

Cameron Peace, Nahla Bassil, Michael Coe, Chad Finn, Ksenija Gasic, Stan Hokanson, Jim Luby, Dorrie Main, Jim McFerson, Jay Norelli, Mercy Olmstead, Vance Whitaker, Chengyan Yue, and Amy Iezzoni

# **Outline of Presentation**



United States Department of Agriculture

National Institute of Food and Agriculture

Amazing Germplasm × Traditional Genetics

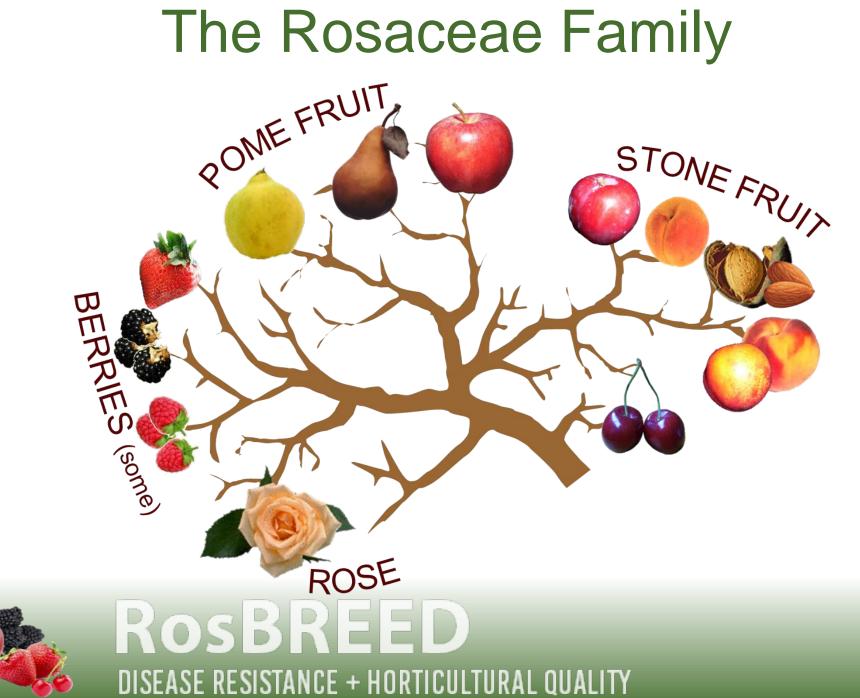
2 International Collaboration × DNA-Based Technologies

Cultivar Development Pipeline × Genomics Research Advances











# Amazing Germplasm!

# Features of Rosaceous Crops

- Product quality is paramount
- Clonally propagated
  - Perennial

Rosbreed

DISEASE RESISTANCE + HORTICULTURAL QUAI

- Often long juvenility
- Often obligate outcrossers
- Often high heterozygosity
- Sometimes polyploid

Few generations in cultivation, much diversity in primary genepools
 → huge genetic gains possible



#### **Traditional Genetics**

- Crossing & selecting mostly by phenotype
- Quantitative genetics rarely used

 Breeders often rely on serendipity

#### Challenges: large plant sizes, long juvenility, disease resistances combined with superior quality

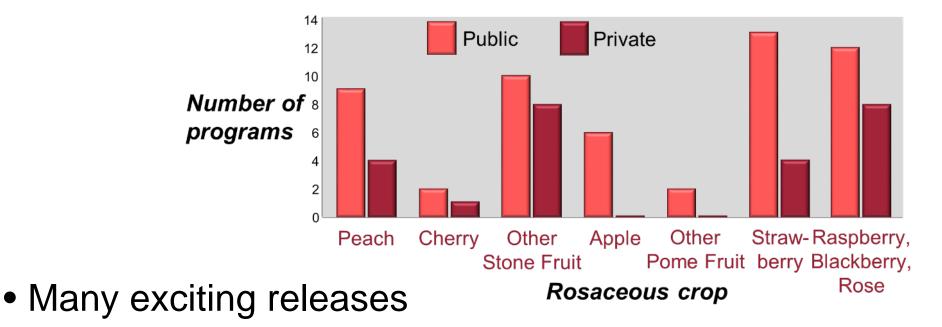


Cosmic Crisp™ brand WA 38, an eye-catching, large, dark red apple

# Oultivar Development Pipelines Oultivar Development Oultivar Development

• Need for much public breeding (some crops/regions not commercially viable?)





Cultivar development pipelines are functional but inefficient

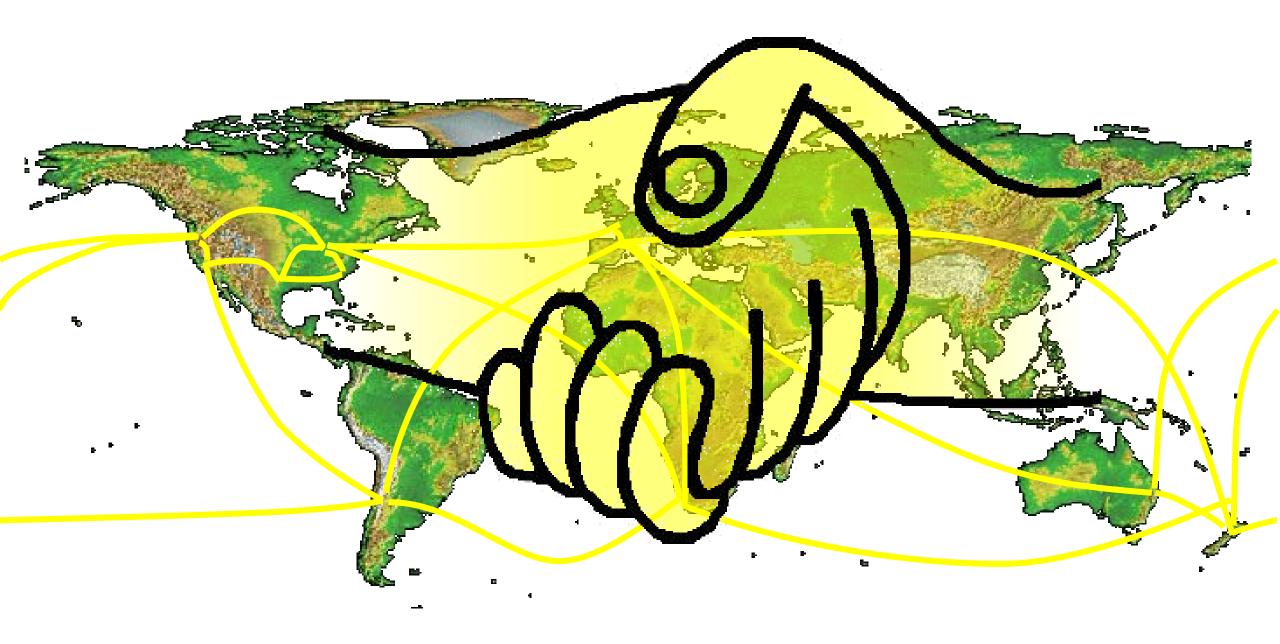
DISEASE RESISTANCE + HORTICULTURAL QUALITY

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# Strong Spirit of Collaboration



#### International Research Community Hub



#### Welcome to the Genome Database for Rosaceae

Initiated in 2003, the Genome Database for Rosaceae (GDR) is a curated and integrated web-based relational database providing centralized access to Rosaceae genomics, genetics and breeding data and analysis tools to facilitate basic, translational and applied Rosaceae research. GDR is supported by grants from the NSF Plant Genome Program (2003-2008), USDA NIFA Specialty Crop Research Program (2009-2019), USDA NIFA National Research Support Project 10 (2014-2019), and the Washington Tree Fruit Research Commission (2008-2016), Clemson University, University of Florida and Washington State University.

#### What's new in GDR?

- Gene Naming Guideline is available (4/23/15)
- F. vesca genome V1.1.a2 and V2.0.a1 and Malus x domestica genome V3.0.a1 available (4/6/15)
- Peach genome V2.0.a1 available (3/18/15)
- New QTL, markers and genetic map data from apple and rose are available (3/15/15)
- New search interfaces for haplotype data available (2/9/15)
- GDR related presentations are available for download (1/15/15)

#### **General Information**

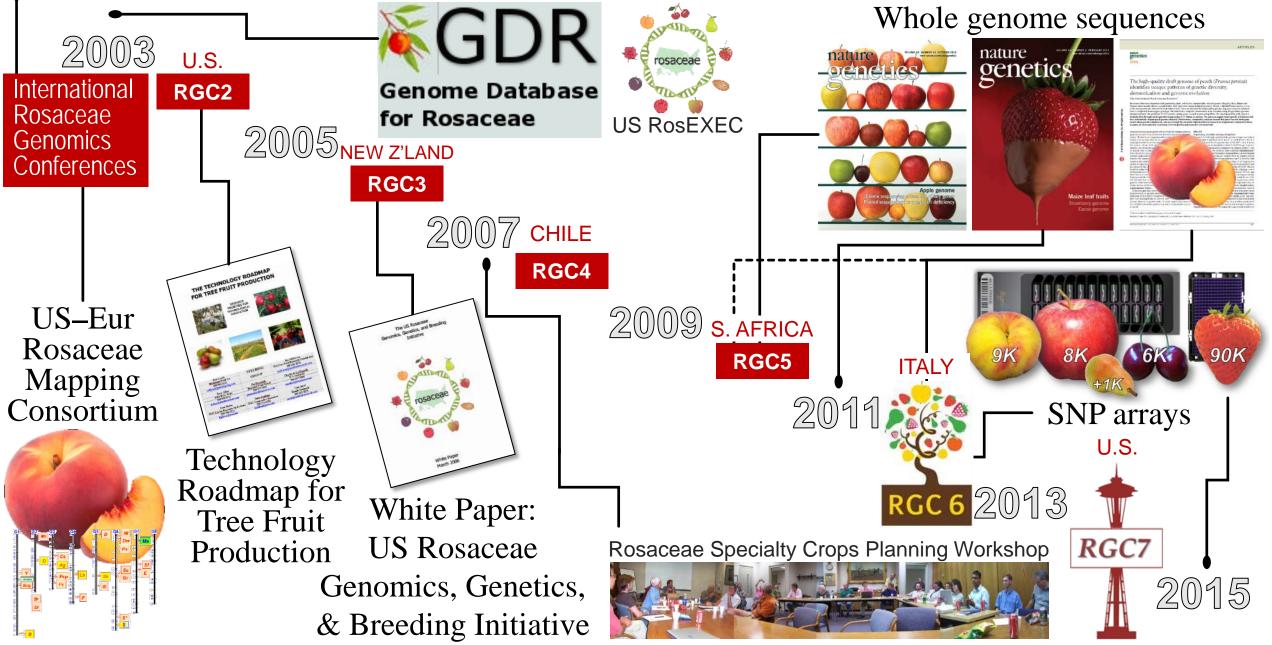
Report a problem | Ask us a question | Post a job | Post a meeting or event | Used GDR data or tools in your research - how to reference us.

#### News

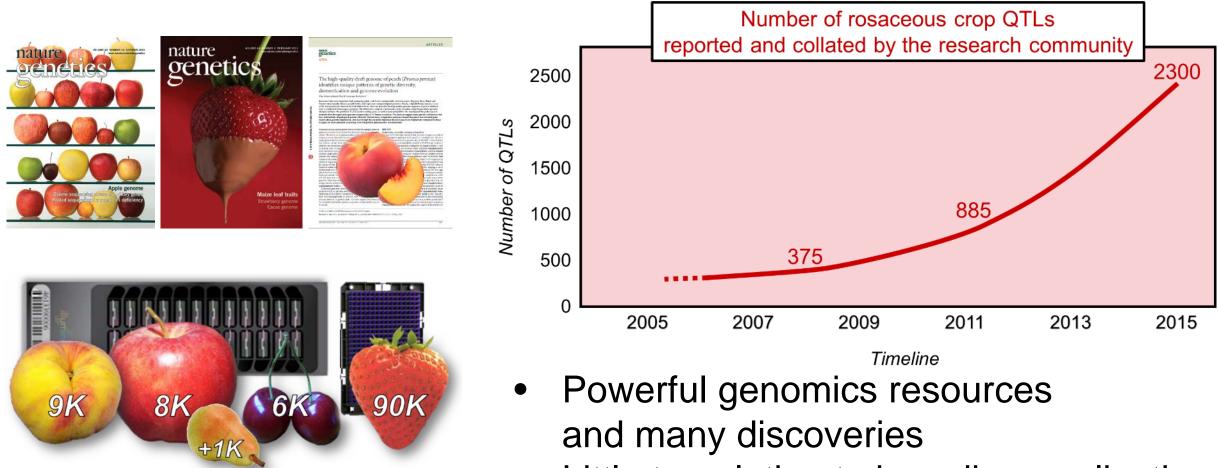
- Gene Naming Guideline available (4/23/15)
- Rose transcriptome data and 68K Axiom SNP array data (Koning-Boucoiran et al. 2015) are available for download (4/23/15)
- GDR Newsletter April 2015 available (4/6/15)
- F. vesca genome V1.1.a2 and V2.0.a1 and Malus x domestica genome V3.0.a1 available (4/6/15)
- Prunus persica genome V2.0.a1 available (3/18/15)
- New QTL, markers and genetic map data for apple and rose (3/15/15)
- New search interfaces for haplotype data (2/9/15)
- GDR related presentations are available for download (1/15/15)
- · Announcements archive

Supported by grants from the NSF Plant Genome Program (2003-2008), USDA NIFA Specialty Crop Research Program (2009-2019), USDA NIFA National Research Support Project 10 (2014-2019), and the Washington Tree Fruit Research Commission (2008-2016). Copyright © 2002-2015. This site is designed to work with IE8, Mozilla, Safari and Opera.

### **Rosaceae Community Collaboration**

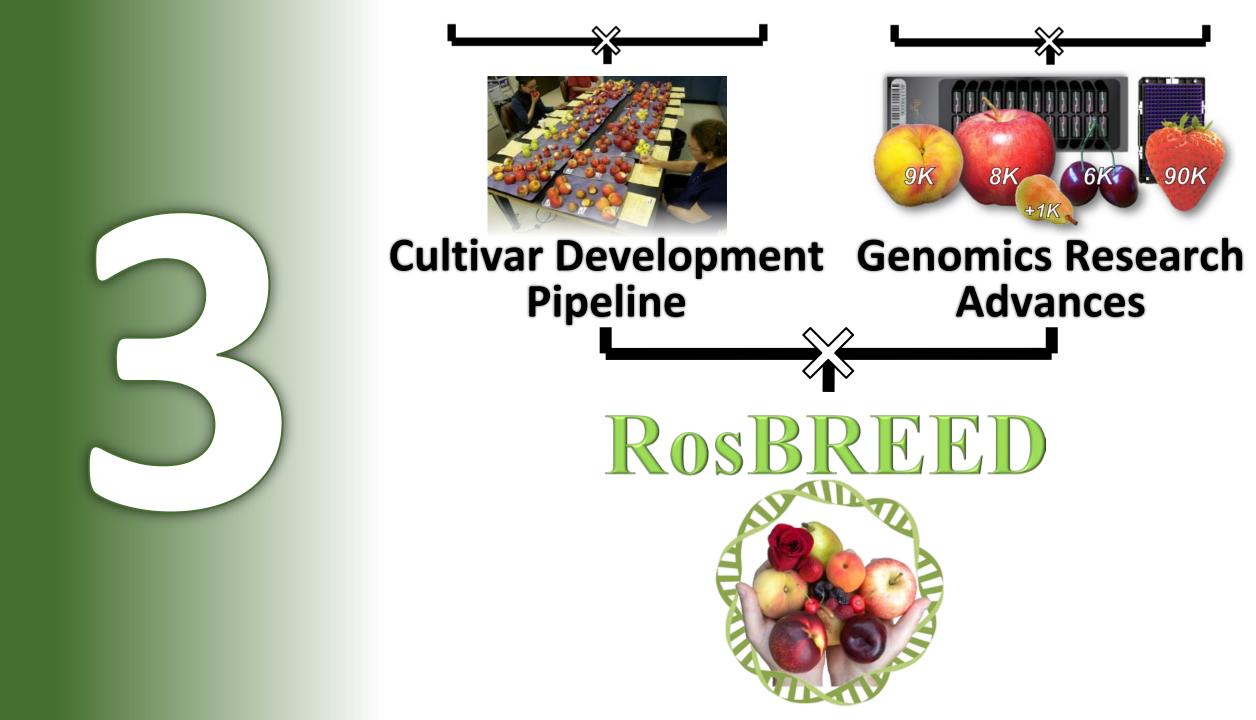


# → Genomics Research Advances



Little translation to breeding application





#### Genomic Shashaeth Breeding

#### **Genomics** resources (e.g., whole genome sequences)

#### Genomics **Research** information

DNA

**Decision** support

Breeding

rograms

#### More efficient development of new cultivars



# **Genomics Research**

DNA

information

# Some Reasons for the Chasm

#### Reported QTLs themselves:

- Trait low priority / low value
- Weak linkage
- Different germplasm
- Unknown functional alleles
- Unknown genetic action
- Unknown linkage relationships
- Unknown environment effects
- Unknown management effects

No suitable DNA testing services Not believed to be cost efficient No training in routine DNA testing

**Decision** support Breeding rograms



<u>MSU</u> Amy lezzoni (PD) Jim Hancock Dechun Wang Cholani Weebadde

Univ. of Arkansas John Clark <u>WSU</u> Cameron Peace (co-PD) Dorrie Main Kate Evans <u>Univ.</u> Karina Gallardo J Raymond Jussaume Che Vicki McCracken Nnadozie Oraguzie <u>Orego</u> Mykel Taylor Alexa

<u>Univ. of Minnesota</u> Jim Luby Chengyan Yue

Oregon State Univ. Alexandra Stone

Enabling marker-assisted breeding in Rosaceae Enabling MSU lezzoni (PD)

> <u>Clemson</u> Ksenija Gasic Gregory Reighard

> > <u>Texas A&M</u> Dave Byrne

**USDA-ARS** 

Nahla Bassil

Gennaro Fazio

Chad Finn

Plant Research Intl,

Netherlands

Eric van de Weg

Marco Bink

Univ. of CA-Davis Tom Gradziel Carlos Crisosto

Univ. of New Hamp. Tom Davis

#### "RosBREED 1"

- Funded by USDA Specialty Crop Research Initiative
- \$7.2 M federal
- Sep 2009 Aug 2014
- Project Directors:
   Amy lezzoni
   Cameron Peace

### **RosBREED 1 Outcomes**

- Socio-economic values for levels of breeding traits
- Genomics resources and DNA information: SNP array genome scans, QTLs

DISEASE RESISTANCE + HORTICULTURAL QUALITY

Rosbreed



### RosBREED 1 Outcomes

 Practical tools for breeding use: DNA tests for many breeding-relevant traits

DISEASE RESISTANCE + HORTICULTURAL QUALITY

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• DNA information about breeding germplasm: genetic potential of parents, seedlings, selections, cultivars



#### DNA Tests in 2009









#### **DNA Tests Now**



#### RosBREED 1 Outcomes

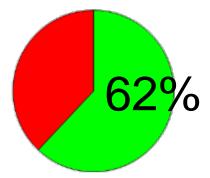
• Trained the next generation of breeders!



### **RosBREED 1 Outcomes**

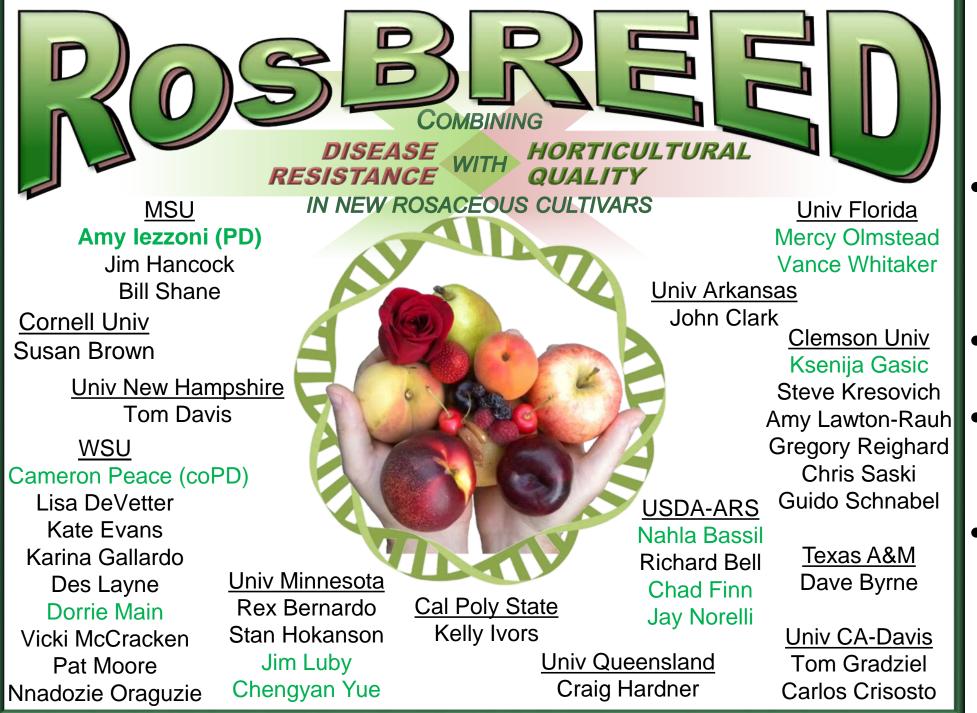
• DNA-informed breeding is now conventional!

U.S.-wide survey of Rosaceae breeders (early 2014):



Agree that "Genetic marker information will be included in 2014 in how my organization addresses plant breeding & selection"





#### "RosBREED 2"

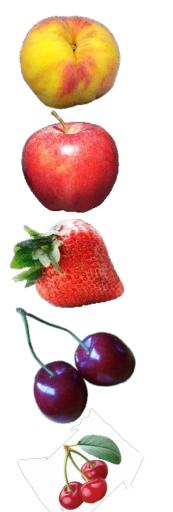
- Funded by USDA Specialty Crop Research Initiative
- \$10.0 M federal
- Sep 2014 Aug 2019
- Project Directors: *Amy lezzoni Cameron Peace*

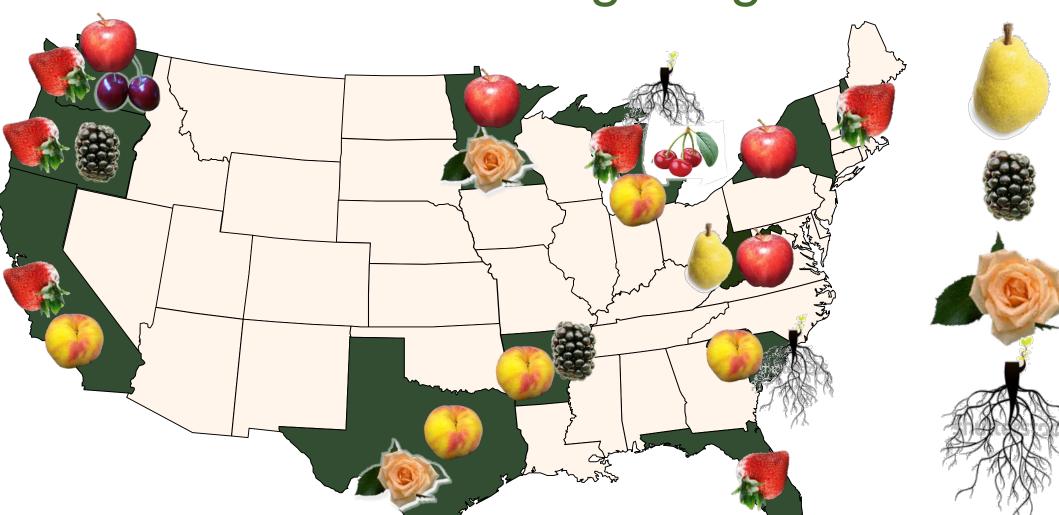
#### **RosBREED 2 Vision**

U.S. rosaceous crop breeding programs will exploit genetic resources, applying modern genomics tools to efficiently, accurately, creatively, and rapidly deliver new cultivars with market-essential horticultural quality and producer-required disease resistances to enhance consumer demand and mitigate stakeholder risk for rosaceous fruit, nut, and floral products.



# **RosBREED 2 Breeding Programs**

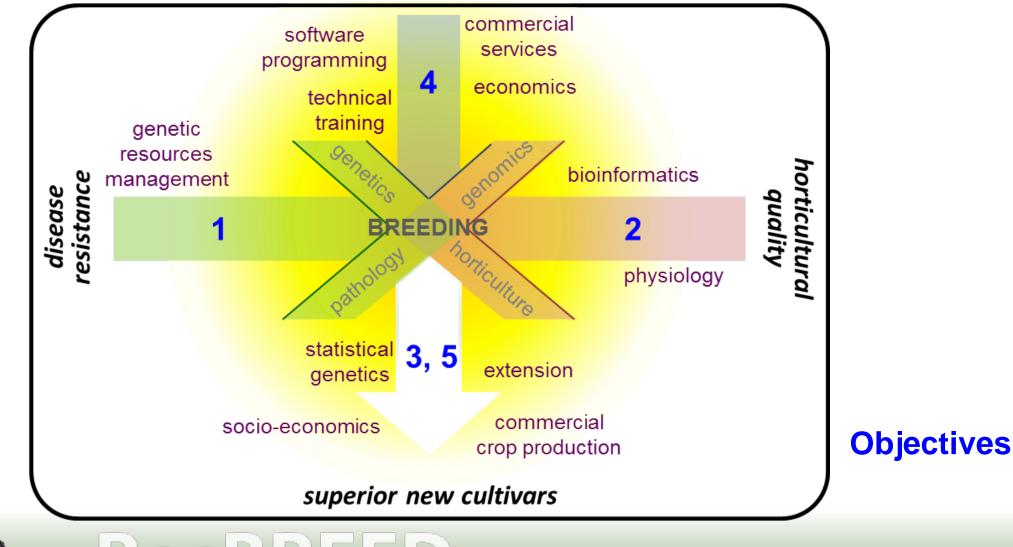






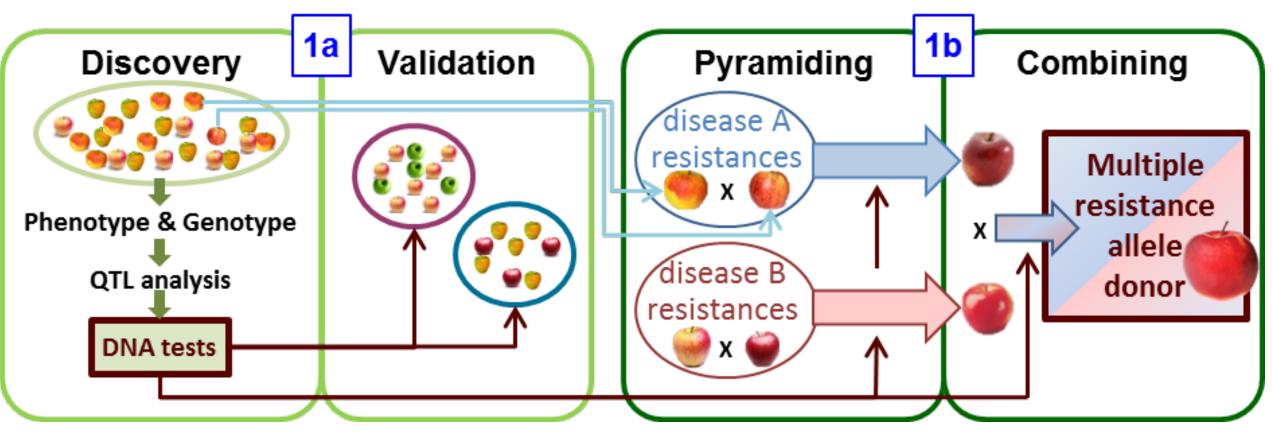
#### **RosBREED 2 Breedging Germplasm Levels** ONSUMPTION MARKETING PRODUCTION accurate predictions Innovative. Elite selection superior, Socioeconomics performance risk-mitigating informationnew cultivars enrich families Creative, efficient, Seedling selection accurate, rapid Parent selection breeding DNA ation information introgression Germplasm Accessible natural diversity development

## **RosBREED 2 Transdisciplinary Approach**



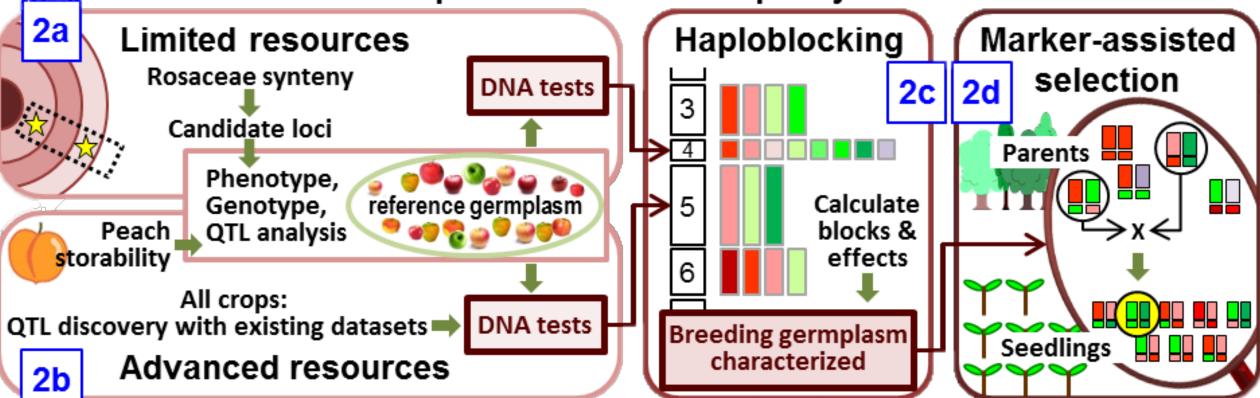


**Obj. 1:** Develop donor parents with multiple alleles for disease resistance



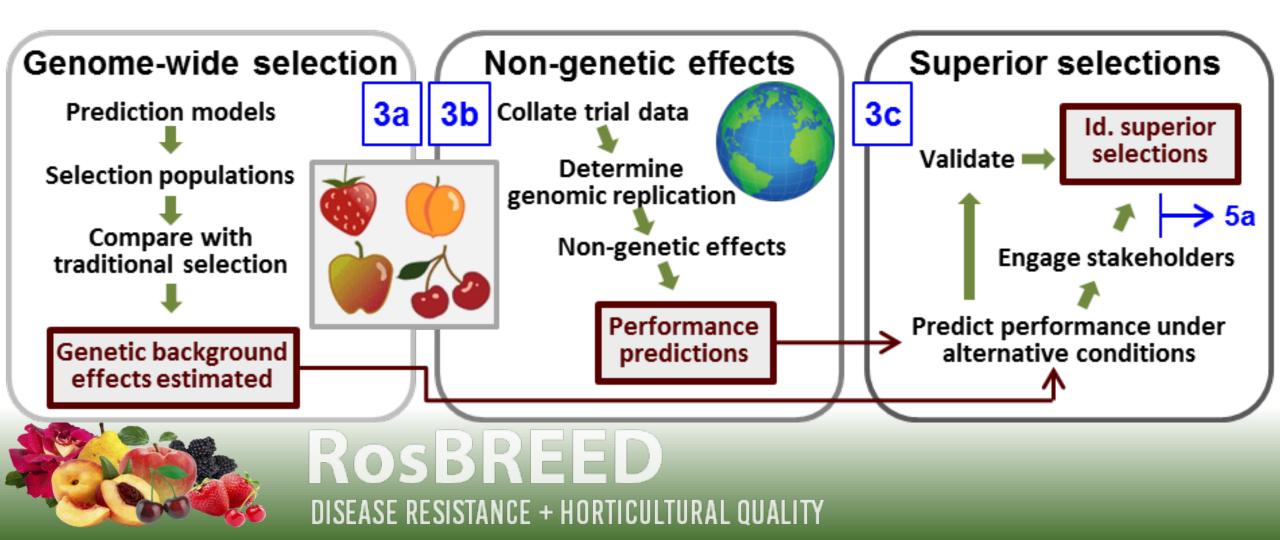


Obj. 2: Enrich breeding families with alleles for disease resistance and superior horticultural quality

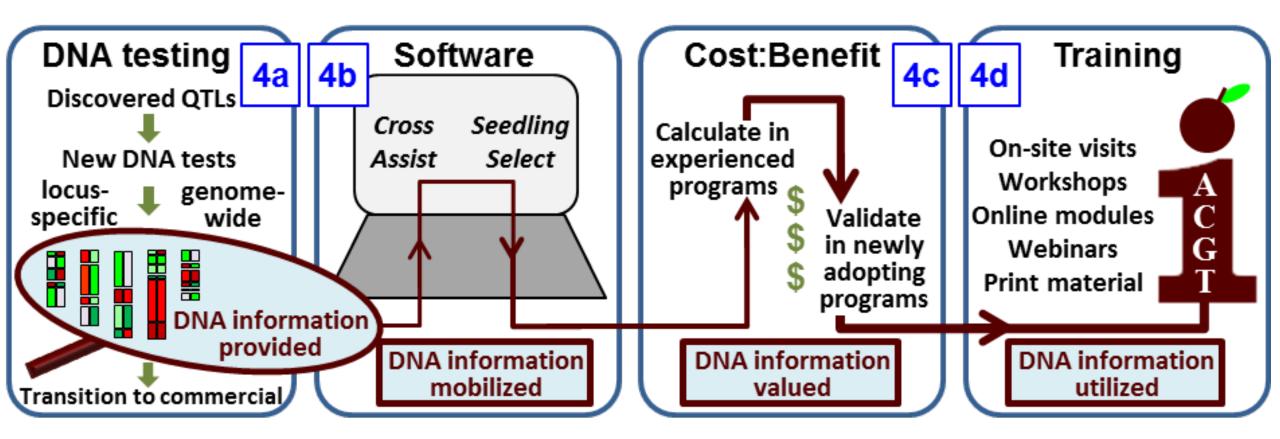




Obj. 3: Advance selections with alleles for superior horticultural quality and disease resistance with improved confidence

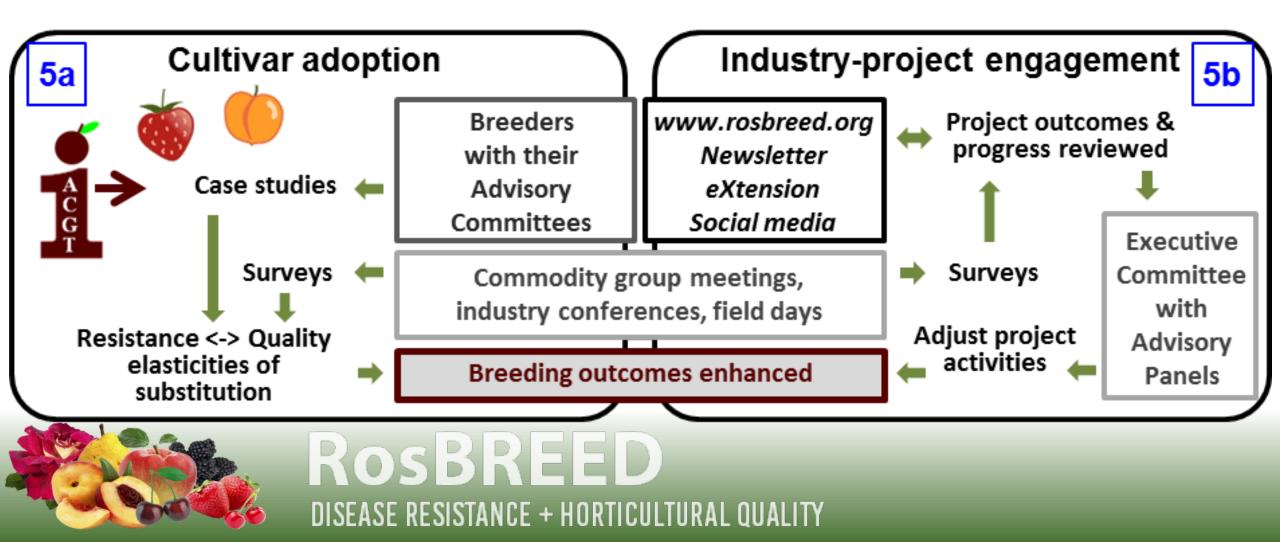


Obj. 4: Increase routine use of DNA information in rosaceous crop breeding

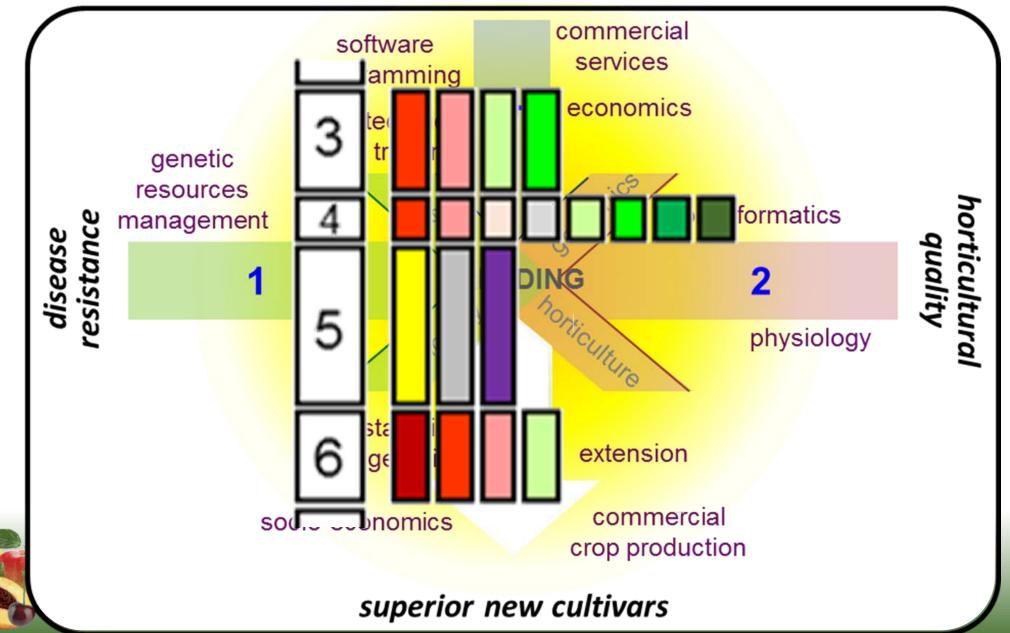




Obj. 5: Engage industry stakeholders in project outcomes, evaluation, and adjustment



# Haploblocking



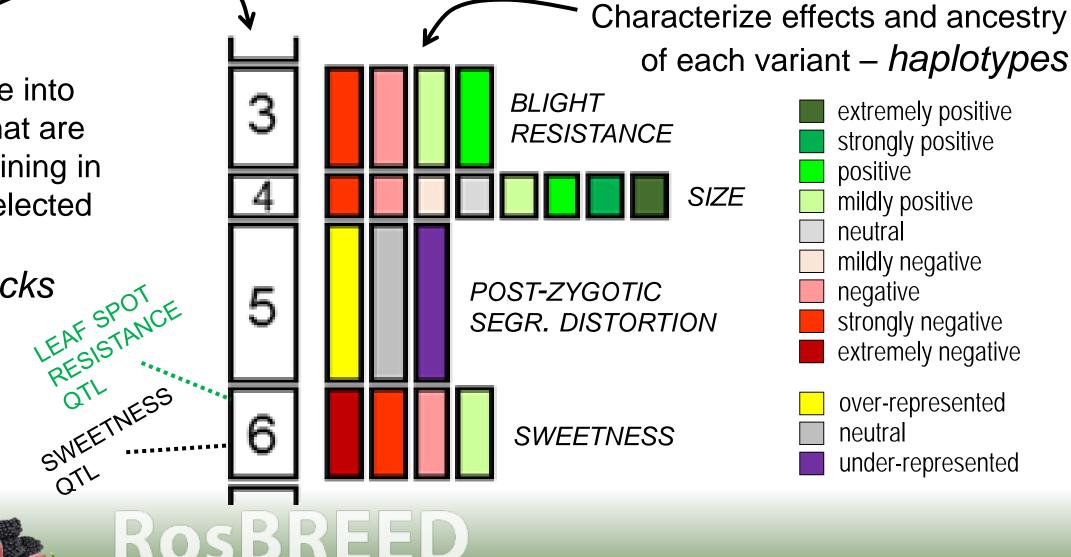
# Haploblocking

DISEASE RESISTANCE + HORTICULTURAL QUALITY

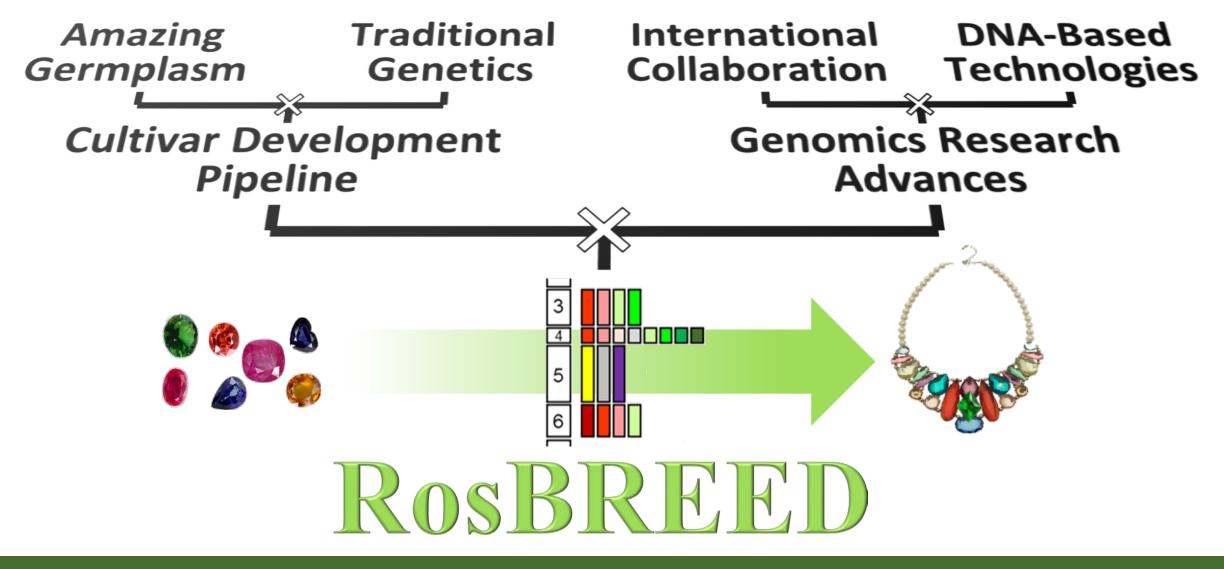
Divide each chromosome into segments that are non-recombining in historical, selected germplasm

haploblocks

Target tight linkages to break, if desired

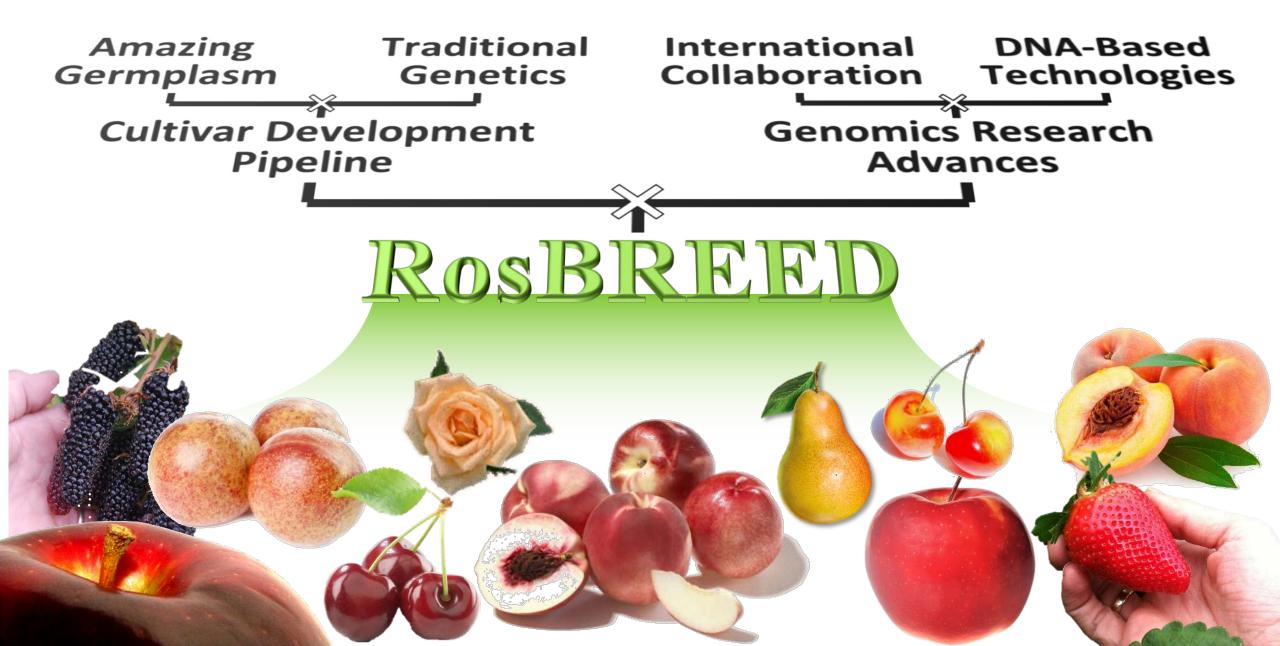


## The Pedigree of RosBREED



→ Enhanced breeding *efficiency*, *accuracy*, *speed*, & *creativity* 

#### The Children of RosBREED



# Acknowledgements





United States Department of Agriculture

National Institute of Food and Agriculture

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#### **Questions?**





United States Department of Agriculture

National Institute of Food and Agriculture