

# Accelerated Domestication and the Circular Carbon Economy

A perspective from perennial cropping systems

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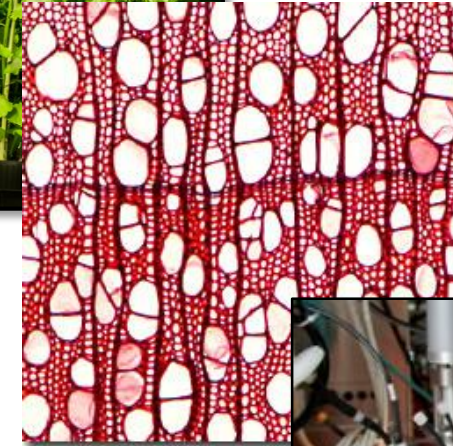


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# Presentation outline

- Poplar Life History
- Concept of Accelerated Domestication
- Genomewide Association Studies
- Targeted Phenotypes
- Epistasis and Pleiotropy
- Genomic Selection
- Poplar Ideotype
- Summary



# Poplar biology and a fully domesticated ideotype

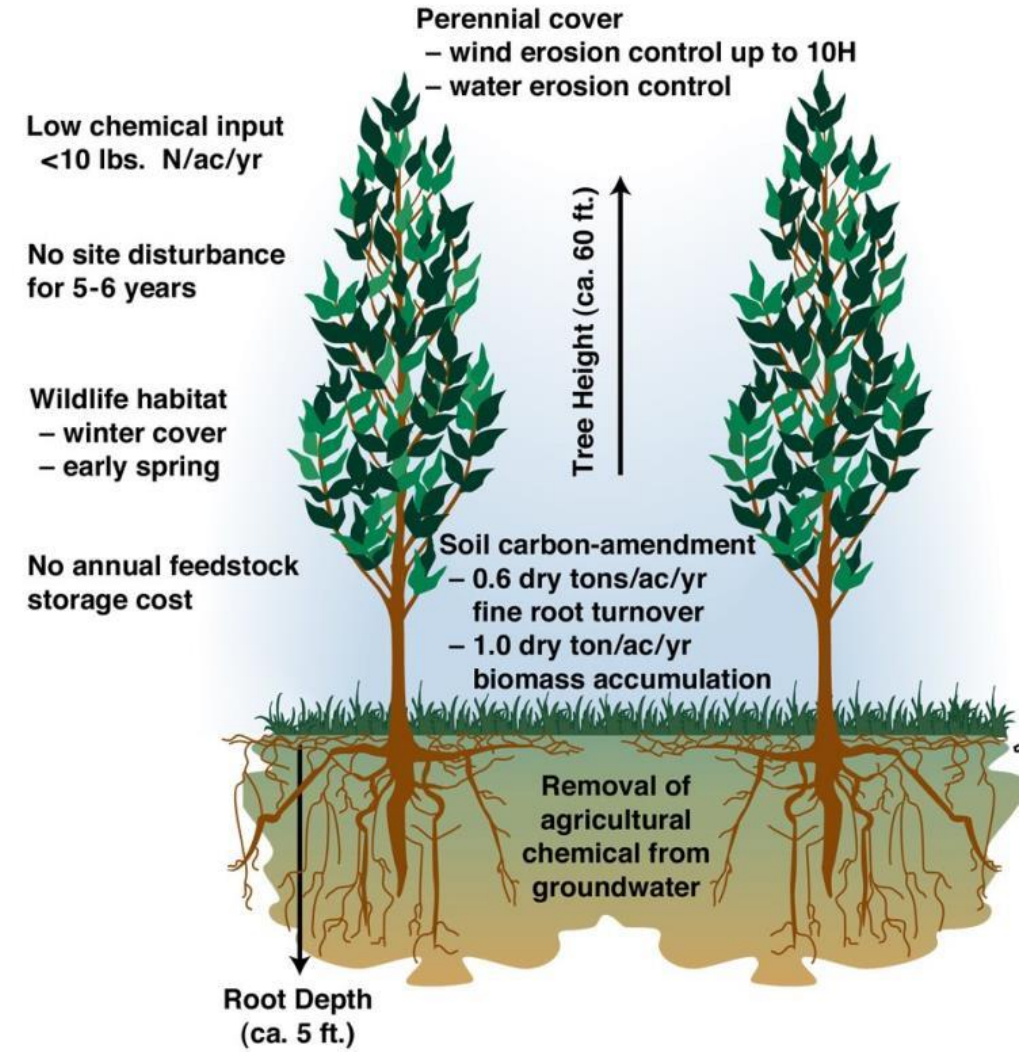
Native poplars are:

- Long-lived
- Dioecious
- Fast growing
- Undomesticated
- Perennial plants



Domesticated poplars also have:

- Higher productivity per unit area
- Greater number of stems per unit area
- Drought/Stress tolerance
- Nutrient-use efficiency
- Reduced recalcitrance of harvested tissues
- Greater product yield

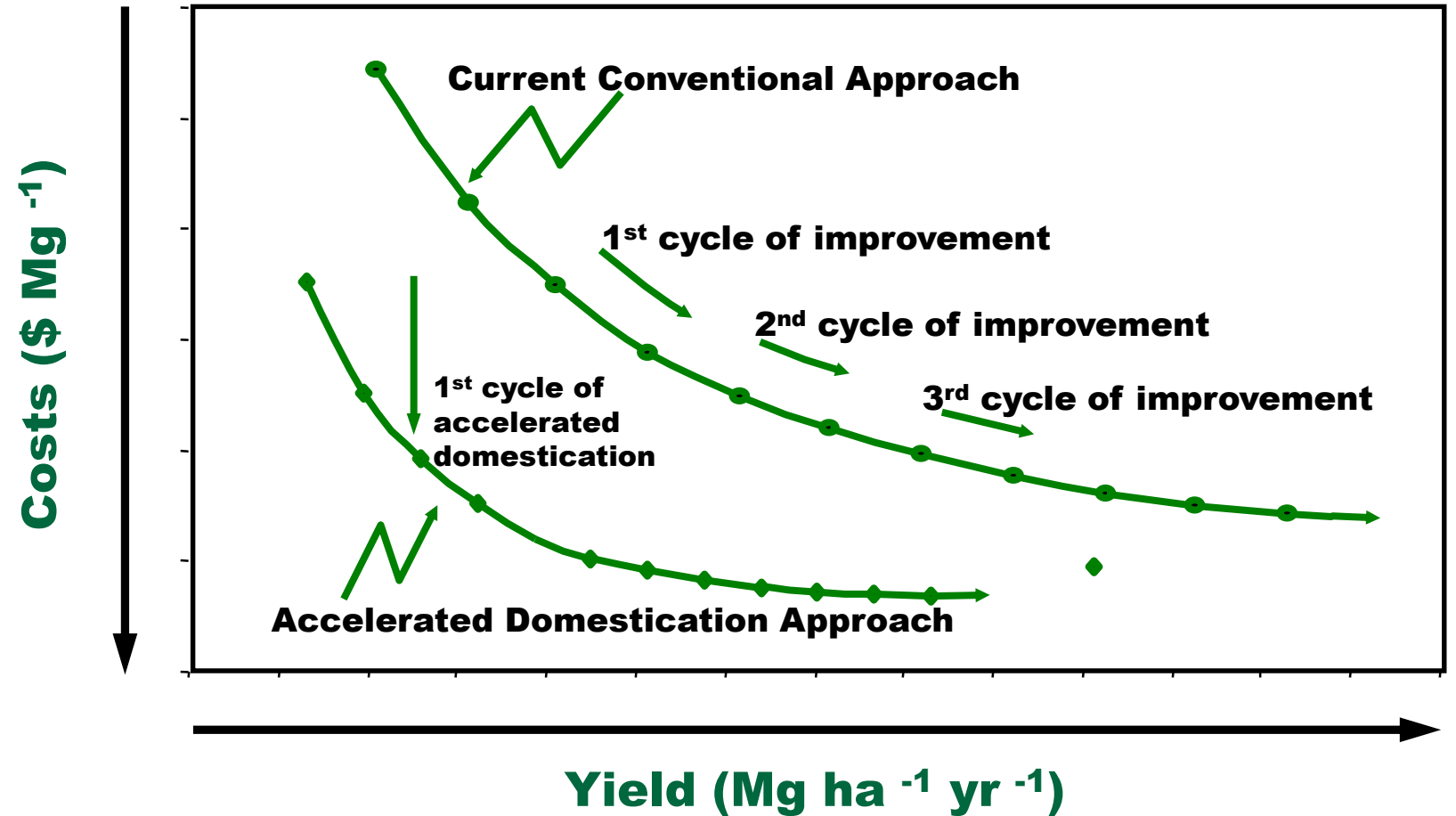




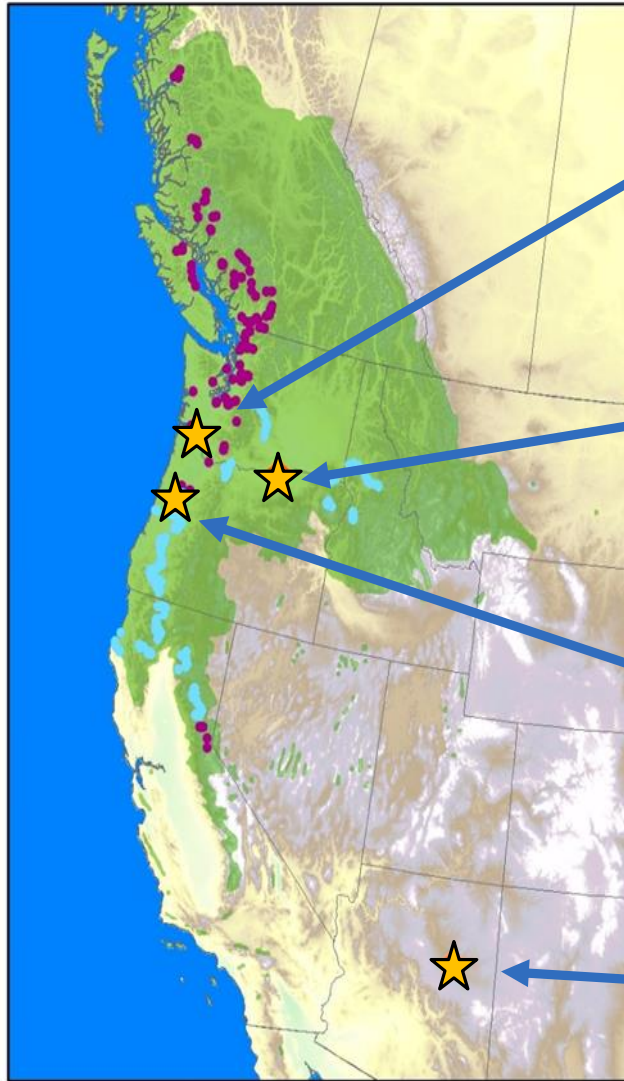
# There is a relationship between yield and cost

Traditional Breeding achieves incremental annual improvement

**Accelerated Domestication**  
disrupts the relationship  
between yield and cost



# Genomewide Association (GWAS) and Common Gardens



Clatskanie, OR (2009) Coastal Mesic



Boardman, OR (2016) Inland Xeric



Corvallis, OR (2009) Inland Mesic

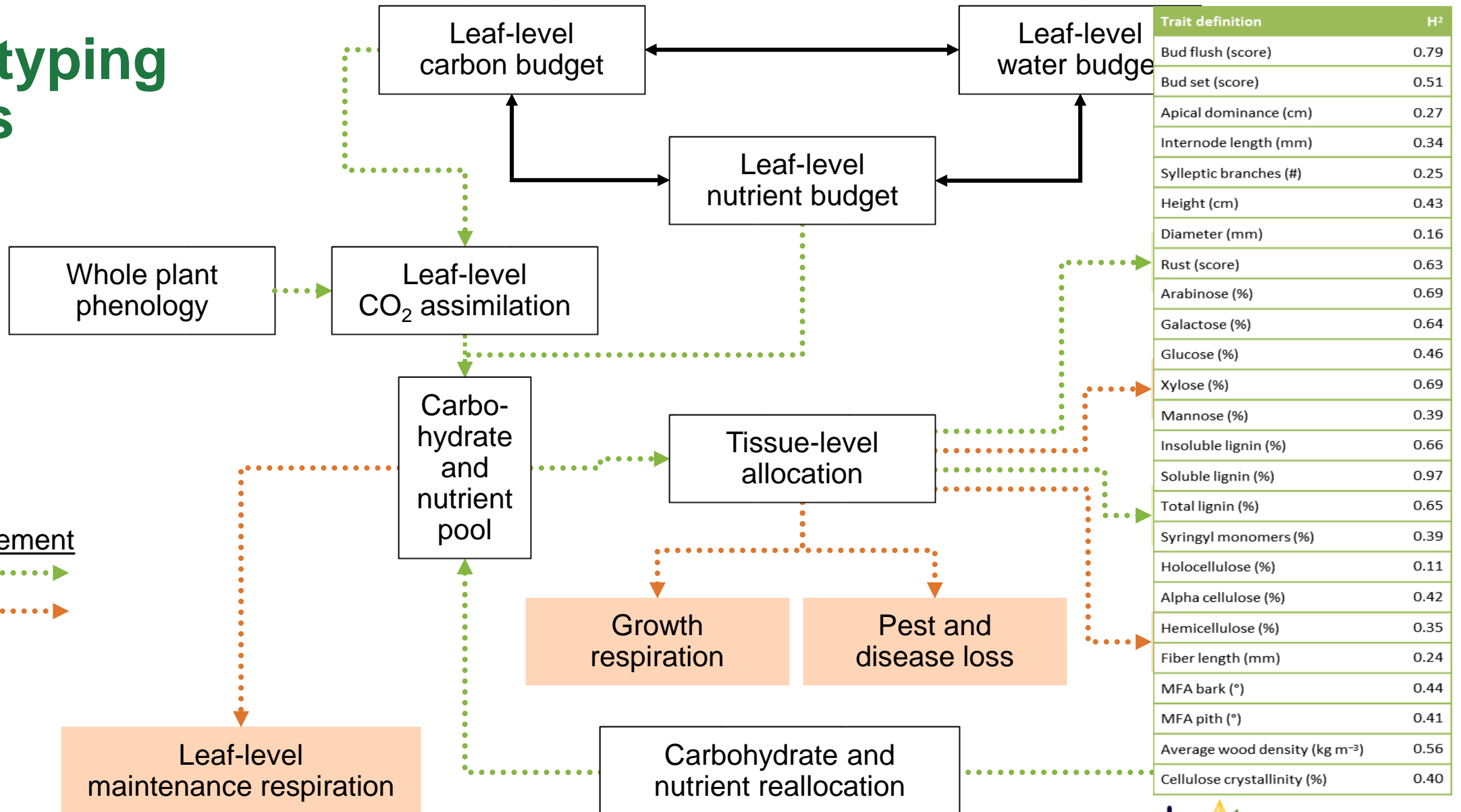


Sedona, AZ (2016) High Desert Xeric



- The selected population / panel
  - 1250 genotypes
  - Southern BC to Northern CA
  - Established in 4 common gardens
- Genotyping
  - Resequenced at a minimum of 18x depth
  - 68 million high-quality SNPs
  - A SNP every 17 bp
  - LD decays on average within 300 bp and in many cases within <20 bp
- Transcriptome
  - RNAseq data was generated for the 1250 genotypes for leaves, xylem and roots

# GWAS phenotyping targets

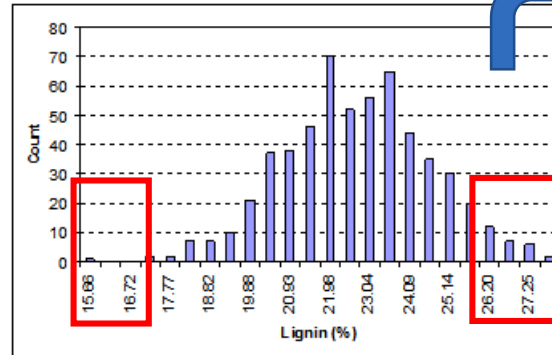


# Genome-wide association approaches

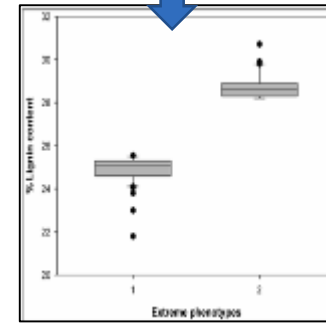
Assemble a population



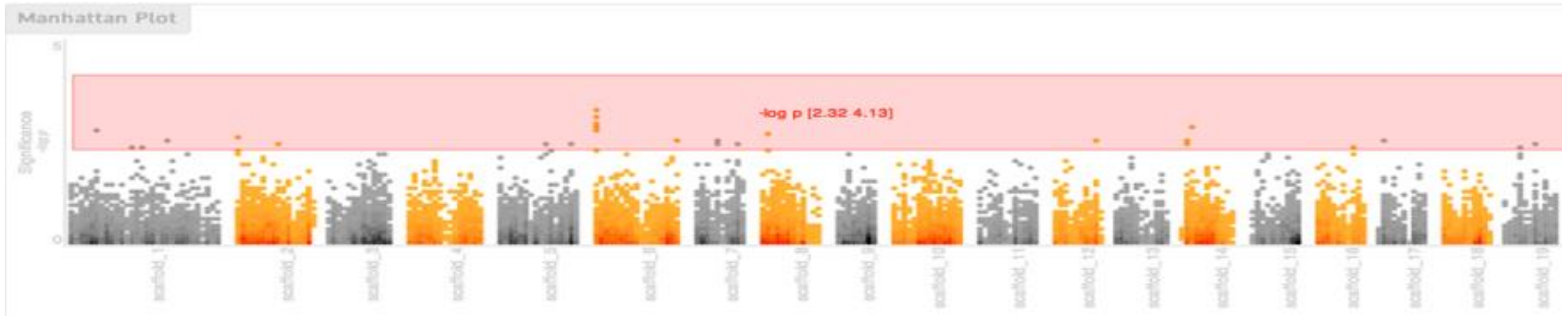
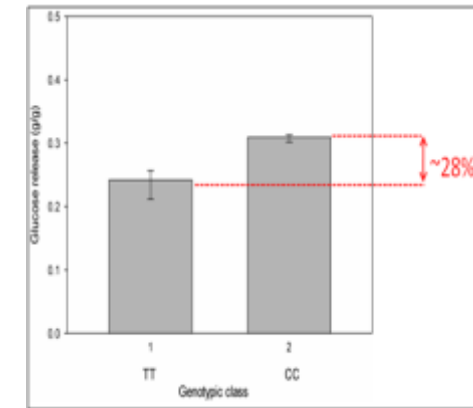
Phenotype the population



Create a SNP library

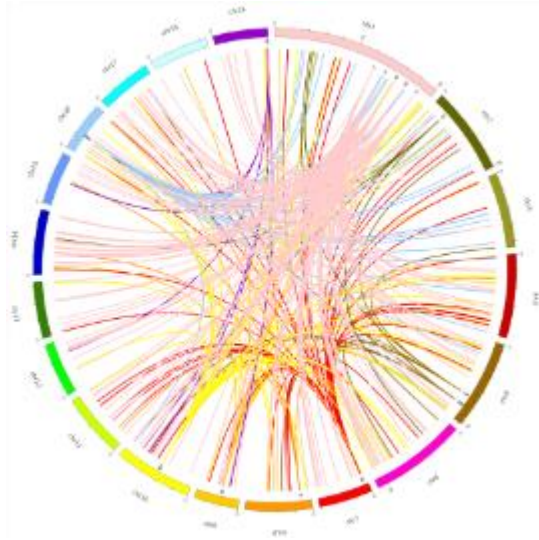


GWAS Analysis

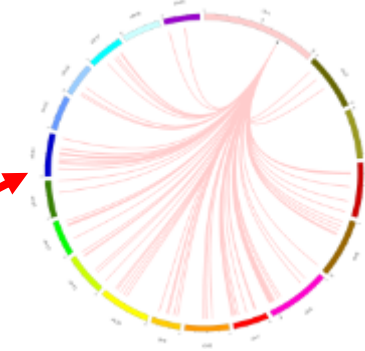


# Epistatic Network: Transcriptional Hotspots

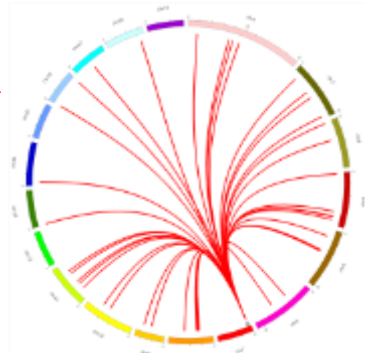
Trans-acting eQTN hotspots  
(i.e. >20 co-expressed loci)



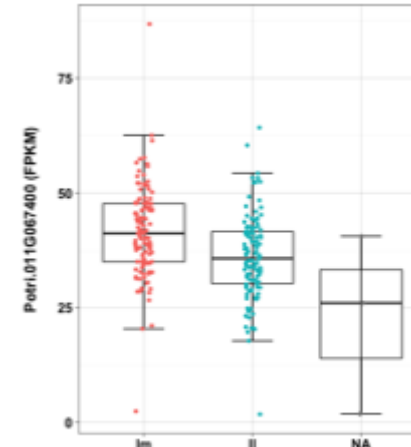
Trans-acting eQTN hotspots  
on chromosome 1



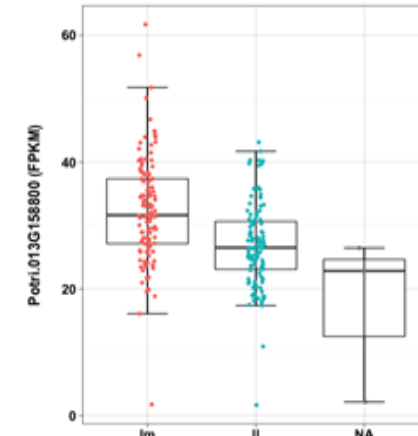
Trans-acting eQTN hotspots  
on chromosome 7



Co-expressed Loci

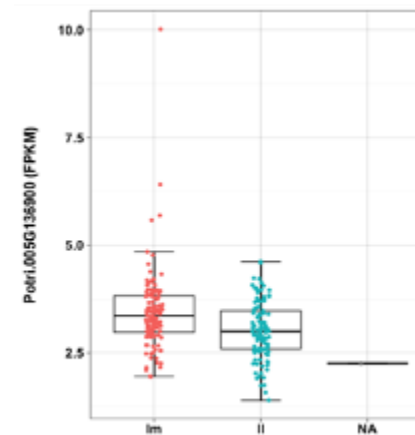


Gene A

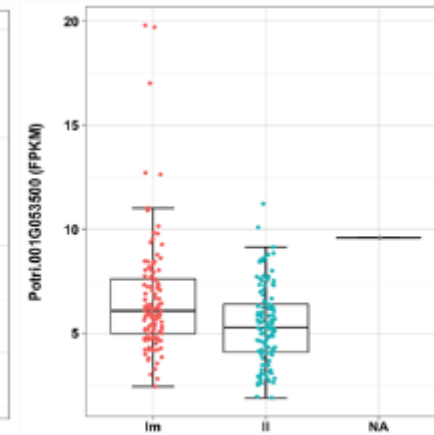


Gene B

Co-expressed Loci



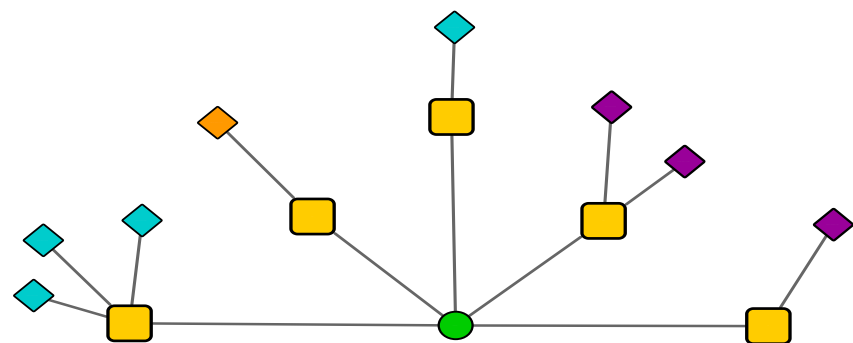
Gene C



Gene D

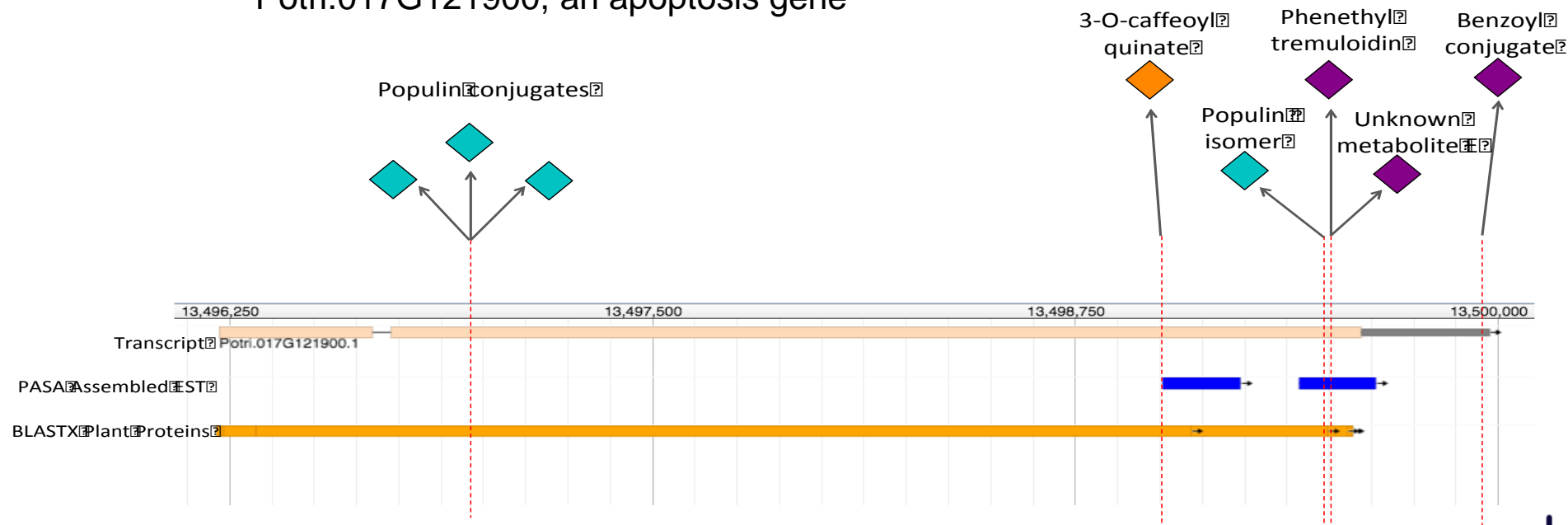


# Example of Complex Pleiotropic Signatures

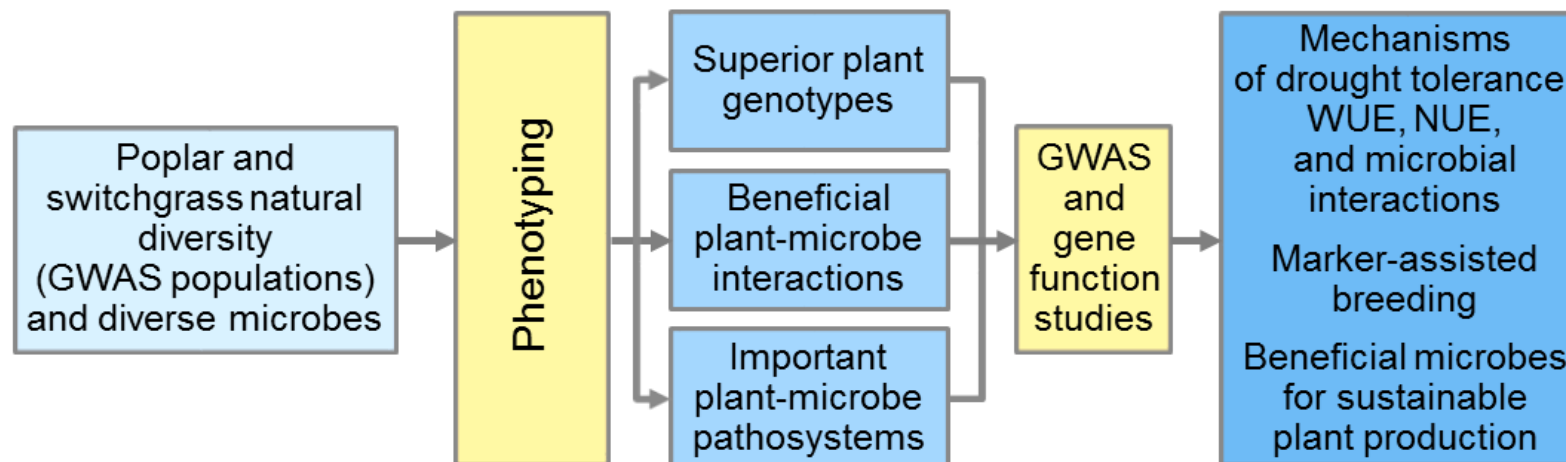
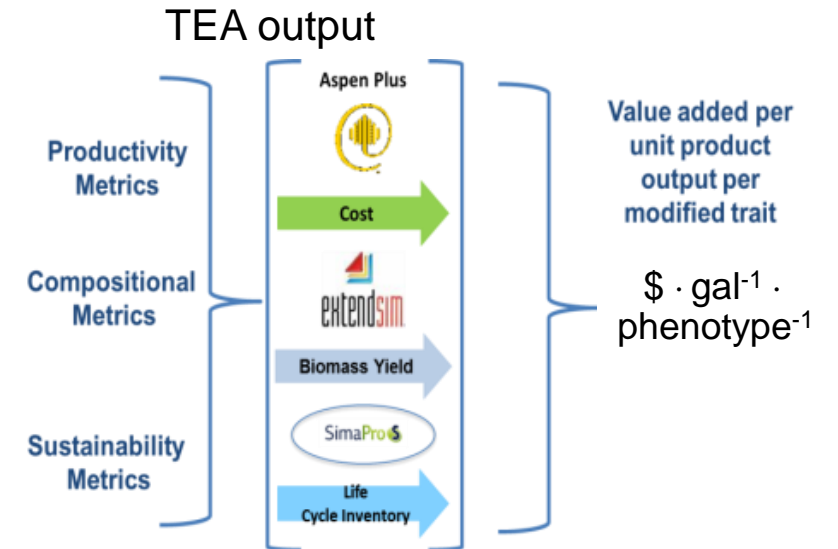
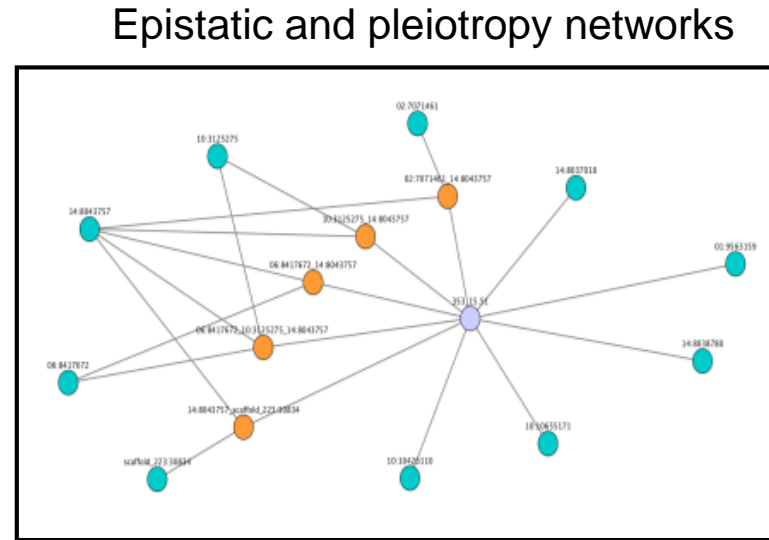
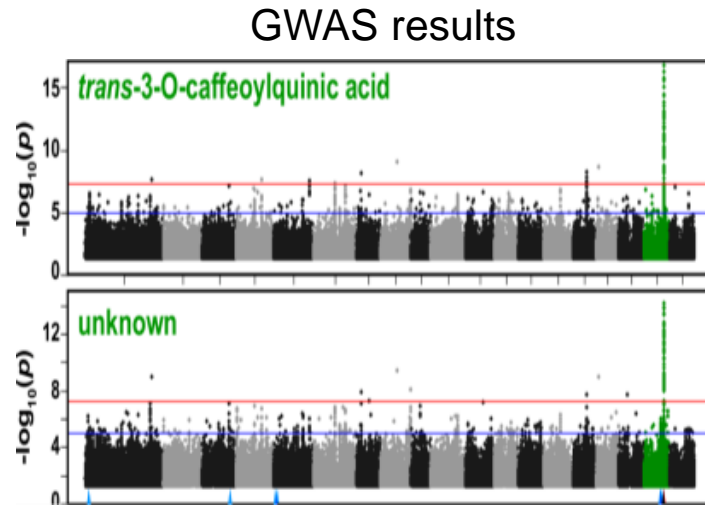


- Module
- Gene
- Phenotype (metabolite)
- Salicylate-related (metabolite)
- Lignin-related (metabolite)

Potri.017G121900, an apoptosis gene



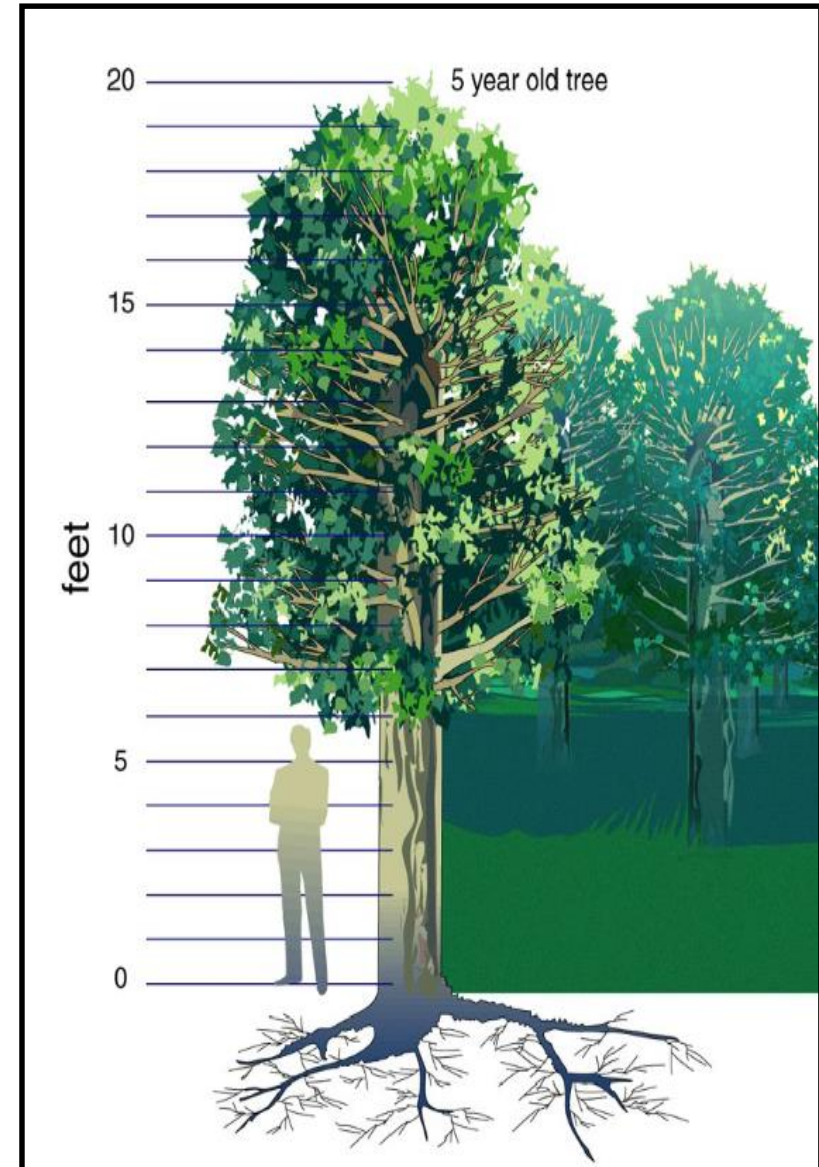
# Genomic selection and technoeconomic analysis



# Genomic selection can disrupt the yield to cost relationship and accelerated the domesticated of poplar

## Domesticated Ideotype

- **Higher productivity per unit area**
  - Reduced flowering
  - Reduced height growth
  - Compact crown
  - Enhanced radial growth
- **Greater number of stems per unit area**
- **Water-use efficiency**
- **Nutrient-use efficiency**
- **Favorable plant-microbe interactions**
- **Greater product yield**
  - Reduced stem biomass recalcitrance
  - increased root recalcitrance
  - Greater rooting depth



# Poplar feedstock, biofuels and bioproduct targets and goals

## Sustainable Feedstocks

**Yield** (biomass per unit area per unit time)

- >50% improvement

### Sustainability

- Water-use efficiency (WUE): <10% yield loss under drought
- Nitrogen-use efficiency (NUE): >20% improvement
- Pest and pathogen resistance
- Favorable microbial associations

### Feedstock uniformity

- Biomass composition over time and environments

## Biofuels Targets

### Fuels portfolio

- >80% carbohydrate solubilization at >120 g/L solids loading
- > 30 g/L iso- and/or n-butanol
- > 1 g/L C6 esters (e.g., ethyl butyrate, butyl butyrate, butyl acetate)

## Bioproduct Targets

### Microbial products

- 2-pyrone-4,6-dicarboxylic acid,  $\beta$ -ketoadipate, cis,cis-muconate

### Catalytic products

- 4-propyl guaiacol and 4-propyl syringol

### Materials

- Polyurethanes and foams from high molecular weight lignins



# CBI research partners and sponsor

## CBI research partners:

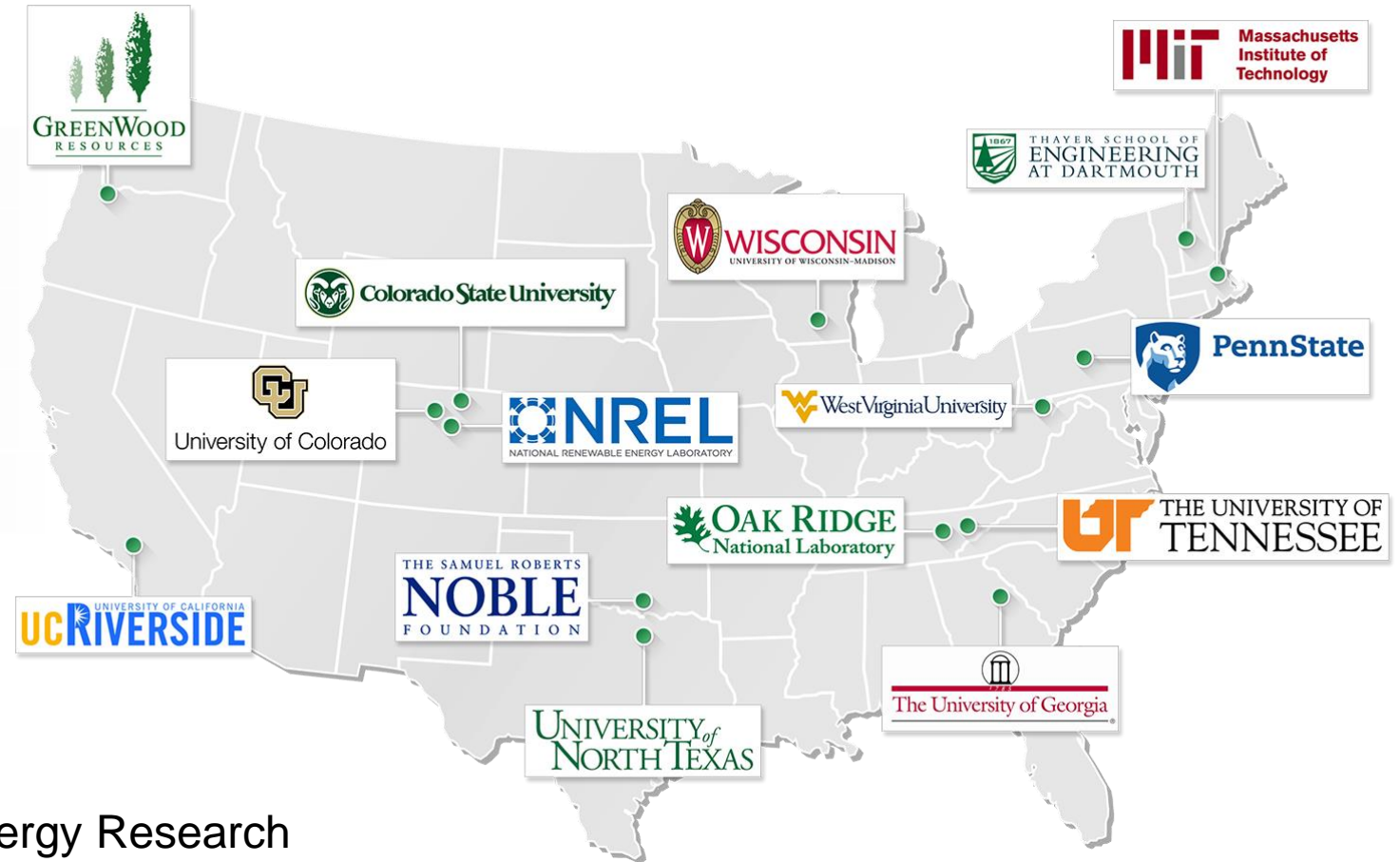
2 national laboratories

11 academic institutions

1 research foundation

1 private company

40% new PIs and/or institutions



CBI is a U.S. Department of Energy Bioenergy Research Center supported by the Office of Biological and Environmental Research in DOE's Office of Science

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Sustainable Cropping Systems



Domesticated Feedstocks



Advanced Biofuels and Bioproducts



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