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2003 POTATO BREEDING AND GENETICS RESEARCH REPORT

David S. Douches, J. Coombs, K. Zarka, S. Cooper, L. Frank, J. Driscoll, and E. Estelle

Department of Crop and Soil Sciences Michigan State University

East Lansing, MI 48824

Cooperators: R.W. Chase, Ray Hammerschmidt, Ed Grafius Willie Kirk, George Bird, Sieg Snapp and Chris Long

INTRODUCTION

At MSU, we conduct a multi-disciplinary program for potato breeding and variety development that integrates traditional and biotechnological approaches. We conduct variety trials of advanced selections and field experiments at MSU research locations (Montcalm Research Farm, Lake City Experiment Station, Muck Soils Research Farm and MSU Soils Farm), we ship seed to other states and Canadian provinces for variety trials, and we cooperate with Chris Long on 17 grower trials throughout Michigan. Through conventional crosses in the greenhouse, we develop new genetic combinations in the breeding program, and also screen and identify exotic germplasm that will enhance the varietal breeding efforts. With each cycle of crossing and selection we are seeing directed improvement towards improved varieties (e.g. combining chip-processing, scab resistance and late blight resistance). In addition, our program has been utilizing genetic engineering as a tool to introduce new genes to improve varieties and advanced germplasm for traits such as solids, insect resistance and disease resistance. We feel that these in-house capacities (both conventional and biotechnological) put us in a unique position to respond to and focus on the most promising directions for variety development and effectively integrate the breeding of improved chip-processing and tablestock potatoes.

The breeding goals at MSU are based upon current and future needs of the Michigan potato industry. Traits of importance include yield potential, disease resistance (scab, late blight and early die), insect (Colorado potato beetle) resistance, chipping (out-of-the-field, storage, and extended cold storage) and cooking quality, bruise resistance, storability, along with shape, internal quality and appearance. We are also developing potato tuber moth resistant lines as a component of our international research project. If these goals can be met, we will be able to reduce the grower's reliance on chemical inputs such as insecticides, fungicides and sprout inhibitors, and improve overall agronomic performance with new potato varieties.

PROCEDURE

I. Varietal Development

Each year, during the winter months, 500-1000 crosses are made using about 150 of the most promising cultivars and advanced breeding lines. The parents are chosen on the

basis of yield potential, tuber shape and appearance, chip quality, specific gravity, disease resistance, adaptation, lack of internal and external defects, etc. These seeds are then used as the breeding base for the program. We also obtain seedling tubers or crosses from other breeding programs in the US. The seedlings are grown annually for visual evaluation (size, shape, set, internal defects) at the Montcalm and Lake City Research Farms as part of the first year selection process of this germplasm each fall. Each selection is then evaluated post harvest for specific gravity and chip processing. These selections each represent a potential variety. This system of generating new seedlings is the initial step in an 8-12 year process to develop new varieties. This step is followed by evaluation and selection at the 8-hill, 20-hill and 30-hill stages. The best selections out of the four-year process are then advanced for testing in replicated trials (Preliminary, Adaptation, Dates-of-Harvest, Grower-cooperator trials, North Central Regional Trials, Snack Food Association Trials, and other out-of-state trials) over time and locations. The agronomic evaluation of the advanced breeding lines in the replicated trials is reported in the annual Potato Variety Evaluation Report.

II. Evaluation of Advanced Selections for Extended Storage

With the Demonstration Storage facility adjacent to the Montcalm Research Farm we are positioned to evaluate advanced selections from the breeding program for chipprocessing over the whole extended storage season (October-June). Tuber samples of our elite chip-processing selections are placed in the demonstration storage facility in October and are sampled monthly to determine their ability to chip-process from colder (42-48°F) and/or 50°F storage.

III. Germplasm Enhancement

To supplement the genetic base of the varietal breeding program, we have a "diploid" (2x = 24 chromosomes) breeding program in an effort to simplify the genetic system in potato (which normally has 4x chromosomes) and exploit more efficient selection of desirable traits. This added approach to breeding represents a large source of valuable germplasm, which can broaden the genetic base of the cultivated potato. The diploid breeding program germplasm base at MSU is a synthesis of seven species: *S. tuberosum* (adaptation, tuber appearance), *S. raphanifolium* (cold chipping), *S. phureja* (cold-chipping, specific gravity, PVY resistance, self-compatability), *S. tarijense* and *S. berthaultii* (tuber appearance, insect resistance) and *S. chacoense* (specific gravity, low sugars, dormancy and leptine-based insect resistance). In general, diploid breeding utilizes haploids (half the chromosomes) from potato varieties, and diploid wild and cultivated tuber-bearing relatives of the potato. Even though these potatoes to transfer the desirable genes by conventional crossing methods via 2n pollen.

IV. Integration of Genetic Engineering with Potato Breeding

Through transgenic approaches we have the opportunity to introduce new genes into our cultivated germplasm that otherwise would not be exploited. It has been used in potato as a tool to improve commercially acceptable cultivars for specific traits. Our laboratory has 11 years experience in *Agrobacterium*-mediated transformation to introduce genes into

important potato cultivars and advanced breeding lines. We are presently using genes in vector constructs that confer resistance to Colorado potato beetle (Bt-cry3A and avidin), potato tuber moth, late blight resistance via the RB gene, lowering glycoalkaloids (STG), and drought resistance (CBF1). We also have the glgC16 gene (ADP-glucose pyrophosphorylase (AGPase or starch gene) from Monsanto to modify starch and sugar levels in potato tubers. Furthermore, we are investing our efforts in developing new vector constructs that use alternative selectable markers and give us the freedom to operate from an intellectual property rights perspective. In addition, we are exploring transformation techniques that eliminate the need for a selectable marker (antibiotic resistance) from the production of transgenic plants.

RESULTS AND DISCUSSION

I. Varietal Development

Breeding

The MSU potato breeding and genetics program is actively producing new germplasm and advanced seedlings that are improved for cold chipping, and resistance to scab, late blight, and Colorado potato beetle. For the 2003 field season, progeny from over 600 crosses were planted and evaluated. Of those, the majority were crosses to select for round whites (chip-processing and tablestock), with the remainder to select for yellow flesh, long/russet types, red-skin, and novelty market classes. In addition to crosses from the MSU breeding program, crosses were planted and evaluated from collaborative germplasm exchange from other breeding programs including North Dakota State University, University of Minnesota, and the USDA/ARS program at the University of Wisconsin as part of the Quad state cooperative effort. During the 2003 harvest, about 1800 selections were made from the 45,000 seedlings grown. Following harvest, specific gravity was measured and potential chip-processing selections were chipped out of the field. All potential chip-processing selections will be tested in January or March 2003 directly out of 42°F and 50°F storage. Atlantic (50°F chipper) and Snowden (45°F chipper) are chipped as check cultivars. Selections have been identified at each stage of the selection process that have desirable agronomic characteristics and chip-processing potential. At the 8-hill and 20hill evaluation state, 400 and 150 selections were made, respectively. Table 1 lists some of the potential lines for grower trials in year 2004.

Chip-Processing

Excellent chip-processing selections have been identified in the breeding program, despite switching to a more stringent screening temperature (42 vs. 45°F storage) a few years ago. Over 70% of the single hill selections have a chip-processing parent in their pedigree. Of those selections, about 75% have a SFA chip score of 1.5 or less. Based upon the pedigrees of the parents we have identified for breeding cold-chipping potato varieties, we have a diverse genetic base. We believe that we have at least eight cultivated sources of cold-chipping. Examination of pedigrees shows up to three different cold-chipping germplasm sources have been combined in these selections.

Our promising chip-processing lines are MSF099-3 (42°F chipper), MSG227-2 (scab resistant 45°F chipper), MSH095-4, MSH094-8, MSH067-3, MSH228-6, MSJ147-1, MSJ126-9Y, MSJ167-1, and late blight resistant chipper MSJ461-1.

Dr. Joe Sowokinos, Univ. of Minnesota, has conducted biochemical analyses of our best chipping lines and has discovered that our lines differ from older varieties in their proteins involved in chipping. His analysis will allow us target specific crosses to find improved chip-processing varieties that will allow processing from colder storage temperatures.

Tablestock

Efforts have been made to identify lines with good appearance, low internal defects, good cooking quality, high marketable yield and resistance to scab and late blight. Our current tablestock development goals now are to continue to improve the frequency of scab resistant lines, incorporate resistance to late blight along with marketable maturity and excellent tuber quality, and select more russet lines. From our breeding efforts we have identified mostly round white lines, but we also have a number of yellow-fleshed and redskinned lines, as well as long, russet type and purple skin selections that carry many of the characteristics mentioned above. We are also selecting for a dual-purpose russet, round white, red-skin, and improved Yukon Gold-type yellow-fleshed potatoes. Some of the tablestock lines were tested in on-farm trials in 2003, while others were tested under replicated conditions at the Montcalm Research Farm. Promising tablestock lines include MSE221-1 as a scab resistant tablestock, while MSE018-1 is a high yielding tablestock with a large oval shape. Boulder (MSF373-8) is a high yielding line with large tubers that also chip out of the field. Michigan Purple also performs well. In addition, all these clones performed well in the dry land trial at Montcalm Research Farm. MSE192-8RUS and MSE202-3RUS are two russet table selections that have excellent type and scab resistance. MSI005-20Y and MSJ033-10Y are yellow-fleshed lines with smooth round appearance and high yield potential.

Disease and Insect Resistance Breeding

Disease screening for scab has been an on-going process since 1988. Results from the 2003 MSU scab nursery indicate that 42 of 190 lines evaluated demonstrated little to no infection to common scab. In addition, 23 other MSU breeding lines showed moderate scab resistance. The limitation of breeding for scab resistance is the reliance on the scab nursery. The environmental conditions can influence the infection each year, thus multiple year data provides more reliable data. A laboratory-based screening process is currently under development that would use thaxtomin in tissue culture to expedite selection of material with potential scab resistance. Secondly, the scab nursery space has been full. In response, we have spent 3 years developing a second scab nursery. In 2004, we will begin early generation evaluation of scab reaction in the breeding program. This additional effort should lead to more clones with scab resistance. Since the mid-1990's we have directed efforts to identify sources of late blight resistance and use this resistance to breed late blight resistant varieties. At MSU, we have also participated in the national late blight trial and we have conducted our own efforts to use field and greenhouse screening to identify additional sources of resistance that can be used by the breeding community. In the past 7 years the MSU breeding program has intensely evaluated over 1200 crosses between late blight resistant x late blight susceptible parents and have identified parents that transmit strong late blight resistance to the highest percentage of the offspring. This year we added an early generation screen which will improve our ability to select late blight resistant lines.

As of 2003, based upon 7 years of inoculated field experiments, we have at least 8 sources of foliar resistance to the US8 genotype of Phytophthora infestans (Mont.) that have different pedigrees from which their resistance is derived. The resistance in Jacqueline Lee has now held resistance for 7 years of testing. MSJ461-1, the chipprocessing selection, has the same late blight resistance source Jacqueline Lee and was resistant to a US-17 genotype of *Phytophthora infestans* in New York this year. Our other promising late blight resistant lines that have been tested in replicated agronomic trials are MSJ317-1, MSI152-A, MSJ453-4, MSK136-2, MSL159-AY, MSL179-AY and MSL211-3 (see Potato Variety Evaluation Report for agronomic data). In each of these lines, the resistance is based on a single resistance source. If we rely on a single source of resistance, the varieties developed from this strategy may be overcome by *P. infestans* at some future date that we cannot predict. Therefore, the most effective breeding strategy is to combine resistance from different pedigrees to build a more durable resistance. Our efforts are now focusing on pyramiding the different resistance sources. This year we added a transgenic strategy using the late blight resistant *RB* gene cloned from *S. bulbocastanum*. We should have initial field tests using transplants in 2004.

With support from GREEEN, we also introduced an early generation Colorado potato beetle screen at the Montcalm Research Farm. From this screen we identified 32 individuals with either transgenic or non-transgenic foliar resistance to Colorado potato beetle. Eighteen lines were categorized as moderately resistant and 36 were susceptible.

Single-hill selections in 2003 also had an exciting number of individuals with pedigrees for potential late blight, Colorado potato beetle or scab resistance or some combination of the three. Of the single hill selections, 75% of progeny have at least one late blight parent, 15% have a Colorado potato beetle resistant parent, and 25% have a scab resistant parent in its pedigree.

II. Evaluation of Advanced Selections for Extended Storage: MSU Potato Breeding Chip-processing Results From the MPIC Demonstration Commercial Storage (October 2001 - June 2002)

The MSU Potato Breeding Program has been conducting chip-processing evaluations each year on potato lines from the MSU breeding program and from other states. For 4 years we have been conducting a storage study to evaluate advanced breeding lines with chip-processing potential in the Dr. B. F. (Burt) Cargill Potato Demonstration Storage facility directly adjacent to the MSU Montcalm Research Farm. In October 2002, tuber samples from 6 lines in the Montcalm Research Farm trials were placed in the bin to be cooled to 48°F. Tubers from another 9 lines were placed in the bin that was to be cooled then held at 51°F. The first samples were chip-processed at MSU in October and then, each month until June 2003. Samples were evaluated for chip-processing color and quality.

Table 2 summarizes the chip-processing color of select lines over the 8-month storage season. In the 48°F bin, Snowden was the check variety. In April the Snowden and MSH095-4 chips began to go off-color. In contrast only MSG227-2, MSH094-8 and MSF099-3 and W1201 maintained acceptable chip color throughout the storage sampling. Of these lines, MSG227-2, MSF099-3 and MSH094-8 maintained the lightest chip color throughout the storage season. MSG227-2 also has scab resistance.

In the 51°F bin Atlantic and Pike were used as check varieties and both varieties chip-processed acceptably until May. Of the 7 advanced breeding lines evaluated Liberator chip-processed acceptably throughout the storage season until June. Liberator offers chip-processing from storage and scab resistance. MSJ461-1 had the most consistent chip color throughout the storage season until May. MSJ461-1 also offers strong foliar late blight resistance along with the chip-processing quality; however the solids content can be lower than other chip-processing lines. UEC also had good chip color until the May sampling.

In addition, Liberator and MSF099-3 was grown by Sandyland Farms and placed in one of the 500 cwt bins. Despite field frost occurring in MSF099-3's harvested tubers, the potatoes chip-processed successfully out of the bin in April 2003 at Utz. The Liberator bin was sent to Shearers in March and produced a good chip product.

III. Germplasm Enhancement

In 2003, about 5% of the populations evaluated as single hills were diploid. From this breeding cycle, we plan to screen the selections chip-processing from storage. In addition, selections were made from over 2,000 progeny that was obtained from the USDA/ARS at the University of Wisconsin. These families represent material from South American potato species and other countries around the world that are potential sources of resistance to Colorado potato beetle, late blight, potato early die, and ability to cold-chip process. About 100 selections were made among the diploid material in 2003. Through GREEEN funding, we were able to initiate a breeding effort to introgress leptine-based insect resistance. From previous research we determined that the leptine-based resistance is effective against Colorado potato beetle. We will continue conducting extensive field screening for resistance to Colorado potato beetle at the Montcalm Research Farm and at the Michigan State University Horticulture Farm in 2004.

Late Blight Breeding and Genetics: Mapping Late Blight Resistance in three Populations

A high priority objective of the breeding program is to identify sources of late blight resistance and use these sources for breeding varieties with late blight resistance. In 1999 we initiated a set of studies (via GREEEN) to identify the genes in potato associated with late blight resistance. If we can identify the genes that contribute to late

blight resistance we feel that we could more effectively breed varieties with durable late blight resistance. A diploid potato population was developed with the objectives to map quantitative trait loci (QTL) conferring resistance to Phytophthora infestans (Mont.) de Bary and other agronomic traits using simple sequence repeats (SSR) and isozymes and to examine associations between late blight resistance and other agronomic traits. The mapping population was a cross between a late blight resistant selection of Solanum *microdontum* Bitter and a susceptible diploid advanced breeding clone. A second diploid population derives its late blight resistance from S. berthaultii. The third population is tetraploid and the resistance comes from Jacqueline Lee. Based upon field trials at the Muck Soils Research Farm, Bath, MI between 1999 and 2002, we have identified major late blight resistance genes in the three populations. Currently, one chromosome region containing the resistance is linked to a genetic marker has been identified in S. *microdontum.* This past year we identified a major QTL associated with late blight resistance was found in the tetraploid population and multiple QTLs for late blight resistance in the S. berthaultii mapping population. These QTLs should be suitable for marker-assisted selection to introgress a new source of resistance to P. infestans to the cultivated tetraploid germplasm of potato.

The tetraploid cross for mapping (Jacqueline Lee x MSG227-2) offers more than just mapping late blight resistance genes. This cross has traits such as late blight resistance, scab resistance, chip-processing, specific gravity, maturity all segregating at one time. This summer we screened a sub sample of the population for scab reaction. A number of the progeny showed little scab. In 2004 we hope to screen a greater number of the population.

IV. Integration of Genetic Engineering with Potato Breeding

Assessment of Natural (Glandular Trichomes and Glycoalkaloid-Based) and Engineered (*Bt-cry3A*) Potato Host Plant Resistance Mechanisms for Control of Colorado potato beetle: Caged no-choice studies.

The Colorado potato beetle, Leptinotarsa decemlineata Say (Coleoptera: Chrysomelidae), is the leading insect pest of potato (Solanum tuberosum L.) in northern latitudes. Host plant resistance is an important tool in an integrated pest management program for controlling insect pests. A field study was conducted in 2003 to compare natural (glandular trichomes (NYL235-4) and glycoalkaloid-based (ND5822C-7)), engineered (Bt-cry3A: NO8.8, Atlantic NewLeaf, Bt-cry1Ia1: Spunta G2) host plant resistance mechanisms of potato for control of Colorado potato beetle. Six different potato lines representing 5 different host plant resistance lines were evaluated in caged studies (no-choice) at the MSU campus farms. Each cage with 10 plants represented one plot. The cages were arranged in a randomized complete block design consisting of three replications. Twenty egg masses were placed on the plants in each cage. Observations were recorded weekly for a visual estimation of percent defoliation by Colorado potato beetles, and the number of egg masses, larvae, and adults. The Bt-cry3A transgenic line and the combined resistance line were effective in controlling feeding by Colorado potato beetle adults and larvae. The high glycoalkaloid line had less feeding, but the beetles clipped the petioles, which led to greater defoliation in the first few weeks. Foliage regrowth occurred by the end of the season. The glandular trichome line suffered less

feeding than the susceptible control. Spunta G2 was effective in limiting defoliation, but larval mortality was not as high as in the *Bt-cry3a* lines. Based on these results, the *Bt-cry3A* gene in combination with glandular trichome mechanism is an effective strategy that could be used to develop potato varieties for use in a resistance management program for control of Colorado potato beetle. **Figure 1** shows the results of caged trial in 2003.

Bt-cry3A-transgenic line Agronomic Trial

In 2001 and 2002, we had extensive field testing for agronomic performance in replicated trials of our most advanced *Bt-cry3A* transgenic lines. Based upon 2001 agronomic performance and 2002 Bt-cry3A protein concentrations in foliage, 12 of 26 transgenic lines were eliminated. In general, the *Bt-cry3A* transgenic lines had similar agronomic and tuber characteristics compared to the non-transgenic parental line. These selections represent a diverse portfolio of *Bt-cry3A* lines that could be commercialized if the intellectual property rights and regulatory requirements could be met. We will maintain these lines in our program. These lines are MSE018-1, NYL235-4, NY123, Jacqueline Lee, Onaway, Norwis and Spunta. If the acceptance of transgenic food crops becomes deregulated, we will consider these lines for commercialization. In 2003 we developed a new Bt-cry3A construct that uses a different gene promoter. We are currently transforming MSJ461-1 and Michigan Purple.

USAID-funded International project to Develop Potato Tuber Moth Resistant Potatoes

Potato tuber moth, *Phthorimaea operculella* (Zeller), is the most serious insect pest of potatoes worldwide. The introduction of the Bacillus thuringiensis (Bt) toxin gene via genetic engineering offers host plant resistance for the management of potato tuber moth. The primary insect pest in Egyptian potato production, like many other countries in the Middle East, is the potato tuber moth. In the field, the moths lay their eggs on the potato foliage and the hatched larvae mine the foliage and the stems. This feeding damage leads to irregular transparent tunnels in the leaves and weakening of the stem. The larvae attack the tubers through infected stems or directly from eggs, which are oviposited on exposed tubers or where soil cracks allow moths to reach the tubers. Larvae mine the tuber in the field and in storage reducing potato quality and increasing the potential for pathogen infection. Field and storage studies were conducted to evaluate Btcry5 potato lines for resistance to potato tuber moth in Egypt under natural infestations and their agronomic performance in both Egypt and Michigan. From 1997-2001, field experiments were conducted at the International Potato Center (CIP) Research Station, Kafr El-Zyat, Egypt and/or Agricultural Genetic Engineering Institute (AGERI), Giza, Egypt to evaluate resistance to tuber moth.

Two transgenic 'Spunta' clones, G2 and G3, have been identified that produced high control levels of mortality in first instars of potato tuber moth in laboratory tuber tests (100% mortality), and field trials in Egypt (99-100% undamaged tubers). Reduced feeding by Colorado potato beetle first instars was also observed in detached-leaf bioassays (80-90% reduction). Field trials in the U.S. demonstrated that the agronomic performance of the two transgenic lines was comparable to 'Spunta'. We are currently working with USAID, Syngenta and South Africa to commercialize the Spunta-G2 and Spunta-G3 lines. We have also transformed Atlantic, Lady Rosetta and Jacqueline Lee with the *Bt-cry5* gene. We hope to have approval to field test these in Mexico some time in the future.

V. Variety Release

The MSU breeding program has now named and released its first varieties and is in the process of licensing the new varieties to the Michigan Potato Industry Commission. Three potato varieties were released in 2001: Jacqueline Lee (MSG274-3), Liberator (MSA091-1), and Michigan Purple. MSU is currently licensing the first 3 varieties to MPIC and working out procedures to market these varieties. Boulder (MSF373-8) was released in 2003. Virus-free tissue culture plantlets are maintained at MSU.

Development of a DNA-based Fingerprint System for Potato Varieties

The ability to quickly and accurately identify potato clones is important to potato breeding programs and to the potato seed industry and commercial growers. Since 1990, the Michigan State University Potato Breeding and Genetics Program has used an isozyme-based fingerprint system to identify potato cultivars. Isozyme analysis has been an economical and effective means of discriminating potato clones; however, they require fresh, healthy tuber or leaf tissue. DNA-based fingerprinting using simple sequence repeats (SSRs or microsatellites) has been shown to discriminate between potato clones. The objective of this study was to identify those SSR primer combinations that accurately and efficiently distinguish clones on polyacrylamide and agarose gels. SSR primer combinations used were based on polymorphism levels in previous tetraploid studies from PCR amplification products. DNA isolated from 17 potato clones representing chip processing, tablestock, russet and red market classes were visualized on both polyacrylamide and low melting point (Metaphor ®) agarose gel systems. Eighteen SSR primer combinations were screened on both gel systems. Polymorphism was observed in all eighteen of the primer combinations on polyacrylamide (PAGE) and fourteen on agarose gel systems. The PAGE system was determined to be the preferred system for variety identification, but agarose can be used to differentiate lines when specific varietal comparisons need to be made. The primer combination STM0031 with STACCAS3 was able to differentiate all 17 clones on agarose. In addition, five different DNA source tissue types were evaluated (fresh foliar, freeze-dried foliar, fresh tuber skin, freeze-dried tuber skin, and freeze-dried tuber tissue). Amplification products were similar for all five tissue sources used for DNA isolation. This ability to isolate DNA from freeze-dried tissue will allow us to fingerprint varieties when fresh tissue is not available. The SSR fingerprinting system presented here can be used as a practical fingerprint system for cultivated potato. This research will be published in the American Journal of Potato Research.

	Pe	digree				
Line	Female	Male	Comments			
Processing						
BOULDER (MSF373-8)	MS702-80	NY88	Chips out of the field, large tubers			
MSF099-3	Snowden	Chaleur	42 °F chipper			
MSG227-2	Prestile	MSC127-3	Scab resistant			
MSH067-3	MSC127-3	W877	Flat, round			
MSH094-8	MSE251-1	W877	45 °F chipper			
MSH095-4	MSE266-2	OP	45 °F chipper			
MSH112-6	Michigold	Zarevo	42 °F chipper, high solids			
MSH112-0 MSH228-6	MSC127-3	OP	Scab tolerant			
MSJ036-A	A7961-1	Zarevo	Scab tolerant chipper			
MSJ080-1	MSC148-A	S440	High yield			
MSJ147-1	Norvalley	S440	cold chipper			
MSJ461-1	Tollocan	NY88	Late blight resistant			
MSK061-4	MSC148-A	ND2676-10	Scab tolerant chipper			
WISK001-4	MSC140-A	ND2070-10	Seab tolerant empper			
Tablestock						
BOULDER (MSF373-8)	MS702-80	NY88	Chips out of the field, large tubers			
MICHIGAN PURPLE	W870	Maris Piper	Bright purple skin, white flesh			
MSE192-8RUS	A8163-8	Russet Norkotah	Scab resistant russet (Norkotah replacement)			
MSE202-3RUS	Frontier Russet	A8469-5	Scab resistant russet			
MSH031-5	MSB110-3	MSC108-2	Bright skin			
MSI005-20Y	MSA097-1Y	Penta	Yukon appearance			
MSI152-A	Mainestay	B0718-3	Late blight resistant, round white			
MSJ033-10Y	MSA097-1	Penta	Yellow, Scab resistant			
MSJ317-1	B0718-3	Prestile	Late blight resistant, round white			
		1 01114				

Table 1. Potential Lines for 2004 On-Farm Grower Trials

Table 2. 2002-2003 DEMONSTRATION STORAGE CHIP RESULTS

Chip Scores represented using SFA Scale[†]

	2002				Sample Dates:							
	DO)H*	2002	2002	11/12/02	12/18/02	01/07/03	02/11/03	03/10/03	04/10/03	05/05/03	06/03/03
	CWT/A DOH* S		$\text{SCAB}^{\dagger\dagger}$	Bin Temperature (°F)								
POTATO LINE	US#1	TOTAL	SP GR	RATING	57 °F	50 °F	48 °F	48 °F	48 °F	48 °F	- °F	- °F
MSF099-3	323	348	1.076	3.7	1.0	1.5	1.0	1.0	1.5	1.5	-	-
MSG227-2 ^{SCABRES}	256	326	1.072	0.5	1.5	1.0	1.5	1.5	1.0	1.5	-	-
MSH094-8	299	324	1.075	2.3	1.0	1.5	1.0	1.5	1.0	1.5	-	-
MSH095-4	326	351	1.076	2.0	1.5	1.5	1.5	1.5	1.0	2.0	-	-
SNOWDEN	262	304	1.073	2.0	1.5	1.5	1.0	1.5	1.5	2.0	-	-
W1201	364	391	1.081	1.3	1.5	1.5	1.5	1.5	1.5	1.5	-	-
					57 °F	55 °F	51 °F	55 °F	55 °F	54 °F	54 °F	56 °F
ATLANTIC	328	352	1.078	2.7	1.0	1.5	1.5	1.5	1.5	2.0	1.0	2.5
UEC	390	407	1.072	1.5	1.0	1.0	1.5	1.5	1.5	1.5	1.0	2.5
LIBERATOR SCABRES	276	309	1.074	0.0	1.5	1.5	1.5	1.5	1.0	1.5	1.5	1.5
MSE018-1	438	470	1.079	3.6	2.0	2.0	1.5	1.5	1.5	2.0	1.0	2.0
MSF373-8	392	401	1.072	2.5	2.0	1.5	1.0	1.5	2.0	1.5	1.5	2.0
MSI002-3	376	420	1.077	4.0	1.5	2.0	1.5	1.5	1.5	2.5	1.5	2.5
MSI083-5	255	288	1.072	3.3	1.5	1.5	1.5	1.5	2.0	1.5	1.5	2.5
MSJ461-1 ^{LBR}	279	330	1.069	2.7	1.5	1.5	1.5	1.5	1.5	1.5	1.5	2.5
PIKE	262	302	1.077	1.1	1.5	1.5	1.5	1.0	1.0	1.5	2.0	2.0

LSD_{0.05} 53

50 0.002

[†]CHIP SCORE: Snack Food Association Scale (Out of the field); Ratings: 1-5; 1: Excellent, 5: Poor.

^{††}SCAB DISEASE RATING: MSU Scab Nursery; 0: No Infection; 1: Low Infection <5%; 3: Intermediate; 5: Highly Susceptible.

*Agronomic data from Date of Harvest, Round-White Late Harvest (DOH) Trial; Montcalm Research Farm, September 21, 2001.

Chip scores were from two-slice samples from five tubers of each line collected at each sample date.

SCABRES Resistant to Common Scab Streptomyces scabies

LBR Resistant to foliar Late Blight, *Phytopthora infestans*

Table 3. Potato Seed Inventory 2003

MSU Potato Breeding Program Introductions Availability of Michigan Certified Seed A Cumulative Inventory

	MINI-				
	TUBERS	FY1	FY2	FY3	FY4
LINE	(UNITS)	(CWT)	(CWT)	(CWT)	(CWT)
JACQUELINE LEE (MSG274-3)	-	-	12	-	-
LIBERATOR (MSA091-1)	750	60	484	-	-
MICHIGAN PURPLE	15,500	123	103	244	-
MSE192-8RUS	1,500	-	25	300	-
MSE202-3RUS	-	-	-	-	-
MSF099-3	-	-	12	-	144
MSG227-2	-	-	750	-	-
MSH031-5	-	95	-	-	-
MSH067-3	-	50	-	-	-
MSH095-4	-	20	-	-	-
MSI152-A	875	-	-	-	-
MSJ461-1	400	53	32	-	-

Information listed above is a cumulative count from Golden Seed Farms, Hanson Farms, Iott Seed Farms Inc., Krueger Seed Farm, Marker Farms, Makarewicz Seed Farm, and Sklarczyk Seed Farm.

Table courtesy of Chris Long.

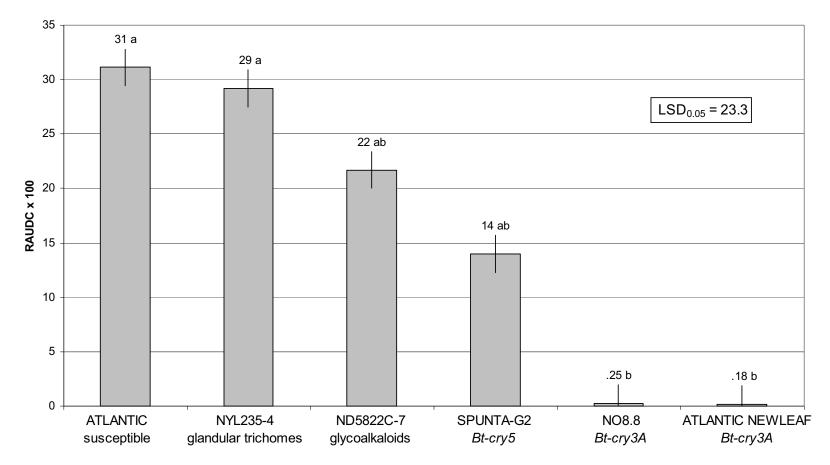


Fig. 1. Colorado Potato Beetle Field Cage No-Choice Trial, Relative Area Under the Defoliation Curve (RAUDC) Results of Host Plant Resistance Potato Lines (2003)

Host Plant Resistant Potato Clone