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Section of Plant Breeding and Seed Production

Via P. Giuria, 15 - 10126 Turin - Italy

Editors
P. Belletti, L. Quagliotti

DI.VA.P.R.A

Section of Plant Breeding and Seed Production
Via P. Giuria, 15 – 10126 Truin-Italy
Fax: int. Code + 11/650.27.54

Scientific Committee

A. Andrasfalvy, Hungary
A. Palloix, France

R. Gil Ortega, Spain
L. Quagliotti, Italy

C. Shifriss, Israel

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The picture in the cover is derived from the “Herbanrio nuovo di Castore Duante”, Venetia,
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APPRECIATION TO DR. EDMOND POCHARD

Dr. E. Pochard recently retired from his activity in INRA, France. ~ We believe that all the members of the EUCARPIA Capsicum group will join us in wishing Dr. Pochard a challenging new post-Capsicum career.

Dr. Pochard is an outstanding pepper agronomist, geneticist, phytopathologist, pepper breeder and above all a generous person who inspired and helped many of us with scientific encouragement and valuable germplasm!

The late Dr. P. Smith named Dr. Pochard as number one pepper geneticist in Europe.

We need just to mention the 'Lamuyo' F1 hybrid and the haploids-dihaploids stories to appreciate the brilliance of his scientific leadership. It is our privilege to wish Dr. Pochard a pleasant retirement in the knowledge that he will always find fresh fields for creative expression.

The Scientific Committee and the Editorial Board

of "Capsicum Newsletter" 3

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FOREWORD

Our faithful readers will find a surprise together with this issue of "Capsicum Newsletter". In fact, we have printed a special volume of our Newsletter. It includes the Proceedings of the EUCARPIA Meeting on Genetics and Breeding of Capsicum and Eggplant, which is going to be held next September in Rome, Italy. We hope that our decision to distribute the Proceedings using the mailing list of "Capsicum Newsletter" will be appreciated.

The invited paper included in this issue is particularly interesting: it deals with the molecular mapping in pepper and has been written by J. Prince and his collaborators. We thank them very much for their kind co-operation. As usual, any suggestions on the subjects and/or authors to be considered for the invited papers in the following issues of "Capsicum Newsletter" will be appreciated.

Although several contributions have not been accepted (the Scientific Committee has started doing its job we have not modified any of the published papers. Therefore, the authors themselves are not only responsible for the scientific content but also for the form of their own reports.

A 'literature review' is again present in this issue. We hope it will be useful and we would like to remind you to send us a copy of your articles, especially those published in journals of limited circulation.

Please, remember that a subscription fee to the Newsletter is requested. The subscription fees have not been changed: 20 U.S.\$ for normal subscribers and 100 U.S.\$ for supporters. Remember also that now it is possible to book your own copy of the journal: just fill in the

order form on page 55 and send it to us. In the meantime your subscription fee should be paid directly to EUCARPIA Secretariat (please note that the address has been modified!). Please, do not send cheques to us in Turin, as we are not allowed by Italian law to run any financial activities.

Again we have to complain about the lack *of* attention paid by many authors to the instructions on the enclosed sample sheet. Please, cooperate with us and follow these instructions very carefully. Otherwise we will not accept the contributions and they will be sent back to the authors.

Lastly, we have to announce that the journal's Editorial Board has decided to enlarge the space available to each contribution. Starting from the next issue, you will have at your disposal four pages per article.

Piero Belletti and Luciana Quagliotti

Turin, 31st May 1992

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MOLECULAR MAPPING IN PEPPER:. AN UPDATE

James P. Prince,* Molly M. Kyle,** Vincent K. Lackney, James R. Blauth, John F. Murphy, and Steven D. Tanksley.

Department of Plant Breeding and Biometry, Cornell University, Ithaca, NY 14853 USA.

*Current address: USDA-ARS-MPPL, Building 011 A, Room 252, BARC-West, Beltsville, MD 20705 USA.

**To whom correspondence should be addressed.

Pepper (Capsicum L.) is an important vegetable crop worldwide. Much classical breeding work has been done to transfer disease resistances and desirable horticultural traits (see Greenleaf, 1986, for a thorough review of pepper breeding, and Pickersgill, 1989, for a review of genetic resources in Capsicum). Many traits, especially those conditioned by genes with incremental effects, are not amenable to classical breeding approaches. In these cases, a marker that is linked and easily selectable could increase the efficiency of gene transfer and selection. Morphological genes as tags are often difficult to use, as major genes may control a wide range of responses, may affect expression of the gene of interest, and may not be dominant and scorable in all genetic backgrounds.

Phenotype-neutral molecular markers such as isozymes and restriction fragment length polymorphisms (RFLPs) can be used to tag genes and to construct genetic maps of the entire genome. Major genes can be tagged by one or more molecular markers for tracking in a breeding program (Tanksley et al., 1989), complex genetic traits can be dissected into their component parts (Osborn et al., 1987, Nienhuis et al., 1987, and Paterson et al., 1990), introgression of genes can be monitored (Young et al., 1988), genetic variation within and between species can be examined (Figdore et al., 1988, Wang and Tanksley, 1989, and Miller and Tanksley, 1990a), and homologies among related genomes can be investigated (Tanksley et al., 1988, and Bonierbale et al., 1988). Livneh et al., (1990) have used an RFLP for hybrid identification in

pepper seed production. For recent reviews of the use of molecular markers in plant improvement, see Paterson et al., (1991) and Tanksley et al., (1989).

For making a molecular map of any crop species, polymorphism between interfertile plants and the segregating population that results from a cross between those plants are needed (in the case of RFLPs, the polymorphisms are different restriction fragments detected by hybridization to cloned pieces of DNA). Many breeding populations are usually available, but they may have inadequate molecular polymorphism, and it is important to measure the amount of polymorphism at the outset of any mapping program. Molecular polymorphism in Capsicum has been detected for chromosomes (Pickersgill, 1977), isozymes et al., (Jensen et al., 1979 and Loaiza-Figueroa et al., 1989), soluble seed protein gel mobility (Panda et al. 1986), and AFLPs (Prince et al., in preparation). The use of these measures of variation (except for RFLPs) and of classical taxonomic characters has been reviewed by Pickersgill (1988).

We present here a brief overview of the history of the development of the genetic map of Capsicum and the current status of some molecular mapping projects.

Molecular maps constructed in pepper

In 1984, Tanksley published the first molecular map of pepper based on the segregation data of fourteen isozyme loci. Four linkage blocks were detected, and five hybrid primary trisomics made by Pochard (1970) were used in dosage analysis to determine the chromosomal locations of three of the isozyme loci. Data suggested that three of the markers were near the site of a reciprocal translocation between the two mapping parents.

Because of the relatively small number of available isozyme markers and the even smaller number that segregate in any given cross, Tanksley et al (1988) decided to use RFLPs, choosing a cDNA library constructed from whole tomato leaf mRNA (Bernatzky & Tanksley, 1986). These clones represented expressed sequences in the genome. A package map consisting of 80 molecular markers in fourteen linkage groups was constructed using an interspecific backcross mapping population. Major results from this preliminary mapping study indicated that pepper and tomato shared the same gene repertoire in that all of the cDNA clones from tomato hybridized to pepper. The order of genes on

the chromosomes were different between the two genera. Based on a comparison of the linkage maps, a minimum of 32 chromosome breakage events had occurred since the pepper and tomato lineages diverged. Some large linkage blocks remain conserved, however, with the largest being 63 cM long. By contrast, the molecular maps of potato and tomato differ by only a few inversions (Bonierbale et al., 1988 and Tanksley et al., in preparation).

In order to improve the linkage map of pepper, we (Prince, Pochard, and Tanksley, in preparation) have constructed a new map using an interspecific F₂ mapping population. The use of an F₂ population allowed improved resolution with a comparable population size. Random tomato genomic clones (Miller & Tanksley, 1990b) and cDNA clones from tomato leaf epidermal tissue (Yu et al., unpublished) were placed onto this map in the hopes of coalescing the map into twelve linkage groups, corresponding to the haploid chromosome complement of pepper (Pickersgill, 1977). Presently (February 1992), eleven large linkage groups and eight small linkage groups have been formed using one hundred and ninety-two molecular markers. The average density of the map is approximately one marker for every 3.5 cM. Again, similarities between the tomato and pepper maps are evident, with large linkage blocks conserved and considerable chromosomal rearrangements that may cluster around centromeric regions. The random tomato genomic clones did not hybridize as readily to pepper as did the cDNA clones, with only about half of them being usable because of low homology or increased copy number in pepper.

Following are the strategies that we are using to attempt to coalesce the map into the expected twelve linkage groups.

Improving the RELP map of pepper.

Random genomic pepper clones. We are currently developing a library of pepper genomic clones to use in more mapping (Blauth et al., unpublished). Based on data from our current mapping work, we have noticed that clones from several chromosome arms of tomato do not hybridize to pepper and that we may be missing corresponding regions of the pepper chromosomes (Prince, Pochard, and Tanksley, in preparation). These regions may be undergoing more rapid sequence divergence and should be detectable with the use of clones from the same species.

PCR-based markers. Gaps in our molecular map may be due to regions of rapid evolution, as noted above, or by stretches of repetitive DNA for which DNA clones cannot be used as markers. The RAPD technique (random amplification of polymorphic DNA, Williams 1990, and Welsh & McClelland, 1990) a polymerase chain reaction-based mapping technique, may allow us to pick up some of these regions. Ample polymorphism is detected and currently we are adding markers to the map (Prince et al. unpublished).

Pulling more clones off of the high-density tomato map. Now that a high density RFLP map of tomato has been completed with over 1000 markers (Tanksley et al., in preparation) we have a much larger pool of tomato clones to draw upon from all regions of the genome. We will target underrepresented regions to see if we can close the gaps between some of our linkage groups.

Current applications of molecular mapping in pepper

We are currently working on mapping the genes for resistance to some of the most devastating viruses affecting pepper: cucumber mosaic virus (CMV), tobacco mosaic virus (TMV), and the related potyviruses pepper mottle virus (PeMV), tobacco etch virus (TEV), and potato virus Y (PVY). Classical breeding to produce multiple virus-resistant plants remains difficult because of the multigenic nature of some of the resistance genes. The difficulty in scoring the phenotype and the difficulty in monitoring more than one virus in a given plant at the same time. We hope to tag resistance genes from several sources to these viral diseases with molecular markers and use these markers to facilitate the breeding of superior lines. .

CMV High levels of resistance have been located in two accessions of *C. frutescens* from southern Mexico by loaiza-Figueroa and Prowidenti (unpublished). Resistance to CMV has been difficult to transfer despite reports of resistance in *C. annuum* cv. 'Perennial' and *C. frutescens* (Loaiza-Figueroa and Prowidenti, unpublished) We are working with several inter- and intra-specific populations in an effort to locate markers linked to resistance from these sources. Segregating populations are being screened for resistance and are being

mapped with our framework of seventy RFLP markers spaced at 5-cM intervals along the map of pepper.

TMV The mapping population used to construct our most recent map of

192 markers also segregates for TMV resistance. Screening of resistance in F3 families has just finished and data are being analyzed.

Potviruses. A new population that is polymorphic at the DNA level and - that segregates for resistance to the potyviruses has been grown and is now being screened for resistance and being mapped with the same set of framework clones as are the CMV populations (Blauth et al., unpublished). In addition, a pepper genomic library has been constructed which should cover areas of the genome that are not well-represented with the libraries used to date.

Work by other groups in mapping projects in pepper. Two other groups are currently pursuing major mapping efforts in pepper. Lefebvre, Palloix, Pochard, and Rives at the Station d'Amelioration des Plantes Maraicheres, INRA, Avignon, France are currently working on resistance genes for several important fungal and viral diseases, and Massoudi, O'Connell, and Bosland at New Mexico State University in Las Cruces, New Mexico, United States, are currently working on Phytophthora resistance.

Coordinating mapping efforts. An effort is underway to coordinate mapping in the genus, and all interested workers are encouraged to contact Molly Kyle. Inter-institutional mapping efforts have been organized for a number of crops and have been especially successful in cereals. We hope that any groups interested in developing programs in this area and contributing to a collective effort will join this consortium, to be organized at the National Pepper Conference in August, 1992, in Monterey, California, USA.

Conclusion

While mapping efforts are still in progress we are very optimistic that the map will develop quickly. A concerted effort is now possible thanks in part to support through the United States Department of Agriculture National Research

Initiative for the Plant Genome. The availability of the high-density tomato RFLP map offers a number of advantages in the construction of a map of pepper and in resolving the twelve linkage groups (Tanksley et al. submitted). The tremendous number of horticultural types in pepper and the difficulty in achieving the most important breeding objectives, despite considerable investment by workers around the world, suggest that marker-assisted selection may be especially useful in Capsicums.

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INDUCED DOUBLE TRISOMIC IN CHILLI Capsicum annuumL.

L.D.Meshram, B.K. Kukade. M.B.Wadodkar, and R.A.Naphade.

Cotton Research Unit, CRS, Punjabrao Krlshi Vidyapeeth,

Akola-444 104, [M.S.] India. '

A double trisomic $[2n + 1 + 11]$ chili plant was recovered 30 Krad dose of gamma irradiated population of chilli cultivar Pusa Jwala' in R^1 generation.

The plant was identified due to its distinct difference. In its morphological characters as compared to normal untreated, plants. Morphologically the plant was having dark green and leathery leaves, big flowers having 70-80 per cent pollen sterility. The plant bore the fruit which were long shriveled in appearance. The fruits turned red in colour ripening and were having 3-4 seeds per fruit as against 30-40 seeds in control.

The flower buds from the plant were fixed in acetic alcohol [1:3] and squashed in 1% aceto-carmin stain. At diakinesis 10 bivalents and two trivalents and $12^{11} + 2^1$ were observed. The trivalents showed three [pan, chain and y-shape] configurations. At metaphase 1 out of 1100 PMCs studied 45 showed $12 + 2$ and quadrivalent was noted indicating that both the univalents present in the plants were of different types. At anaphase 1 the chromosome disjunction was 13-13, 12-14 and 11-15, while in some PMC univalents remained as laggard. Meshram et al. [1980, 81] also reported that radiation induced trisomic and triploid in chilli.

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INHERITANCE OF ANTHOCYANIN PIGMENTATION IN CHILLI (Capsicum annum L.)

Nazeer Atuned and H. I. Tanki Division of Olericulture, Sher-e-Xashmir University of Agricultural Sciences and Technology, Shalimar campus, Srinagar - 191121 (India)

To minimize varietal deterioration in a developed variety there is a need to have a marker to isolate true to type plants in seed production programmed. Since anthocyanin pigment which imparts purple colour in plant may be used as the marker for effective varietal identification, it is therefore important to know the type and number of gene (8) responsible for the trait.

Genetics of pigmentation was studied from six generations . (P1, P2, 1'1, 1'2,BC1 and BC2) ,of an intervarietal cross SPE-1 (Purple stem) x 'Shalinar Long (Green stem) during kharief 1989 at Vegetable Experimental Farm, Sher-e-Kashmir University of Agricultural Sciences and Technology, Shalimar, Srinagar, India. The anthocyanin pigmentation from the stem of each plant' was determined following the method as described by Swains and Hills (1959).

The mean anthocyanin content in F1 was slightly more than the mid parent value indicatino partial dominance of purple stem over green stem (Table 1). The continuous variation observed in F2 and Backcross generations suggested its polygenic nature. The simple additive-dominance model was inadequate, thereby indicatino the presence of non-allelic interactions. The six parameter model which included interaction components revealed only components additive(d) and dominance x dominance(l) to be significant. However in both perfect fit solution and 8imple additive-dominance model, the magnitude of adaitive component was much higher (Table 1) indicating the Importance of additive gene action in the inheritance of anthocyojn pigmentation. This fixable addition component can therefore be exploited through simple selection.

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Table 1: Mean anthocyanin content (mg/g) of different generation and content of generation means of cross SPE –1x Shelimar long

Means (mg/m)	Three parameter Medal	Six parameter model
P1=1.32 = 0.015	M=0.77 + 0.008	M= 0.88= 0.135
P2= 0.28+ 0.0010	[d]= 0.48+0.008	[d]= 0.52+0.009
F1= 0.88+ 0.016	[h]= 0.02+ 0.017	[h]= 0.47+ 0.310
F2 0.75+ 0.029		[I]= 0.08+ 0.134
B1= 0.97+0.025		[j] = 0.06+0.069
B2= 0.48+0.021	(d.f)= 37.86	[I]= 0.43+0.182

** Significant at 5% and 1 % level respectively

A giant chilli plant as a transgressive variant of

Capsicum Frutescens

N. Lakshmi, T. Srivalli and V.V ramachandra Rao

Cytogenetic Laboratory, Department of Botany, Nagarjuna University, Nagarjuna Nagar -
522510 Guntur Dt. (A.P.) India

Capsicum Frutescens popularly called 'seema mirapa' in Andhra Pradesh is a highly pungent species with very small fruits and maximum capsaicin content. Based on fruit colour, three cultivars are recognized namely green, white and yellow, the first two are immature fruit colours while the third represents fruit colour at maturity.

All the three cultivars are erect, tall, vigorous with good spread and big green ovate leaves. Flowers are small with greenish white corolla and bluish green anthers. Fruits are erect, tiny, spindle-shaped, green or white when immature turning into yellow or red at maturity depending on the cultivar.

Intraspecific hybridizations have been made in this laboratory in six combinations involving the three cultivars in order to know the compatibility and cytogenesis relations of the three cultivars. In the hybrid progeny of the green and white cultivars a very tall chili plant with very good spread, profuse branching and high yield was noticed in the year 1990. The morphometrics of this plant along with the two parental cultivars is set out in Table 1. Data revealed that there is three to four fold increase in plant height, spread, number of branches and chlorophyll content, two fold increase in leaf area, and 5-6 fold increase in yield than those of the parental species. However there is decrease in fruit girth and ascorbic acid content. Cytological studies revealed no appreciable change in meiotic metrics like Chiasma frequency per cell, per bivalent and percentage of ring and rod bivalents. Progeny studies revealed the recurrence of green and white cultivars suggesting that the parent plant is a hybrid involving the two cultivars. Since the hybrid exceeds the parents in several parameters it is considered as a transgressive variant formed as a result of recombination of genetic factors.

TABLE 1. Morphometrics of parents and giant chili hybrid				
S.No.	Character	C.f.g.	C.f.w.	Giant Hybrid
1.	Plant Height (cm)	59.60	61.33	205.00
2.	Plant Spread (cm)	89.00	83.67	290.00
3.	No. of branches per plant	149.00	112.00	448.00
4.	Leaf length (cm)	7.50	7.14	10.50
5.	Leaf breadth (cm)	2.80	2.65	3.42
6.	Leaf area (cm)	8.65	8.49	18.80
7.	No. of fruits per plant	125.00	148.00	750.00
8.	Fruit length (cm)	6.00	5.17	6.02
9.	Fruit girth (cm)	3.20	3.10	2.22
10.	Chlorophyll content mg/gm	2.306	2.071	6.500
11.	Ascorbic acid in unripe fruits (mg/100gm)	104.00	78.00	40.00
12.	Ascorbic acid in ripe fruits (mg/100gm)	161.20	150.80	64.00
13.	Chiasma frequency per cell	20.13	20.50	20.27
14.	Chiasma frequency per bivalent	1.68	1.67	1.69
15.	Percentage of ring bivalents	67.78	65.08	69.32
16.	Percentage of rod bivalents	32.22	34.92	30.68
C.f.g = <i>Capsicum frutescens</i> (green)				
C.f.w = <i>Capsicum frutescens</i> (white)				

Induction of Callus and Organogenesis In Pepper R. Pundeva and N. Simeonova
Institute of Genetics, BAS, Sofia 1113, Bulgaria

Donor material – *Capsicum annuum* F. Macrocarpum (cv. 'Albena') *C. annuum* F. microcarpum (No 9), *C. baccatum* var. Pendulum (F. Smith) Cultured explants – hypocotyl, cotyledon and root segments of in vitro grown seedling in cotyledon stage.

Nutrient media- PNS medium enriched with different supplements (Table 1). Growing conditions – temperature 25°C photoperiod – 16 h light. Table 1 shows the morphogenetic response of the explants treated with different hormone concentration. NAA induced direct root formation, sometimes attended with slight callus initiation. KEW with Bap stimulated shoot bud development. 2,4-D combined with KIN evoked abundant friable callus growth, while media containing IAA and Bap promoted the appearance of compact callus and differentiation of shoots and roots on it. Basal Hypocotyl segments has higher predilection to rooting, while middle and apical ones preliminary produced shoots. This tendency significantly decreased on more effective media (NO6,7,8) whole hypocotyl some. Three sectioned cotyledons also demonstrated different morphogenetic response. Higher callus and organogenic capacity has basal segment relatively good capability showed middle ones and less productive were apical sections.

Root explants possessed similar tendency. Basal segments were more effective regarding the rest of root parts but were comparatively less active towards the other tested organs.

The genotypes included in this study showed difference in the speed of their reaction to the corresponding nutrients media with respect to the callus mass and the number of meristematic buds per segment. *C. baccatum* var. pendulum was more effective than *C. annuum* accessions.

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Table 1. Morphogenetic response of meristematic explants to MS medium with different combinations of supplements (mg/l).

Hormones	NAA 0.5	IAA 0.5	BAP 2.0	DROP 0.5	DROP 2 IAA 0.5	IAA 1 BAP 2	IAA 1 BAP 5	IAA 0.5 BAP 1	2,4D 0.5 KIN 0.5	KYN 1 BAP 3
Explants	1	2	3	4	5	6	7	8	9	10
cv. 'Albena'										
hypocotyl	R	C	C	C	C	C+R	C+R	C+R	C	C+R+S
cotyledon	R	C+R	C	C	G+S	C+S	C+S	C+R	C	C+S+R
root	R	C+R	C+R	C+R	C+R	C+R	C+R	C+R	C	C+R
No 9										
hypocotyl	R	C	C		C	C+S+R	C+S+R	C+R+S	C+R	C+R+S
cotyledon	R+C	C+R	C	C	C+S	C	C+R+S	C+R+S	C	C+S
root	R	C+R	C	-	-	C	C+R	C+R	C+R	C+R
C. pendulum										
hypocotyl	R	C	C	C+S	C	C+S+R	C+S+R	C+S+R	C+S	C+S+R
cotyledon	R	C+R	C	C	C	C+S+R	C+S	C+S+R	C+S	C+S+R
root	R	C+R	-	C	C+R	C+S+R	C	C+S+R	C+S	C+S+R

C = callus; R = roots; S = shoots

IN VITRO ANDROGENESIS IN CUBAN F1 HYBRIDS OF SWEET PEPPER

O. Gonez and D. Chambonnet

1. LIH "Liliana Dimitrova", Carr. Quivican km 331/2 La Salud, Habana, Cuba .
2. Station d'Amelioration des Plantes Maraicheres, BP 94, 84143 Montfavet, France 2

During the summer of 1989 the technique proposed by Chambonnet in 1988 for the obtention of haploid plant through in vitro androgenesis was applied to seven cuban F1 hybrids of sweet pepper. The plants were grown in the field, no phytosanitary treatment was applied for aphyds control. The floral bottoms were removed in August at the morphological state in which the length of the petals and the sepals were equal, corresponding to the beginning of the first pollinic mitosis. The anthers were put on the induction medium (P) recommended in the technique; it was enriched with ampicillin (10 mg/l), ampicillin plus streptomycin (10 mg/l) or rifocin (5 mg/l).

The Petri dishes containing the anthers were placed at +35°C during 1-2 or 8 days in the darkness and then transferred .at +25°C and 12 hours of photoperiod; the technique was continued as e.stablished. We could prove feasibility of using in vitro androgenesis from cuban hybrids. Timing for treatment at +35°C and darkness can be variable, from 1-2 to 8 days according to genotype.

A preliminary field trial was caried out in Cuba in 1991 including 19 doubled-haploid (DH) lines, outstanding for compact plant habit, earliness, productivity and disease tolerance.

REFERENCE

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A Rapid Method For the Screening of Salinity resistance in Red Pepper Varieties
N. LAKSHMI, V.V. RAMA CHANDRA RAO and T. Srivalli, Cytogenetics Laboratory,
Department of Botany, Nagarjuna University, Nagerjuna Nagar 522 510 Guntur Dist. (A.P.)
India

Determination of salinity resistance is one of the thrust areas in any crop improvement programmes and there is greater need for the development of varieties resistant to saline soils. So conducting the field trials a rapid screening can be done by testing salinity resistance on the basis germination counts. In the present study an attempt has been made experimentally to screen some red pepper varieties or salinity resistance.

Seed materials of five different varieties of *Capsicum annum* viz., Santaka, 197, 960, jawahar and 235 and one of *Capsicum Chinese* were used in the study of NaCl salinity effect on seed germination.

The seeds of these varieties were soaked in NaCl solution in different concentration (1-10%) at four time intervals (6, 12, 18 and 24 hours) Control were maintained for all samples by soaking them in distilled water. After Treatment the seeds were germinated in petriplates. The percentage interval in record.

Data revealed that in all varieties germination percentage is inversely related to increase in concentration.

Capsicum Chinense and Santaka variety of *Capsicum annum* showed high salinity tolerance since the germination good even at higher concentrations (10%) and longer duration of soaking 24 hours Table I. Hence these two varieties are considered to be showing higher salinity resistance. However their stress tolerance is to be tested under field condition also.

The remaining four varieties are considered to be salt sensitive ones since their germination percentage is affected greatly at higher concentrations of NaCl.

TABLE 1. Percentage seed germination of some red pepper varieties under the influence of NaCl salinity

Name	Period of treatment (hours)	Concentration of NaCl									
		C	1%	2%	3%	4%	5%	6%	8%	10%	
<u>C. annuum</u> variety '960'	6	92	96	92	84	80	84	72	68	68	
	12	88	88	80	84	80	76	76	68	60	
	18	92	84	92	84	80	76	72	72	64	
	24	96	88	84	76	76	72	72	68	60	
<u>C. annuum</u> variety 'Jewehar'	6	100	100	96	96	88	88	76	72	68	
	12	96	96	92	84	84	80	76	76	72	
	18	92	92	88	88	84	76	76	72	68	
	24	92	88	88	84	84	80	76	68	64	
<u>C. annuum</u> variety 'Santaka'	6	100	100	100	92	88	88	84	80	80	
	12	100	96	92	92	84	84	80	76	80	
	18	92	92	88	88	84	84	80	76	76	
	24	92	92	84	88	84	80	80	80	76	
<u>C. annuum</u> variety '197'	6	96	88	88	84	80	72	76	72	68	
	12	96	96	92	84	84	80	72	68	64	
	18	92	88	88	80	76	76	72	68	60	
	24	76	80	80	72	76	72	68	64	60	
<u>C. annuum</u> variety '235'	6	96	96	92	84	88	84	80	72	68	
	12	92	88	88	80	84	76	76	72	64	
	18	92	84	88	84	84	72	72	68	60	
	24	88	88	84	76	76	72	68	64	60	
<u>C. chinense</u>	6	100	96	92	96	92	88	84	80	80	
	12	100	100	100	96	96	92	88	84	80	
	18	96	88	88	84	76	80	76	76	72	
	24	92	92	84	84	80	76	76	72	72	

BIOLOGICAL CHARACTERIZATION OF SPANISH ISOLATES OF TOBAMOVIRUS.

M. Luis Arteaga and R. Gil Ortega

S.I.A. - D.G.A., Apartado 727, E-50080 Zaragoza, Spain

Characterization of nine tobamovirus isolates collected in Spain from pepper (seven) and tomato (two) was accomplished by studying the responses of Capsicum spp. genotypes which carry the L+, Ll, L2, L3 and L4 resistance genes (BOUKEMA, 1983)

According to biological reactions on a series of test plants (Table 1), three of the nine isolates were classified as ToMV (Tomato Mosaic Virus) while the other six belonged to the PMMV (Pepper Mild Mottle Virus). On the basis of the interactions with the L gene for resistance in pepper genotypes we have found that five of those PMMV isolates belonged to the PI-2 pathotype and the sixth one to the PI-2-3 pathotype (Table 2).

In Spain the number of commercial F1 hybrids which carry the L3 allele is still very low. That would be the reason for the scarcely proportion of PI-2-3 pathotype found in this work and by other researchers in Spain (ALONSO et al., 1989; CUADRADO, 1991, personal communication).

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Table 1. Response of a set of test species to nine tobamovirus isolates.

Test species	Virus isolates								
	T-1-75	T-5-84	P-3-85	P-10-85	P-11-85	P-12-85	P-8-88	P-9-88	P-3-90
<i>Chenopodium amaranticolor</i>	lln/o	lln/o	lln/o	llcn/o	llcn/o	llcn/o	llcn/o	llcn/o	llcn/o
<i>Chenopodium quinoa</i>	lln/o	lln/o	lln/o	llcn/o	llcn/o	llcn/o	llcn/o	llcn/o	llcn/o
<i>Datura stramonium</i>	lln/o	lln/o	lln/o	lln/o	lln/o	lln/o	lln/o	lln/o	lln/o
<i>Gomphrena globosa</i>	---	---	---	---	---	---	o/o(-)	---	o/o(-)
<i>Lycopersicon esculentum</i> 'Montserrat'	o/No	---	---	o/o(-)	---	---	---	---	o/o(-)
<i>Nicotiana clevelandii</i>	o/NoSt	o/NoSt	o/NoSt	o/No	o/No	o/No	o/No	o/No	o/No
<i>N. glutinosa</i> , <i>N. tabacum</i> 'Xanthi nc'	lln/o	lln/o	lln/o	lln/o	lln/o	lln/o	lln/o	lln/o	lln/o
<i>N. tabacum</i> 'Paraguay'	lln/o	lln/o	lln/o	lln/o	lln/o	lln/o	lln/o	lln/o	lln/o
<i>N. tabacum</i> 'Samsun'	o/No	o/No	o/No	o/o(+)	o/o(+)	o/o	o/o	o/o	o/o(-)
<i>N. rustica</i>	---	---	---	lln/o(-)	---	---	---	---	lln/o(-)
<i>Petunia hybrida</i> 'Rose du Ciel'	lln/o(-)	lln/o	lln/o	o/o	---	---	---	---	o/No
<i>Physalis floridana</i>	o/No	o/No	o/No	o/No	o/No	o/No	o/No	o/No	o/No

Legend: Local reaction/Systemic reaction

a = Abscission

c = Chlorotic

cn= Chloronecrotic

ll= Big local lesion (2-3 mm)

ll= Small local lesion (1 mm)

Mo= Mosaic

n = necrotic

St = Stunting

Tn = Top Necrosis

o = No reaction

---= Non-inoculated species

(+)= Positive infection by backinoculation

(-)= Negative infection by backinoculation

T = Tomato

P = Pepper

Table 2. Response of *Capsicum* spp. genotypes to nine tobamovirus isolates.

Test species	Virus Isolates								
	T-1-75	T-5-84	P-3-85	P-10-85	P-11-85	P-12-85	P-8-88	P-9-88	P-3-90
<i>Capsicum annuum</i> 'Doux des Landes'	o/No	o/No	o/No	o/No	o/No	o/No	o/No	o/No	o/No
<i>C. annuum</i> 'Tolo Wonder'	llna/o	llna/o	a/o	o/No	o/No	o/No	o/No	o/No	o/No
<i>C. frutescens</i> 'Tabasco'	llna/TN	llna/TN	a/TN	o/No	o/No	o/No	o/No	o/No	o/No
<i>C. chinense</i> F.I. 159236	lln/o	lln/o	lln/o	lln/o	lln/o	lln/o	lln/o	lln/o(-)	o/No(+)
<i>C. chacoense</i> F.I. 260429	lln/o	---	---	a/o	---	---	a/o	---	lln/o

DEVELOPMENT OF A SCREENING TECHNIQUE FOR EVALUATION OF RESISTANCE TO PSEUDOMONAS SOLANACEARUM IN PEPPER K.D.A. Pererao, G.L. Hartman+, j. M.

Poulos+

°Central Agricultural Research Institute, P.O. Box 11, Peradeniya, Sri Lanka

+Asian Vegetable Research and Development Center, P.O. Box 42, Shanhua, Tainan, Taiwan 74199, R.O.C.

Bacterial wilt caused by Pseudomonas solanacearum is an important disease of pepper (Capsicums spp.), especially sweet pepper, under certain tropical conditions. There is little information about the extent of damage caused by this disease. Resistant varieties offer an effective means for its control. Development of a standardized technique in evaluating pepper germ plasm is necessary, as uniform field epidemics are often difficult to create for pepper crops. Three inoculation techniques toothpick stabbing with a bacterial culture, soil drenching with root wounding and soil drenching without root wounding - were tested using 20- and 35- day old seedlings with two isolates (PS 71 and PS 4) of f solanacearum. Soil drenching was applied with a 107 ceil/mi inoculum suspension of the bacterium. The susceptible variety 'Giant Bell (F1)' was used. Toothpick stabbing and soil drenching with root wounding gave a similarly high percentage of wilt (Fig. 1). The root wounding technique, however, was relatively easy to apply and resembled natural inoculation most. The percentage wilt was not significantly affected by the age at which the plants were inoculated. Although there was no significant difference in percentage wilt caused by the two strains, the pepper isolate PS 71 was slightly more aggressive than the tomato isolate PS 4 (Fig. 2). Other studies to confirm resistant genotypes using the same inoculation techniques are in progress.

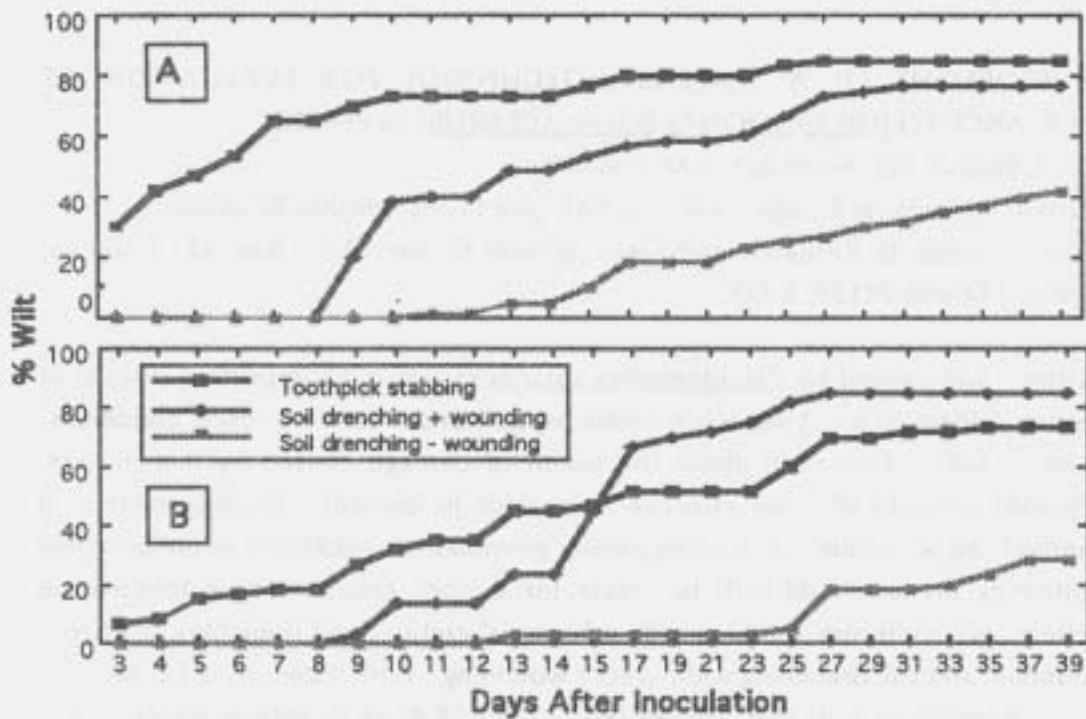


Fig. 1. Bacterial wilt on 20- (A) and 35-day old peppers (B) inoculated with three inoculation techniques.

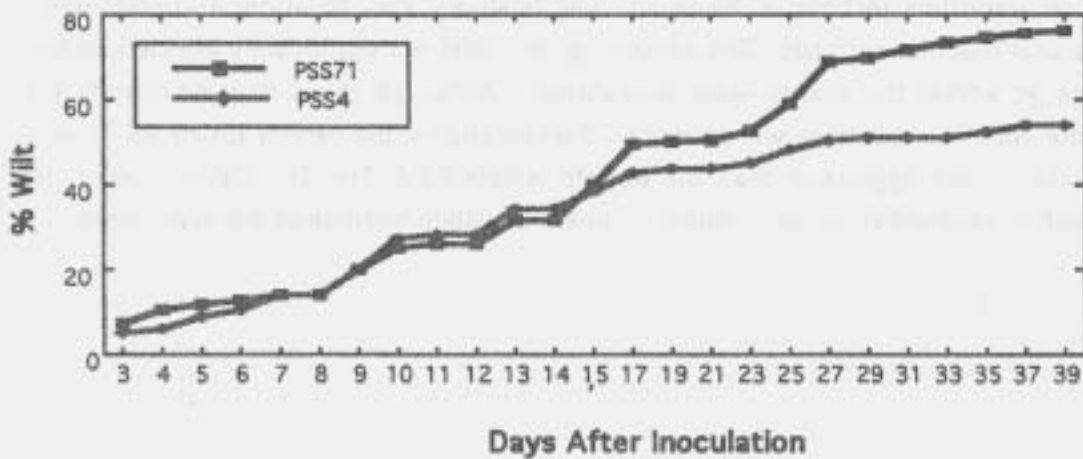


Fig. 2. Percentage bacterial wilt combined over 20- and 35-days and over inoculation techniques for two strains of *Pseudomonas solanacearum*.

SOURCES OF RESISTANCE AMONG CAPSICUM SPP. TO FUSARIUM WILT OF PEPPER
M. M. Jones and L. L. Black, Dept. of Plant Pathol. and Crop Physiol., La. Agric. Expt. Sta., La'. State Univ. Agric. Ctr., Baton Rouge, Louisiana 70803, U.S.A.

Fusarium oxysporum f. sp. capsici (FOC) was recently described as the causal agent of a vascular wilt of pepper. The pathogen was isolated from wilted 'Tabasco' Capsicum Frutescens plants and from soil from production fields at Avery Island, Louisiana. Occurrence of the disease is associated with periods of high rainfall, and is more prevalent in poorly drained areas of the fields. Disease symptoms in the field include flagging of the leaves, slight yellowing of the foliage, vascular discoloration of the roots and crown, and a permanent wilt of the plant. Isolates were obtained by plating surface sterilized tissue on acidified PDA, or air dried field soil on Komada's selective medium. Plant inoculation studies were conducted to confirm pathogenicity of isolates. Inoculum consisted of two 7-day-old cultures propagated by streaking conidia on PDA in 10cm petri plates and incubating at room temperature under continuous fluorescent light. Contents of both plates were blended for about 10 seconds in a Waring blender with 250ml of distilled H₂O to form a slurry consisting of conidia, mycelium, and agar bits. Two to 3-wk-old seedlings of two susceptible cultivars, 'Yolo Wonder', and 'Tabasco', and a resistant accession, PI 188803, representing C. annum, C. frutescens, and C. baccatum, respectively, were root dip inoculated (3.5ml in the slurry). The plants were immediately transplanted to planter flats with 5-cm cells, Todd Planter Flats, Model 200 (Speedling, Inc., Sun City, FL), containing a combination of peatlite soil mix and pasteurized sand 3:2 (v/v). Plants inoculated with different isolates were kept separate on greenhouse benches to avoid cross contamination from water splash. Greenhouse temperatures were maintained at 28:±4C. The same inoculation procedure was used to assess the response of Capsicum accessions to a highly virulent isolate, PS-1 a, of f.Q.C.; Assays to identify wilt resistance among accessions of several species of Capsicum showed that high levels of resistance occur in baccatum. Of the baccatum accessions assayed, 16 of 27, were homogeneous in their response to inoculation and showed no wilt and little or no evidence of root damage. Among accessions of C. annum, C. chacoense, C. frutescens, C. microcarpum, and C. pubescens tested, single accessions of C. annum and C. chacoense were found to be highly resistant. Plants of most other accessions tested among these species were highly susceptible, but plants of a few accessions survived inoculation while showing stunt and root damage, indicating an intermediate type reaction. All accessions of C. frutescens tested were highly susceptible. Using nitrate nonutilizing mutants (nit), all pathogenic isolates of FOC were found to be in the same vegetative compatibility group. No compatibility was observed when nits of nonpathogenic isolates from tissue and soil were paired with the pathogenic nit testers. Isolates of EQ.C (A TCC 66420/66421) may be obtained from the American Type Culture Collection, 12301 Parklawn Drive, Rockville, Maryland 20852.

LITERATURE

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Rivelli, V. 1989. A wilt of pepper incited by Fusarium oxysporum f. sp. capsici forma specialis nova. M. S. Thesis. Louisiana State University. Baton Rouge. 71 pp.

Table: Capsicum accessions resistant to Fusarium oxysporum f. sp. Capsici isolate PS-1a (ATCC 66420)

Capsicum annuum (62 accessions tested)

C00758

Capsicum chacoense (8 accessions tested)

C01553

Capsicum baccatum

PI 188803

PBC 151

C00302

C00313

C00313-1

C00475

C00754

C00755

C00774

C00803

C00941

C01218

C01300

C01526

C01527

C01543

Genetic Resources and Seed Unit, Asian Vegetable Research and Development Center, P.O. box 42, Shanhua, Tainan 74199, Taiwan, ROC

USDA-ARS South Atlantic Area, Regional Plant introduction Station, Experiment, Georgia 30212, USA

TEST FOR RESISTANCE TO VERTICILLIUM DAHLIAE KLEB. IN CAPSICUM ANNUUM L.

J. Barriuso Vargas, R. Gil Ortega and C. Palazon Espanol
S.I.A. - D.G.A., Apartado 727, 50080, Zaragoza, Spain

In 1991 a test for resistance to *V. dahliae* on *Capsicum annuum* was made. Ten genotypes reported-by several authors as resistant to that parasite were included. 'Yolo Wonder' and the Spanish pimiento type culti.var 'R,iguel' were used as susceptible controls (Table 1).

The roots of 9-week-old seedlings were washed and then dipped into a suspension of 3.2×10^6 propagules/ml for 3 minutes (KENDRICK and MIDDLETON, 1959). Fifteen to thirty seedlings per variety were transplanted into pots containing a pasteurized mixture of 1 Humin-Substrat, 1 peat, 1 perlite, 1 sand, 1 clay-loam soil (by volume). Pots were placed on greenhouse benches where average of minima and maxima air temperatures were 13°C and 30°C.

Plants were evaluated for *Verticillium* resistance by weighing their aerial parts three months and a half after inoculation. In order to establish a index which could be regarded as an estimation of resistance (IR), each plant weight was referred to the average weight of six to ten non-inoculated plants of the same variety. Data was treated by analysis of variance.

It was difficult to establish statistical differences between genotypes, with the exception of 'Yolo Wonder' which behave as the most susceptible one (Table 1). Within the rest of genotypes 'Riguel' and 'Buketén' as the more susceptible ones and 'T1-1' and 'New Mexico Accesion' as the more resistant can be pointed out.

These results obtained by artificial inoculation on greenhouse do not exactly agree with the response of same genotypes grown in *V. dahliae* highly infected fields. Classification of varieties according to their IR is in some way influenced by the inoculation method used and by the climate where the assay is made (PALLOIX et al., 1990).

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Recurrent selection for resistance to *verticillium dahliae* in pepper. *EuphytJ.ca* 47: 79-89.

Table 1. Genotypes of Capsicum annuum ordered by resistance (IR) to Verticillium dahliae.

Genotype	IR (0-100)	Source and/or Reference
Yolo Wonder	26.9 a	'Bell' type Cultivar (USA)
Riguel	59.6 b	Pimiento type cultivar (Spain)
Buketen	69.1 bc	Christov and Popova (Bulgaria)
Podarok Moldavii	76.8 bcd	Lobb (USSR)
Luesia	82.3 bcd	Gil Ortega (Spain)
Serrano C.M. 334	85.3 bcd	Guerrero (México) Palloix <u>et al.</u> (France)
C-169 (<u>C. minimum</u>)	88.5 bcd	IVT (The Netherlands)
Pallagi (WYR4618)	92.6 cd	Inst. Vavilov (USSR)
L-25	93.0 cd	Marinkovic <u>et al.</u> (Yugoslavia)
Line 724	93.1 cd	Aleksic <u>et al.</u> (Yugoslavia)
New Mexico Accession	96.0 d	Bosland (USA)
T1-1	97.3 d	Marinkovic <u>et al.</u> (Yugoslavia)

Means separations according to Newman-Keuls test, level 5%.

CHARACTERIZATION OF VERTICILLIUM DAHLIAE KLEB. ISOLATES ON A SET OF Capsicum SPP GENOTYPES.

J. Barriuso Vargas, C. Palazon Espaftol, R. Gil Ortega and I. Delgado Izquierdo.

S.I.A. - D.G.A., Apartado 727, 50080, Zaragoza, Spain

The pathogen behaviour of nineteen verticillium dahliae Kleb. isolates on five Capsicum genotypes during two years has been evaluated. The 19 isolates were evaluated in 1989 while only the 5 most representative were considered in 1990. The isolates had been obtained from diseased plants of pepper. To carry out the assays two-pepper varieties, which are susceptible to *V. dahliae* ('Yolo Wonder' and 'Riquel') and three partially rest Stan genotypes ('Podarok Moldovy', 'Luesia' and *C. minimun* 'C169') have been used.

The roots of 9-week-old seedlings were washed and dipped into a suspension of 3.2×10^6 propagules ml for 3 minutes (KENDRICK and MIDDLETON, 1959). Five seedlings per variety in 1989, and 8 seedlings in 1990 were transplanted to pots containing a pasteurized mixture of 1 Humin-Substrat, 1 peat, 1 perlite, 1 sand, 1 clay-loam soil (b I volume). Pots were placed on greenhouse benches where average a r temperatures ranged from 17°C to 22°C in 1989 and 18°C to 26°C in 1990.

Plants were evaluated for Verticillium resistance by weighting their aerial parts two months after inoculation in 1989, and four months in 1990. In order to establish a percentage which could be regarded as an index of resistance, each plant weight was referred to the average weight of five non- inoculated plants of the same variety in 1989, and eight in 1990~ The data was treated by analysis of variance.

1989 and 1990 trials showed that only six isolates were able to differentiate between pepper genotypes which generally speaking, were classified by resistance as expected. Nevertheless, some of the interactions found in 1989 could not be confirmed in 1990. The non-existence of a clear host-parasite cx interaction joined to the slight importance of some specific c- reactions did not permit the definition of vertical pathotypes. Nevertheless, isolates could be ranked by order of aggressiveness. Data from 1990 experiment are shown in Table 1.

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Table 1. Average percentage of fresh weight of <i>V. dahliae</i> inoculated pepper plants referred to controls					
		Pepper Genotypes			
<i>V. dahliae</i> Isolated	Yolo W.	Riguel	Podarok	Luesia	C-169
VD08	16,2a	12,2a	37,3a	25,1a	67,8a
VD 13	8,9a	13,0a	36,7a	29,1a	77,0ab
VD05	13,6a	14,5a	31,0a	32,5a	72,8ab
VD12	54,5ab	48,6ab	62,9a	67,7b	111,1b
VD03	68,3b	71,2b	74,4a	103,6b	42,3a

Mean separation by Newman-Keuls test, 1% level (for the graphic expression of such separation, letters in the vertical direction and straight lines in the horizontal direction were used).

RESULTS OF SEED TESTS. XIV. - ON GERMINATION PARAMETERS REGISTERED FOR SEED SAMPLES OF CAPSICUM ANNUUM L.

Endre I. SIMAY* and Zsuzsanna HORVATHI ++

I.A.Q. Research Centre for Agrobotany, Tapioszele

Present address: +; Enterprise for Ext. and Res in Fruit Grow. and Ornaments, B-1223 Budapest, Park 2. ++; Univ. of Veterinary Sci., Dept. Botany, B-1J.400 Budapest, POBox 2.

Paprika /Capsicum annuum L./ is a traditional vegetable in Hungary, and numerous old varieties, and races etc. are collected in the gene bank of Research Center for Agrobotany /formerly the National Institute of Agrobotany/. Shmidt/1959/ and /1974/ reported the results of agrobotanical investigations made on some stocks and Szabo/1968/ investigated the cardinal points of germination temperature. Szabo and Vima Yi/1970/ tested seeds stored in unconditioned environment, and the seeds loss their germinability rapidly. However we have not data about connections of germination parameters registered in these tests.

In course of our trials 134 - 134 seed samples stored for 1 and 7 years were investigated respectively. The seeds were stored in conditioned environment at 0-4 °c in hermetically sealed containers' with 7-8 % initial moisture content. The germination tests were' carried out according to . /1985/ suggestions. Data of percentage of germination and of rates of seed molding and seedling abnormality were registered by samples, and correlation matrices of these parameters were computed. .

Significant differences were not stated for means of variables' registered in tests of seeds samples stored for 1 or 7 years. The mean value of germ inability was 92.13 ± 3.17 after 7 years. The correlations computed for the two seed populations /Table 1. 2. / show significant connections and similar tendencies at seed samples stored for different time periods. The computed correlations are signaling negative connections in pairs of germination-molding and germination-abnormality, and positive one in pair of molding-abnormality showing the possible tonic effect of the fungi colonizing the seeds. The connections were similar to those that were reported in case of lupins and pea earlier maY and HortvAth, 1990; 1991/.

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Table 1: Correlations of germination parameters after 1 year storage

Germinating	Molding	Abnormality
/G/ 1	-.6791+++	-.03062+++
/M/	1	.2598++
/A/		1

++; P=0.01,+++; P=0.001

Tables 2: Correlations of germination parameter after 7 year storage

Germinating	Molding	Abnormality
/G/ 1	-.6081++	-.2741++
/M/ 1	1	.2319
/A/		1

+ P= 0.05, ++ P=0.001

Screening of Germplasm For Yield and Quality Egg Plant in The Punjab.

Dr. Altaf hussain, M. Magir A. Chaudary, Mansoor Ahmad, Vegetable Research Institute, Faisalabad- Pakistan.

Egg plant is a very important vegetable crop in the Punjab province of Pakistan. Three crops are grown in this part of the globe during the year. Evaluation of available germplasm to screen out high yielding varieties was conducted Trial No. 1 March –July, 1989 and Trial 2 August 1989 – January, 1990

In the first trial 9 varieties were included in the studies. Nursery was sown in Feb under plastic tunnel, to protect it from the frost, whereas the transplanting was done in the 2nd week of March, RCBD system of layout with four replications was followed. Net Plot size was (8x3) m², row spacing 1.5m and within rows 0.5 m. was maintained.

In second trial, in addition to the varieties included in the first, one more variety Local Selection was Asses. In and within rows 0.5 m. was kept

Nursery was sown in the 2nd week of January and transplanting was done during the 2nd week of March, 1989 for second experiment the nursery was sown in last week of August, 1989, yield data recorded in both the cases are given in the table-1.

The data collected on yield of edible fruit in both the cases revealed that the variety “Neelum” produced included in the trials. Even all the varieties included in the trial. Even the standard variety Multan Selection could not compete it. Multan selection got the 2nd position on merit list in the first experiment, and fourth in the 2nd. The variety Black King produced significantly the lowest yield. In addition to the seasonal effect the plant population also affected the yield significantly. In first trial the density of the plant population was lesser whereas in the second experiment it was more i.e. row spacing 1.5m and 1. Respectively. The yield recorded was such higher in the 2nd experiment as compared with that of 1st experiment except in case of No. 7.

Table: Snowing The Fruit Field of Different Varieties During Spring and Autumn, 1989-90

S.No. Name of the variety

Yield in tomes per hectare

Trial No. 1 Trial No. 2

March-July, 89. Aug-Jan 1990

1. Nealum	12.59a	18.31a
2. Multan Gelection	11.53b	16.16b
3. Qaisar	11.4b	17.02ab
4. White Egg.	8.22c	12.06 cd
5. Sarhandi Long.	7.85c	11.40cd
6. Choraium Purple	7.76c	13.83c
7. Shah Jamal	7.72c	7.21e
8. Pusa Purple Long	7.66c	1322cd
9. Black King.	4.57d	6.83e
10. Local selection		17.68ab
Cd1	0.882	2.031
Cd2	1.196	2.742

EFFECT OF GELLING AGENTS AND ACTIVATED CHARCOAL ON SOLANUM MELONGENA PLANT REGENERATION

D. Perrone, V. Iannaccone and G.L. Rotino. Research Institute for Vegetable Crops, 20075 Contano L. (KI), Italy
Research Institute for vegetable Crops, P.O.BOX 48 - ~ Pontecagnano
(SA), Italy.

In *S.melongena* several previous studies showed that both genotypes and growth regulators influenced plant regeneration through organogenesis. The aim of this study was to evaluate the effect of agar and gerlite as gelling agents and the influence of activated charcoal on the morphogenetic response of two eggplant genotypes (Giant of China and WIR 769).

Murashige and Skoog medium supplemented with 2 mg/l glycine was used as basal medium. Cytokinins 6-benzylaminopurine (BAP) and zeatin (ZEA) were tested singly at 0.5, 1, 1.5, 5 and 10 mg/l or in combination using 0.5/1, 0.5/5, 1/1 and 1/5 mg/l of BAP/ZEA. All these media were solidified with 0.8% purified Agar-agar (MERK) or 0.25% gerlite (KELCO). Besides, agar-containing media were also tested with addition of 0.5% activated charcoal (MERK). The pH of all media was adjusted to 5.8 prior to autoclaving at 120°C for 20 min. All the cultures were kept in a 2 -1 controlled environment chamber at 25 + 2°C under 50 uEm s for a 16h day cycle.

For plant regeneration, leaf disks (1 cm) were prepared from young fully expanded leaves of *in vitro* grown-plants. Leaf-disks were placed with the lower side in contact with the medium in 100 x 15 mm petri dish. A total of 21 explants (7/plate) were used for each genotype and treatment. After 4 weeks the number of shoots and shoot buds per explant was collected under a dissecting microscope.

For both genotypes the gelling agent had a strong effect on the frequency of the shoot regeneration process. Agar was superior to gerlite in all the media tested (tab. 1 and 2). The positive effect of agar was more dramatic in BAP-containing media and for the cv. Giant of China where it allowed up to a twenty fold increment in the regeneration frequency (tab. 1). Surprisingly activated charcoal completely inhibited shoot regeneration in both genotypes. At the concentrations tested. ZEA was more effective than BAP to achieve shoots and shoot-buds differentiation. On average, WIR 769 showed an higher morphogenetic response than Giant of China. These results might be utilized in genetic transformation experiments to introduce, *Arsrobacterium tumefaciens*, useful foreign genes in the eggplant genetic pool.

Tab. 1 - Number of shoots and shoot-buds per leaf-disk (mean \pm SE) of cv Giant of China as influenced by growth regulators, gelling agents and activated charcoal.

Growth regulators (mg/l)		Gerlite	Agar	Agar + activated charcoal
ZEA	BAP			
0.5	-	6.9 \pm 4.4	19.0 \pm 2.6	0
1	-	2.3 \pm 2.7	15.5 \pm 4.0	0
1.5	-	2.0 \pm 2.6	18.0 \pm 2.4	0
-	1	6.3 \pm 5.7	15.2 \pm 1.1	0
-	5	0.8 \pm 1.1	14.4 \pm 1.3	0
-	10	0.6 \pm 0.9	13.6 \pm 2.9	0
0.5	1	-(a)	23.3 \pm 2.1	0
0.5	5	-	21.6 \pm 4.5	0
1	1	-	16.9 \pm 5.0	0
1	5	-	19.8 \pm 2.8	0

a, - = media not tested

Tab. 2 - Number of shoots and shoot-buds per leaf-disk (mean \pm SE) of WIR 769 genotype as influenced by growth regulators, gelling agents and activated charcoal.

Growth regulators (mg/l)		Gerlite	Agar	Agar + activated charcoal
ZEA	BAP			
0.5	-	19.0 \pm 9.9	30.3 \pm 4.5	0
1	-	5.2 \pm 4.1	25.0 \pm 5.8	0
1.5	-	21.2 \pm 4.9	24.3 \pm 7.0	0
-	1	6.6 \pm 4.0	12.3 \pm 6.7	0
-	5	8.2 \pm 6.1	23.1 \pm 9.9	0
-	10	1.6 \pm 1.2	7.4 \pm 5.1	0
0.5	1	-(a)	23.5 \pm 9.9	0
0.5	5	-	21.8 \pm 10.6	0
1	1	-	26.7 \pm 7.9	0
1	5	-	28.2 \pm 7.5	0

a, - = media not tested

INHERITANCE OF NPTII GENE AND SCREENING FOR KANAMYCIN RESISTANCE IN TRANSGENIC SOLANUM MELONGENA PLANT

F. Sunseri and G.L. Rotino.* Metapontum Agrobios - SS Jonica 106 kII 488.2, 75011 Metapontum (NT). Italy.
Research Institute for Vegetable Crops. 20075 Montanaso L. (MI). Italy

The aim of this study is to carry out genetic and molecular analysis of NPTII gene segregation in progenies of transgenic eggplant cv. 'Picientia' previously obtained (Rotino and Gleddie. 1990). Besides an easy and not-destructive screening method was also set up to select for functional NPTII gene as proposed by Weide et al., (1989) for tomato.

Seven RO plants derived from independent transformation events were used screening was carried out using three R1 progenies of T1, T14, T20 transgenic plants and untransformed control. Sterile seeds and plantlets were grown on selective MS basal medium containing 100-mg/l kanamycin. Plantlets exhibiting normal growth were considered to be kanamycin resistant. Two plants showed a 3 km :1 km ratio as expected in the case of one NPTII gene insertion, plant T14 gave a segregation ratio (2 kmR:1 kmS) that fit with the hypothesis of a lethal mutation, acting at the homozygous state caused by T-DNA insertion (Tab. 1). Resistant plants were randomly selected and selfed to obtain R2 seeds assay was performed on six R1 progenies two R2 progenies and untransformed control. To this end, greenhouse-grown plantlets at the fourth true leaf were sprayed twice a day (at 8 hours interval) for three consecutive days with a solution of 300 mg/l kanamycin dissolved in water. After 3 weeks plants were scored as resistant/sensitive to kanamycin on the basis of absence/presence of bleaching sectors in the sprayed leaves. The in vivo assay worked equally well as the in vitro screening. In fact, the R1 progenies of plant T1 and plant T20 gave the same segregation ratio in both in vitro and in vivo assay. All the R1 progenies segregated as one active copy of NPTII gene was inserted in the genome (Tab. 2). The T1-R2 progenies did not segregate because, probably, the selfed resistant T1-R1 plant was homozygous for NPTII gene; while the T20-R2 progenies showed a 3 kmR:1km ratio, this suggests that it derived from a heterozygous plant for NPTII gene.

All the plants gave segregation ratios that indicated inheritance of the * Km phenotype in a Mendelian fashion. The transgene was stably expressed through two sexual generations. Work is, now in progress to characterize the transgenic progenies. at molecular and enzymatic level. Rotino G.L. and Gleddie S., 1999 - Transformation of eggplant (*S. melongena* L.) using a binary agrobacterium tumefaciens vector. Plant Cell Rep., 9: 26-29.

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Tab. 1 - In vitro assay for kanamycin resistance segregation in R₁ progenies derived from S three independent transgenic plant (km^R=kanamycin resistant; km^S=kanamycin sensitive).

Transgenic plant	Total No. of seeds	Germination (%)	No. of seedlings tested	Phenotype		χ^2 value (P=0.05)	Segregation km ^R :km ^S
				km ^R	km ^S		
T1-R1	153	100	153	116	37	0.054	3:1
T14-R1	219	100	219	147	72	0.020	2:1
T20-R1	123	60	88	73	15	2.970	3:1
Control	140	97	136	0	136	-	-

Tab. 2 - In vivo assay for kanamycin resistance segregation in six R1 and two R₂ progenies derived from S six independent transgenic plant (km^R=kanamycin resistant; km^S=kanamycin sensitive).

Transgenic plant	Germination (%)	No. of seedlings tested	Phenotype		χ^2 value (P=0.05)	Segregation km ^R :km ^S
			km ^R	km ^S		
T1-R1	100	145	114	31	1.013	3:1
T1-R2	88	135	135	0	-	-
T10-R1	46	66	53	13	0.990	3:1
T12-R1	91	120	92	28	0.178	3:1
T16-R1	85	118	96	22	2.542	3:1
T20-R1	95	102	84	18	2.941	3:1
T20-R2	98	141	110	31	0.683	3:1
T22-R1	67	100	81	19	1.920	3:1
Control	90	135	0	135	-	-

ANNOUNCEMENT

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Kimio ITO, Vegetable Breeding Nagano, Chushin Agricultural, Shiojiri - NAGANO
399-64
Laboratory of Vegetable and Ornamental Horticulture, Faculty of Agriculture - Kyoto
University, KYOTO 606 Morioka Branch, V.O.C.R.S., MORIOKA 020-01
National Inst. of Agrobiological Resources, YATABE IBARAKI
National Research Inst. of Veget. Ornamental, Plants and Tea, Lab. of Breeding Soian.
Vegetables, ANO AGE-GUN MIE 514-23
Nihon Horticultural Production Institute, 207 Kamishiki, MATSUDO-SHI CHIBA-KEN
271
NIPPON DEL MONTE Corp., Research and Development, NUMATA, GUMMA 378
OHTA Yasuo, TSUKUBA - SHI 305
SAKATA SEED Corp., Kakegawa Breeding Station, Dr. K. Miyoshi, KAKEGAWA -
SHIZUOKA 436-01
Sakata Seed Corp., Plant Biotechnology Center, Toshio Shiga, SODEGAURA, CHIBA,
299-02
Shizuoka Agricultural Experimental Station, SHIZUOKA
The Nippon Shinyaku, Institute for Botanical Research, Oyake Sakanotsuji-cho 39,
KYOTO 607 YUKURA Yasuo 46.7 3-Chome, TOKYO

KOREA

Pilot Greenhouse Farm, PYONGYANG
Pyongyang Vegetable Research Center, PYONGYANG
KOREA,
Department of Agricultural Biology, College of Agriculture and Life Sciences, Seoul
National University, SUWON 441- 744
Department of Horticulture, College of Agriculture Seoul National University, SUWEON
170
Dept. of Horticulture, College of Agriculture, Kyungpook National University, TAEGU
702-701 Div. of Vegetable Breeding, Horticultural Experiment Station, SUWON 441-440
Horticultural Experiment Station, PUSAN 57111
NONG-WOO SEEDS, Plant Breeding Research Institute, 387-2 Sasa-2Ri, HWASONG
445-820 OSAN, Breeding Institute, Choong-Ang Seeds Co. Ltd., HWASUNG

LEBANON Plant Breeding Dept., Agricultural Research Inst., TRIPOLI ..
Faculty of Agricultural and Food Sciences, BEIRUT .
Institut de Recherche Agronomique du Liban (IRAL)

LIBERIA

CARI, Central Agricultural Research Institute, GBARNGA-BONG COUNTY CARL,
Central Agricultural Research Institute, MONROVIA, SUAKOKO

LIBYA

Agricultural Research Station, TRIPOLI nI:~ National Bureau for Agricultural Consultations and Studies, TRIPOLI

MALAYSIA

Dept. of Agronomy and Horticulture, University of Agriculture Malaysia, SERDANG - SELANGOR Dept. of Genetics & Cellular Biology, University of Malaya, KUALA LUMPUR 22-11 MARDI, KUALA LUMPUR MARDI, Research Station JALAN KEBUN, SERDANG-SELANGOR MARDI, Tanah Rata 39007, Cameron Highlands - PAHANG

MARTINIQUE , I.R.A.T.-C.I.R.A.D., FORT DE FRANCE

MAURITANIA

Centre Nat. de Recherche Agronomique et de Developpement de l'Agriculture, Ministere de l'Agriculture, Dept. de l'Horticulture, NOUAKCHON

MAURITIUS

Ministry of Agriculture and Natural Resources, Agricultural Service, PORT LOUIS

MEXICO

Centro de Botanica, Colegio de Postgraduados, 56230 CHAPINGO-Estado de Mexico

Centro de Investigaciones Agricolas del Nord, INIA-SARH, 3300 CD. DELICIAS -

CHIH. Experimental Station Celaya, INIFAP, CELAYA-GTO 38000

Genetic Center, College of Postgraduate, 56230 MONTECILLO

IBPGR, Oficina para Latinoamerica, c/o CIMMYT, MEXICO 06600 D.F.

Instituto Nacional de Investigaciones Forestales y Agropecuarias (INIFAP), Dr. Juan Hernandez Hernandez, 93400 PAPANTLA-VERACRUZ

Instituto Nacional de Investigaciones Forestales, Agricolas y Pecuarias (INIFAP),

MEXICO CITY Instituto Nacional de. Investigaciones Agricolas, Apartado Postal C-1,

TAMPICO Jose A. Laborde, CELAYA-GTO. 38040

Library, C.I.F.A.P., Campo Exper. del Sur de Tamaulipas Apartado Postal C-1,

Institut National de la Recherche Agronomique, INRA, RABAT Societe du

Developpement Agricole (SODEA), RABAT

MOZAMBIQUE

Facultad de Agricultura, Universidade Eduardo Mondlane, MAPUTO

MYANMAR

Institute of Agriculture, University of Yangon, University Estate, YANGON
NEW ZELAND

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The Librarian (serials), Massey University, PALMERSTON NORTH
NICARAGUA

Istituto Superior Ciencias Agropecuarias, REGEN, MANAGUA
NIGERIA

Dept. of Agronomy, Inst. for Agricultural Research, Ahmadu Bello University,
SAMARU - ZARIA

Dept of Crop Science, University of Nigeria, NSUKKA

National Horticultural Research Inst, Idi-Ishin Jericho Reservation Area, IBADAN

NORWAY

Dept of Vegetable Crops, the Agricultural University of Norway, 1432 AA8-NLH IF;
PAKISTAN

Pakistan Agricultural Research Council, ISLAMABAD

Vegetable Research Institute, FAISALABAD 38950

PERU

Dept. de Ciencias Agropecuarias, Universidad Nacional de San Agustin, AREQUIPA

Dept de Horticultura, Universidad Nacional Agraria, LA MOLINA-LIMA

Experiment Station LA MOLINA, LA MOLINA -LIMA HOLLE Miguel, LIMA 18

Instituto Nacional de Investigacion, Promocion Agropecuaria (INIPA), Sinchi Roca 2727
- Lince, LIMA 14

Julio A. QUEA, T ACNA

PHILIPPINES

College of Agriculture, Inst of Plant Breeding, Univ. of the Philippines at Los Banos,
LAGUNA 3720 EAST-WEST SEED Company Inc., MAKATI (Manila)

POLAND

Academy of Agriculture, Inst of Genetics and Plant Breeding, 60-625 POZNAN

Department of Plant Genetics, Breeding and Biotechnology, 02766 WARSAWA

Dept of Genetics and Plant Breeding, University of Agriculture, 60-198 POZNAN

Dept of Genetics, A. Mickiewicz University, 60-594 POZNAN

Inst of Plant Genetics, Polish Academy of Sciences, 60-479 POZNAN

POIAN Krakowska Hodowta, I Nasiennictwo Ogronicze, 30-130 KRAKOW Research

Institute of Vegetable Crops, 96-100 SKIERNIEWICE

PORTUGAL

I.N.IA, Estacao Agronomica Nacional, OEIRAS

ROMANIA

Research Inst for Vegetable and Flower Growth, 8268 VIORA JUD. GIURGIU

RWANDA

Institut des Sciences Agronomiques du Rwanda (ISAR), BUTARE

SOUTH AFRICA

Division of Plant and Seed Control, PRETORIA 0001

SPAIN

C.S.I.C., Estacion Experimental La Mayora, ALGARROBO-COSTA MALAGA Centro de Investigaciones Agrarias, 26080 LOGRONO Clause Iberica S.A., PATERNA (Valencia)

Departamento de Bioquimica y Biologia Molecular, Universidad de Almeria, 04120 LA CANADA DE SAN URBANO Dept. de Biologia Animal y Vegetal, F. Merino de Caceres, Universidad de La Coruna, 15701 LA CORUÑA Diputacion General de Aragon, Servicio de Investigacion Agraria, Seccion Documentacion y Bibliotheca, 50080 ZARAGOZA

Escuela de Capacitación Agraria, DON BENITO (Badajoz) GIL ORTEGA R., D.G.A. - S.I.A., 50080 ZARAGOZA Horticulture Department, C.R.I.A., MURCIA

I.N.I.A., Dr.F.Ponz, 28080 MADRID

Instituto Nacional Investigaciones Agrarias., Cit. Centro Inv. y Tecn. Biblioteca, 28040 MADRID :-, Plant Protection Department, C.R.I.A., MURCIA

Polytechnical University of Valencia, Biotechnology Department, Dr. F. Nuez, 46020 VALENCIA Polytechnical University of Valencia, Plant Protection Department Pathology, 46020 VALENCIA Semi lias Fit6, BELLPUIG (Lerida)

Semillas Pioneer S.A., J. Riado Abad, 04710 EL EJIDO (ALMERIA) SEMILLAS RAMIRO ARNEADO, CALAHORRA-LOGRONO Sluis & Groot Semi lias, EL EJIDO-ALMERIA

SRI LANKA

Agricultural Research Station, MAHAILLUPPALLAMA

Central Agricultural Research Institute, GANNORUVA, PERADENIYA

Food Technology Section, Ceylon Inst. of Scientific and Industrial Research, COLOMBO Government's Department of Agriculture, PERADENIYA

ST. LUCIA

C.A.R.D.I., CASTRIES

SUDAN

Agricultural Research Corporation, Horticulture Germplasm Unit, WAD MEDANI Department of Horticulture, Faculty of Agriculture, University of Khartoum, SHAMBAT, KHARTOUM University of Gesira, Faculty of Agricultural Sciences, Dept. of Horticulture, WAD MEDANI

SUISSE

Nestec S.A, CH-1800 VEVEY "

THAILAND

APTA, Phaya Thai Court, BANGKOK 10400

AVRDC, Thailand Outreach Program, Kasetsart University, (Kasetsart) BANGKOK 10903 CHIA TAI Company Limited, BANGKOK 10100

Div. of Horticulture, Dept. of Agriculture, Bagkhen - BANGKOK

EAST-WEST Seed Co. Ltd., Mr. S. J. de Joop, CHIANG MAI 50290 Faculty of Agriculture, Chiang Mai University, CHIANG MAI 50002

Horticulture Research Institute, Headquarters, Dept. of Agriculture, Ministry of Agriculture and Cooperatives,

BANGKOK 10900 r Thep Watana Seed Co. Ltd., BANGKOK 10500

THE NETHERLANDS

BEL AGRO Handelmaatschappij b.v., 1001 AS AMSTERDAM

Bruinsma Seed b.v., Jurko Leij, 2670 AA NAALDWIJK

Chronica Horticulturae, CH-ISHS, 6703 BC WAGENINGEN

De Ruiter Seed, 2665 BLEISWIJK

DE RUITER ZONEN, Dr. D. Vreugdenhil, 2691 RA 'S GRAVENZANDE

Enza laden, De Enkhuizer Zaadkandel b.v., R.Kuijsten, 1600 AA ENKHUIZEN 78

EUCARPIA Secretariat, 6700 AH WAGENINGEN

Gebr. Bakker Zaadteelt Zaadhandel, 1723 LM NOORDSCHARWOUDE

Glasshouse Crops Research and Experiment Station, 2670 AA NAALDWIJK

Institute of Horticultural Plant Breeding, 6700 AA WAGENINGEN

Keygene NV, 8700 AE WAGENINGEN "

Kniphorst International Booksellers, Postbus 67

Landbouwwuniversiteit 99458, Bibliotheek, 6700 HA WAGENINGEN Leen de Mos

Groentezaden BV, 2690 AB 's GRAVENZANDE Nickerson Zwaan b.v., 2990 AA

BANRENDRECHT Nunhems Zaden BV, 6080 AA HAELEN

Rijk Zwaan Zaadteelt en Zaadhandel B.V., 2678 ZG DE LIER Royal Sluis, 1600 AA ENKHUIZEN

S & G Seeds, J. van Deursen, 2678 LV DE LIER

SAKATA SEED EUROPE B. V., Dr. Y. Kobayashi, 1435 DD RIJSENHOUT Scientia Horticulturae, AMSTERDAM

Sluis en Groot Seeds B. V., 2678 LW DE LIER

Swets & Zeitlinger B.V., Backsets Department, 2160 SZ LISSE .. Van der Zaden B. V., Beethovenstraat 42, 5102 XB DONGEN

TOGO

Department de l'Agriculture, Ministere de l'Agriculture, YAOUNDE

TUNISIA

Department of Biological Sciences, Faculty of Sciences, Library Research II, 1060

TUNIS ' Ecole Superieure d'Horticulture, CHOTT -MARIEM-SOUSSE

INRAT, Inst. National Recherche Agronomique de Tunisie, 2080 ARIANA INRAT, Laboratoire de Cryptogamie, 2080 ARIANA

Inst. National Agronomique de Tunisie, Lab. Cultures Maraicheres et Florales, 1002-

TUNIS BELVEDERE Station d'Appui de la Medjerda, 2010 MANOUBA

TURKEY

Aegean Regional, Agricultural Research Inst., MENEMEN-IZMIR

Ankara University, Faculty of Agriculture, Department of Horticulture, ANKARA-

Diskapi Atatork Horticultural Research, Yalova Inst., ISTAMBUL

Department of Horticulture, Fac. Agriculture, Univ. Of Cukurova, 01330 ADANA

Department of Plant Pathology, Fac. Agriculture, Univ. Of Cukurova, 01330 ADANA

Ege Universitesi, Ziraat FakOltesi Bitki Koruma B610mO, BORNOVA 35100-IZMIR

Uludag Univ., Faculty of Agric., Dept. of Horticulture, BURSA Vegetable Research

Institute, 07110 ANTALYA

U.S.A.

ASGROW E.P.G., KALAMAZOO - Michigan 49001

ASGROW SEED Company, Dr.R.Heisey, SAN JUAN BAUTISTA- CA 95045 Chili-Queen, TUCSON - ARIZONA 85704

College of Agricultural Sciences, University of Delaware, Department of Plant Sciences, NEWARK - Delaware 19717. Cornell University, Albert R. Mann, Library Acquisition Division, ITHACA - New York 14853 DE MARS Lawrence, MINNEAPOLIS-Minn 55410

Dept. of Agr. Engineering, Michigan State University, EAST LANSING -Michigan 48824-1323 ., Dept. of Biology, Indiana University, BLOOMINGTON - IN 47405 ...

Dept. of Botany, Miami University, OXFORD - Ohio 45056

Dept. of Horticultural Sciences, New York Agricultural Expt. Stat., Prof. R.W.Robinson, GENEVA-New York ~ Dept. of Horticulture, Louisiana Agricultural Exp. Stat., BATON ROUGE - LA 70803-2120 Dept. of Horticulture, Michigan State University, EAST LANSING - Michigan 48824

Dept. of Vegetable Crops, Cornell University, Plant Science Building, ITHACA - N. Y. 14853-0327 Dept. of Vegetable Crops, University of California, DAVIS - California 95616

Dept. Plant Breeding & Biometry, Cornell University, ITHACA - N.Y. 14853-1902

Dept. Plant Pathol. And Crop Physiol., Louisiana State University, BATON ROUGE - Louisiana 70803 Desert Botanical Garden, Richter Library, PHOENIX - ARIZONA 85008 Dr. Bradley Boese, SAGINAW - MICHIGAN 48601 Dr. Jean ANDREWS, AUSTIN - TX 78703

Extension-Research Center, ATTAPULGUS-Georgia 31715

Genetics Department, North Carolina State University, RALEIGH-NC 27695

Hortinova Research Inc., Joseph Stem, SAN JUAN BAUTISTA -California 95045 FAS, University of Florida, Agric. Research and Education Center, FORT PIERCE - Florida 33454 IFAS, University of Florida, Agronomy Department, GAINESVILLE - Florida 32611-0621 IFAS, University of Florida, ARES BELLE GLADE - Florida 33430 IFAS, University of Florida, DELRAY BEACH-Florida 33444

tFAS, University of Florida, Prof. Robert E. Stall, Plant Pathology Department, GAINESVILLE - Florida 32611-0513 Library, CORNELL University, New York State Agricultural, GENEVA-New York 14456

National Seed Storage Laboratory, U.S. Dept. Agriculture, Colorado State University, FORT COLLINS-Colorado 80523

New Mexico State University, Dept. of Agronomy and Horticulture, Dr. P. W. Bosland, LAS CRUCES-New Mexico 88003 New York Botanical Garden, Library, Serials & Exchange, BRONX-N.Y. 10458 - 5126 Pepper Research Inc., Dr. S.R.Subramanya, BELLE GLADE-FL 33430

PETOSEED Co, Dr. Kenneth Owens, WOODLAND - CA 95695

Petoseed Florida Research, Dr. Dale S. Kammerlohr, FELDA - FLORIDA 33930-0249

Rogers Food Co., Chili Products, Dr. P.A. Gniffke, 39502 Cypress Ave., GREENFIELD-CA 93927 ..

ROGERS SEED Co., Steven J. Czaplewski, BOISE -IDAHO 83711-4188 -"- Suburban Experiment Station, University of Massachussets, WAL THAM-Ma 02254

Texas Agr. Exp. Station, The Texas University, WESLACO - Texas 78596-8399.

The Pepper Gal National Hot Pepper Association, Dr. Elisabeth A. Payton, FORT LAUDERDALE - FL 33311 The Pillsbury Company, Technology Center, Att. Bonnie Moore, MINNEAPOLIS - MN 55414-2198

USDA, Agr. Mark. Serv., Plant Variety Protection Office, NAL Bldg.-Room 500, BELTSVILLE - Maryland 20705-2351

USDA, National Agricultural Library, Current Serial Records Rm 002, BELTSVILLE - Maryland 20705 USDA-ARS Southern Regional, Plant Introduction Station, EXPERIMENT-Georgia 30212 USDA-ARS-SM, Gilbert R.Lovell, GRIFFIN-GA 30223-1797

UGANDA

Library, Kawanda Agricultural Research Station, KAMPALA

YUGOSLAVIA

Vegetable Research Institute, Palanka, 11420 SMEDEREVSKA PALANKA VITAMIN, Spice Pepper Breeding Station, 24420 HORGOS

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