Disease- confirmed through field screening, graft testing and progeny studies.
- Preferred by farmers for its CMD – resistant character in OFT-conducted in various districts in Kerala and Tamil Nadu.
- Observed as free from Tuber rot even in cent percent soil moisture for prolonged period. (Papyreddipatty, Dharmapuri dist.)
- Reported as resistant to Cassava Bacterial Blight. (CBB) - IITA, Nigeria.
- Susceptible to drought, spidermites and white fly.

**Al CRP Trials: IET Ca 99.**

**Yield (t/ ha) - 3 years**

<table>
<thead>
<tr>
<th>Varieties</th>
<th>2001-'02</th>
<th>2002-'03</th>
<th>2003-'04</th>
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<tr>
<td>CI-848</td>
<td>15.80</td>
<td>38.13</td>
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<tr>
<td>Sree Prabha</td>
<td>15.80</td>
<td>39.93</td>
<td>30.46</td>
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<td>CI-850</td>
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<td>MNga-1</td>
<td>16.9</td>
<td>35.93</td>
<td>35.26</td>
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<tr>
<td>H-282</td>
<td>17.21</td>
<td>19.27</td>
<td>17.9</td>
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<tr>
<td>H-152</td>
<td>19.3</td>
<td>29.23</td>
<td>20.7</td>
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<tr>
<td>H-740</td>
<td>20.90</td>
<td>34.03</td>
<td>27.4</td>
</tr>
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</table>

**OFT in Tamil Nadu plains : 2004-'05**

<table>
<thead>
<tr>
<th>Lines</th>
<th>L1 yield</th>
<th>L1 starch</th>
<th>L2 yield</th>
<th>L2 starch</th>
<th>L3 yield</th>
<th>L3 starch</th>
<th>L4 yield</th>
<th>L4 starch</th>
<th>L5 yield</th>
<th>L5 starch</th>
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<td>31.2</td>
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<td>24.00</td>
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<td>Y-6-2</td>
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<td>Y-4-4</td>
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<td>29.6</td>
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<td>23.25</td>
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<td>Y-3-1</td>
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<td>15.00</td>
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<td>MNga-1</td>
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<td>12.5</td>
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<td>28.0</td>
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</tbody>
</table>
Mnga-1-Belukurichi, Namakkal

Mnga-1 at PR Patty, Dharmapuri

Mnga-1 at SPAC, Erode.

PROGENIES EVALUATED IN FIELD: SEEDLING & CLONAL

First Filial Generation: 7,500 seedlings 162 selected
Back cross-1 generation: 185 Lines evaluated
Selfed generation: 17 lines evaluated
Open pollinated (Half sib Generation): 120 lines evaluated
Total lines evaluated: 484
Evaluation of lines selected for resistance and yield characters

F1 - Lines in RBD trial : 162
F1 - In location trial:   10 (3 locations in Kerala, Tamil Nadu & AP)
F1 - In farmers participatory evaluation : 25 (6 locations in Tamil Nadu & Kerala)

<table>
<thead>
<tr>
<th>Sl. No.</th>
<th>Lines</th>
<th>Starch percentage</th>
<th>Mean Tuber yield</th>
<th>Resistance To CMD</th>
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<tbody>
<tr>
<td>1</td>
<td>CMR-1</td>
<td>27.7</td>
<td>34.5 t/ha</td>
<td>RV</td>
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<tr>
<td>2</td>
<td>CMR-2</td>
<td>27.9</td>
<td>27.5 t/ha</td>
<td>RV +</td>
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<tr>
<td>3</td>
<td>CMR-3</td>
<td>28.8</td>
<td>29.7 t/ha</td>
<td>RV</td>
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<tr>
<td>4</td>
<td>CMR-4</td>
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<td>47.1 t/ha</td>
<td>RV</td>
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<tr>
<td>5</td>
<td>CMR-5</td>
<td>24.7</td>
<td>19.0 t/ha</td>
<td>RV</td>
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<td>CMR-6</td>
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<tr>
<td>7</td>
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<td>9</td>
<td>Mnga-1</td>
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<td>37.4 t/ha</td>
<td>R</td>
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<tr>
<td>10</td>
<td>Sree Vijaya</td>
<td>21.1</td>
<td>27.1 t/ha</td>
<td>S</td>
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</tbody>
</table>

CD 0.5 = 11.85

RV= No symptom expressing.  RV+ = symptom expressing but recovers completely
RV+ = symptom expressing but shows partial recovery

IET 1: Evaluation trial of CMD resistant lines.

IET 2: Evaluation trial of CMD resistant lines.

Cassava Mosaic Resistant lines selected
HIGHLY RESISTANT SELECTION WITH NORMAL YIELD

Novel variation induced in crosses with MNgla-1

Open pollination progeny

CMR LINES IN SANKAGIRI
Genetics of CMD resistance in Mnga-1

Mapping population: CO2 X Mnga-1

141 F1 progenies

Susceptible lines: 68
Resistant lines: 73

CMD resistance segregation ratio = 1:1

CMD resistance controlled by single dominant gene

Frequency distribution disease in the F1 progeny:
Phenotypic score

1-Symptomless plant
5-Severe CMD symptom

Frequency distribution of phenotypic values of CMD resistant F1 mapping population
Enhancement of CMD resistance through pyramiding of genes.

- Fifty percent gene expression in first F1 generation was found enhanced to 70 to 89% in selfed, open pollinated and backcrossed generation of selection made from Mnga-1 crosses.
- 374 lines selected based on resistance and tuber characters planted for clonal evaluation in 2005-06.

Interspecific hybridization for cassava mosaic resistance

- H. 99/14 - 3 - BC line with M. glaziovii genome.
- CMD resistant & High Yielding (> 50t/ha)

Leaf yield : 6kg/plant
No. of leaves : 2529
Tuber Dry matter : 42.46
Tuber protein : 2.9%

High Leaf retention
Ten times higher than top branching
Three times higher than spreading
Canopy spread : 2.1m

Interspecific hybridization for protein enrichment M. esculenta x M. tristis

TRUE SEED PROGRAMME:

- Rapid multiplication of cassava lines having disease resistance and yield attributes in traditional and new areas using true seeds
- Minimizing progeny variation using a male sterile line (Ambakadan) in pollination block planted in mixed rows with the disease resistant line Mnga-1
- Seedlings and clones evaluated in diverse environment for selection of potential lines suitable to the location
- Realization of location specific lines within minimum time spent for selection and multiplication
Conclusion and future prospects

- **Strength**: The strength of the programme is the genetic variations created using a potential donor parent MNga-1 (TMS 30001) thanks to IITA, Nigeria, and CIAT, Cali, Colombia. Our strength lies also in the enthusiastic farmers.

- **Weakness**: Lack of MAS (Marker assisted Selection) methods for rapid selection process. HRD (Human Resource Development) to be increased along with funding to quicken the results of selection and multiplication.

- **Opportunity**: There is opportunity in the production of more varieties with CMD resistance, root yield, high starch, nutrient content and industrial qualities along with culinary traits. There is pressing demand for new varieties from farmers, industrialists and traders.

- **Threat**: Natural appearance of new strains of virus is the major threat. Fresh genes, or gene combination are to be developed from exotic, or pathogen derived or from wild source.