



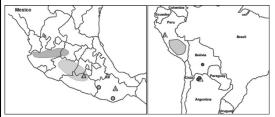
## Phaseolus vulgaris L. Common Bean

- One of 5 species of *Phaseolus* domesticated
- Found from Mexico to Argentina
- An annual legume in subfamily Fabaceae
- A species variable for growth habit, seed color and ecological adaptation
- An important food crop, a source of protein
- Nitrogen fixer similar to other grain legumes
- Includes horticultural garden, green, snap beans

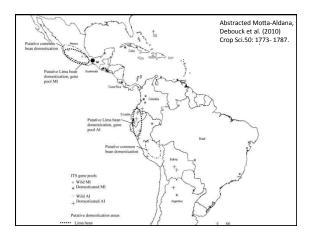


Where was common-be	ean domesticated?
- M	
	- Parking
3 m	
© Cristina Menéridez and Paul Gepts - PLB143 - L05	April 17,1996 11

# New Origin of Common Bean



Common Bean Origin & Domestication - green area – Jalisco, Mexico Common Bean Domestication in Andes – violet area – Southern Peru Maize Domestication - orange area – Michocan, Mexico, Guerrero Abstracted: Bitocchi, Papa et al. 2012



## Two Gene Pools and Six Races

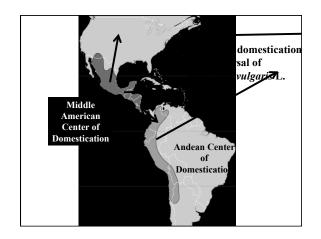
### Andean Gene Pool Larger Seeded Types

- Race Nueva Granada--White & Red Kidneys
- Race Peru--Yellow
- · Race Chile--Cranberry

### Middle American Gene Pool Small & Medium Seeded Types

- Race Durango--Pinto and Great Northern
- Mesoamerica--Navy and Black
- · Race Jalisco--Reds and Pinks









# Dry Beans in the US Today

- The US is the 6th leading producer of dry beans
- Production in Northern & Higher Altitudes Zones
- Michigan and North Dakota together produce half of the total US production rainfed conditions
- Western production irrigated
- 20% of production is exported
- Nationwide value of \$790 million in 2011

Source USDA/ERS accessed April 2012

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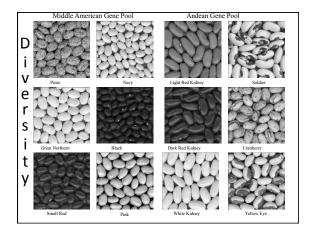
## Dry Beans in Michigan

- Production value of \$134 million in 2011
- Michigan is the leading producer of:
- · --Black beans
- --Cranberry
- --Small Reds
- Michigan is second leading producer of:
- --Navy
- --Kidney
- Non GMO Crop

Source USDA/ERS accessed April 2012

# Need for Diversification

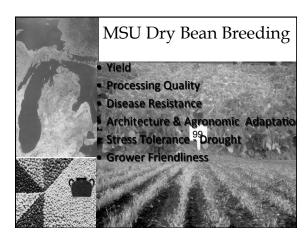




## **Common Bean Genetics**

- Diploid, Self Pollinating
- Annual Species, 2n=2x=22
- 11 chromosomes Pv01-Pv11
- No natural outcrossing
- Near Homozygous Pure Line Varieties
- Bean Genome Sequenced
- Small Genome ~ 587 Mb
- Genetic Synteny with Soybean





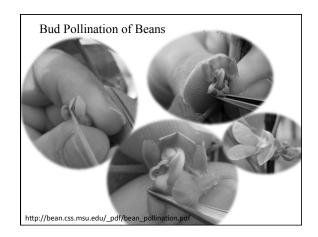
# Breeding Systems - Traits

- Twelve different commercial seed classes
- · Pedigree breeding Yield & Quality
- Backcross -Disease Resistance
- Recurrent Selection -Architecture
- Inbred BC Introduce Diversity
- Marker-Assisted Selection: MAS – Disease Resistance
- QTL Analysis for Quantitative Traits
- Genome Wide Association Mapping

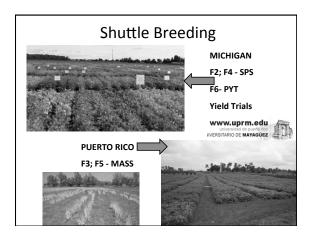


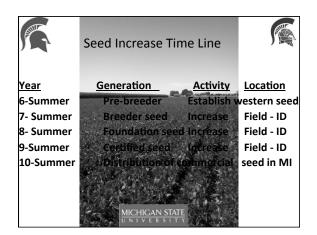
# Pield Improvement Growth Habit Maturity Disease Resistance Market class Seed Traits Canning Quality Broad Foundation Adequate Genetic Variability Adapted from Kelly et al. (1998) Breeding for yield in dry bean. Euphylica 102:343-356

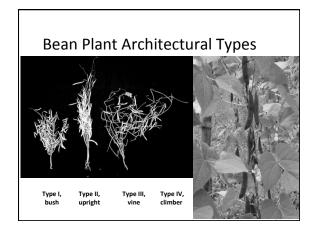


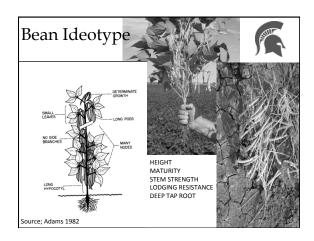


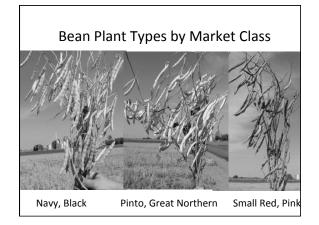
	Breeding Time	Line			
<u>Year</u>	Generation	<u>Activity</u>	<b>Location</b>		
1- Fall	Parents	Crossing	GH		
1- Spring	F1 plants	Selfing	GH		
1- Summer	F2 plants	SPS	Field - MI		
2-Winter	F3 plant rows	MS	Field - PR		
2- Summer	F4 plant rows	SPS	Field - MI		
3- Winter	F5 plant rows	MS	Field - PR		
3- Summer	F6 yield trial	PYT	Field - MI		
4- Summer	F7 yield trials	AYT	Field - MI		
5- Summer	F8 yield trials	RYT	Field MI+		
6- Winter	Pre -Release of B	Pre -Release of Breeding line as new Variety			
SPS – Single Plant Selection, MS – Mass Selection: PYT – Prelimin Yield Trial, Advanced & Regional					

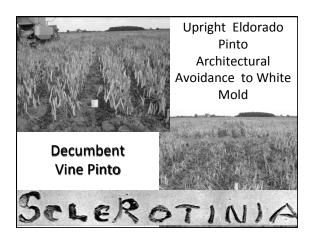






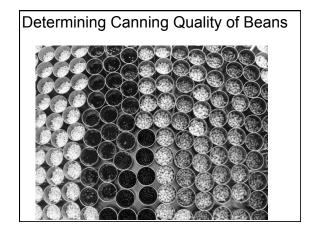








# "Straight cutting" --Clipping --Clipping



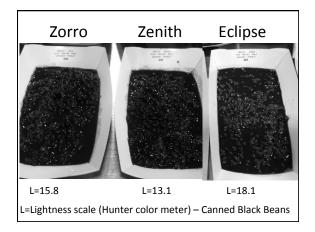
# Selection for Canning Quality Traits

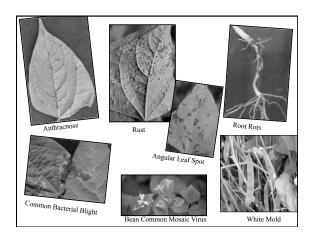
- Visual selection trained judges
- Whole seed integrity
- Color retention, uniformity
- Clear brine no starch extrusion
- Measure color of canned product
- Hydration ratios, drained weights
- Texture Kramer Shear Press
- Commercial evaluation elite lines

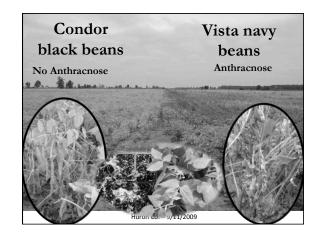


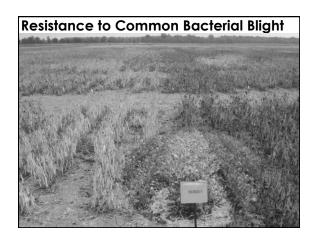
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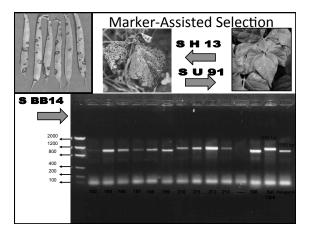




# Can we Improve Phenotypic Selection with Marker Assisted Selection – MAS?

# Resistance Genes in Common Bean with Tightly Linked Markers

Gene	Disease	Gene	Disease
Co-1	Anthracnose	Mp-1 N	<b>Aacrophomi</b> i
Co-2	"	Mp-2	"
Co-4	"	Ur-3	Rust
Co-42	"	<i>Ur-4</i>	"
Co-5	"	Ur-5	"
Co-6	"	Ur-7	"
Co-9	"	Ur-9	"
I	BCMV	Ur-11	"
bc-12	"	bgm-1	BGMV
bc-3	"	Phg-1	ALS
		Phg-2	ALS



## Marker - Aided Bean Breeding

Using molecular markers to select for resistance to:

- Anthracnose
   Colletotrichum lindemuthianum
- Bean Common Mosaic Virus
- Rust

Uromyces appendiculatus

• Common Bacterial Blight Xanthomonas axonopodis



# Bean Germplasm Releases (52)



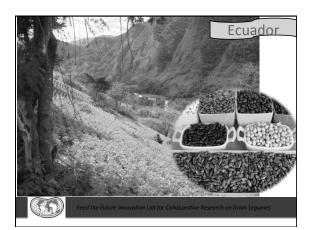
Illumina Infinium Genechip BARCBEAN6K\_3 SNP 2,687 InDel Markers NDSU\_IND\_3\_10.8903

www.beancap.org



United States Department of Agriculture National Institute of Food and Agriculture

This project is funded by the USDA National Institute of Food and Agriculture





## Beans are Nutrient Dense Food Great to Eat!!

Bean Frijol

Frejol Feijao

Poroto Judia

Haba









## Additional Information - Bean Breeding

Bean Improvement Cooperative <a href="http://bic.css.msu.edu/">http://bic.css.msu.edu/</a>

BeanCAP Plant Architecture

http://www.youtube.com/watch?v=wf\_nOs7DP-o

Plant Breeding Science + Creative Problem Solving <a href="http://www.youtube.com/watch?v=wiMl-">http://www.youtube.com/watch?v=wiMl-</a>

uGcsIk&feature=youtu.be

Bean Genome, Schmutz et al. Nature Genetics (2014) doi:10.1038/ng.3008





## Thanks for joining us today.

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http://www.extension.org/pages/60426/webinar-registration-and-archive

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