



Imperial College
London



Poplar for Bioenergy: The ENERGYPOPLAR PROJECT

9 Partners, 3M (EU) Total 4.1M

Gail Taylor,
24 March, 2009
Madrid

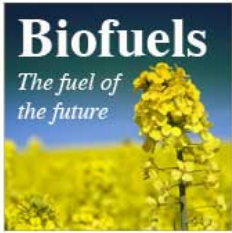


UNIVERSITY OF
Southampton
School of Biological Sciences

Policy Drivers for increased bioenergy

- 10% Renewable energy by 2010 and 20% by 2020. UK, European SET plan.
- 60% reduction in CO₂ emissions and first nation to have a climate change bill
- RTFO – 5% by volume liquid fuels from renewable source by 2010 and 10% by 2020
- Bioenergy currently contributes
 - 4.1 % to heat and electricity
 - 2.5 % to liquid transportation (April 2008, from 0.5%)

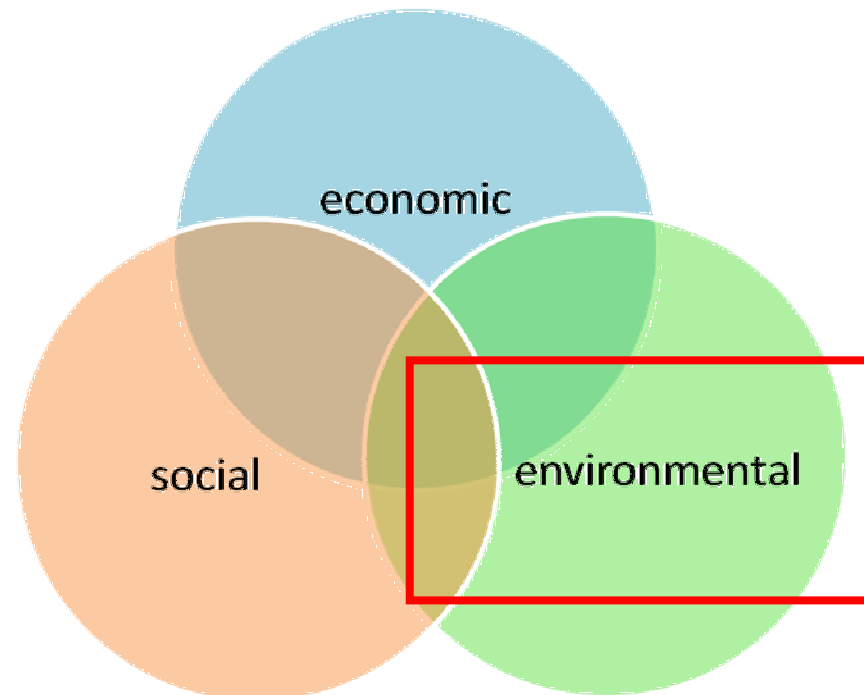
Taylor (2008), *Energy Policy In Press*



Bioenergy -concerns

- Greenhouse gas balance?
- Biodiversity? Loss of species?
- Other impacts?
- Food crop displacement and prices?
- Other indirect effects... tropical rain forests?
- Societal concerns? GM?
- Landscape-scale changes?
- Water use and management?

Bioenergy Research – what are the show stoppers?



Feedstocks for bioenergy?

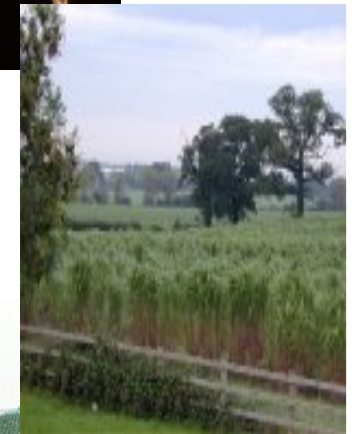
First generation crops

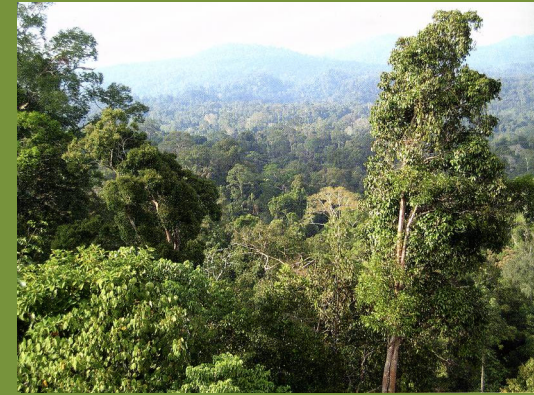
- Maize
- OSR
- Wheat grain
- Sugar beet
- Other oil crops from tropics -import
- Sugar cane (Brazil) - import



Second generation dedicated lignocellulose

- Trees – willow and poplar
- Grasses- miscanthus, switchgrass ,
- Others





Forest ecosystems – 50% global vegetation C

Estimated total amounts of C in soil & vegetation in different terrestrial systems (Royal Society 2001). Pg C or Gt C

croplands	131	boreal forests	559
wetlands	240	temperate forests	159
tundra	127	tropical forests	428
deserts & semides.	199		
temperate grasslands	304		
tropical savannas	330		

Total non forest 1,331

Total forests 1,146



Feedstock: Woody lignocellulosic biomass (2GC)

- Heat, power and liquid fuels (biological and thermochemical)
- Vast UK, EU and global resource
- Good GHG balance?
- Grown on marginal and abandoned land..... No food v fuel
- Biodiversity? Water use?



What feedstock traits need improving? Ideotype breeding?

Biomass Quantity

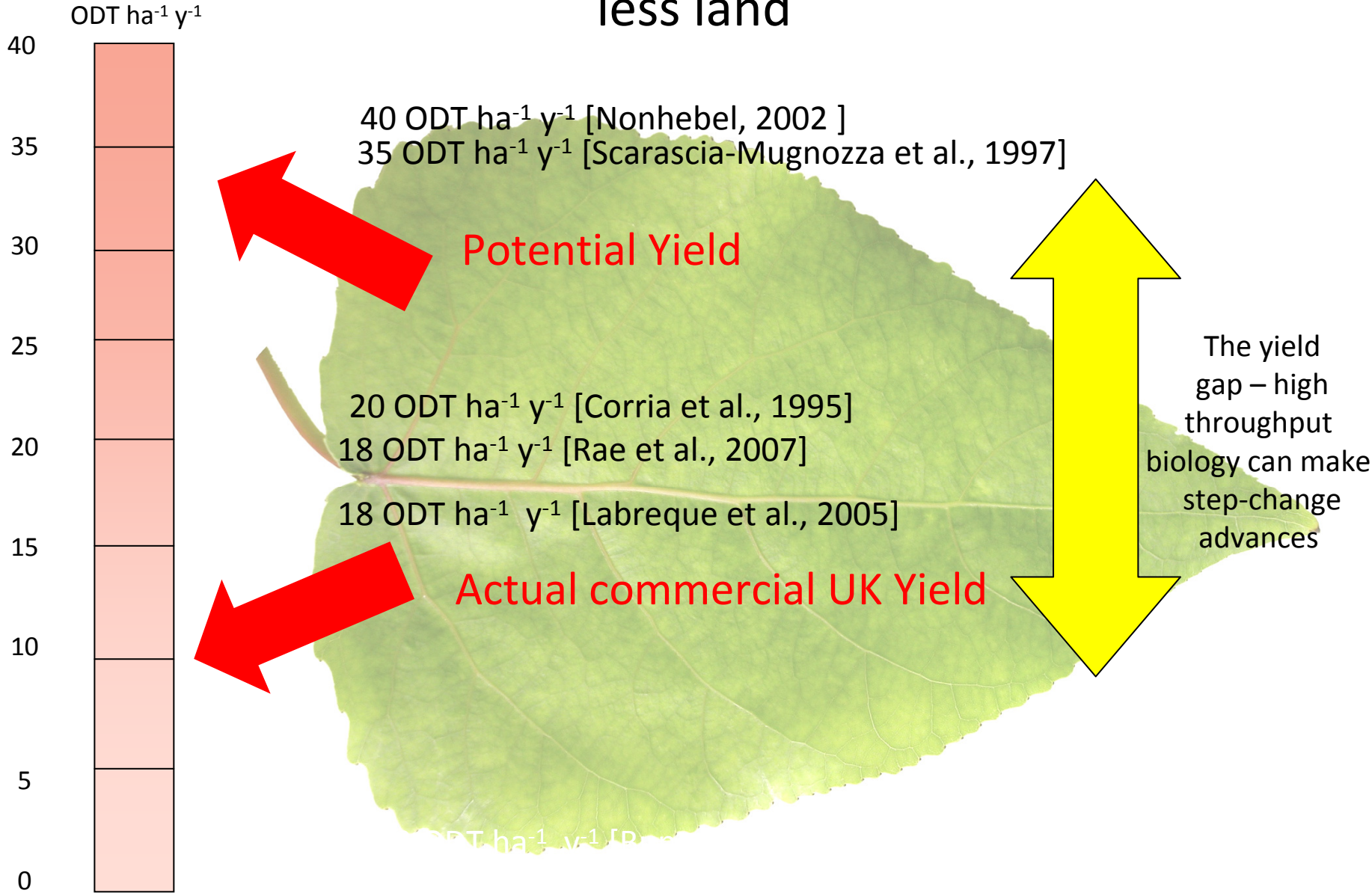
- More biomass per hectare
- Limited N and water inputs

Biomass Quality

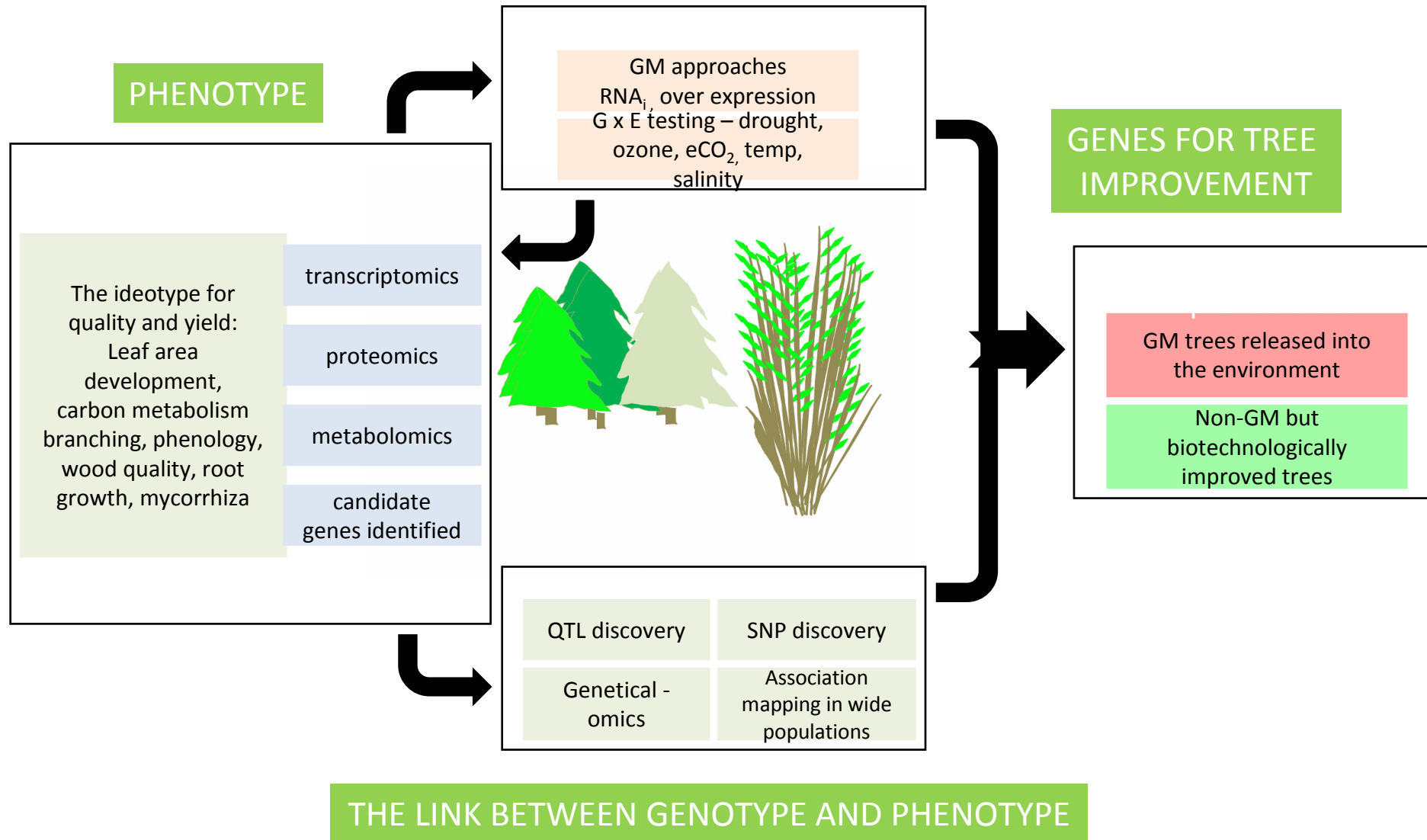
- Improved quality
 - lignin
 - cellulose
 - hemicellulose
- saccharification
- calorific value

Sustainable – GHG balance, biodiversity, water use,

Addressing the yield gap in energy poplar – more biomass on less land



Systems Biology for optimised yield and lignocellulosic quality in energy



Poplar for bioenergy – globally significant

More than 30 species, wide range of climates and soils

Large genetic diversity, large natural provenances (91%)

Interspecific hybridisation



P. SIGAUD



FAO; Ball, Carle and Del Lungo

J. CARLE

Why is poplar the ideal bioenergy model?

Full genome sequence (15 Sept 2006)



Small genome (similar to rice)

Diversity of species

Fast growing and perennial

Clonal

Transformation system

Genetic collections and molecular maps

Activation tagged populations

EST and BAC libraries

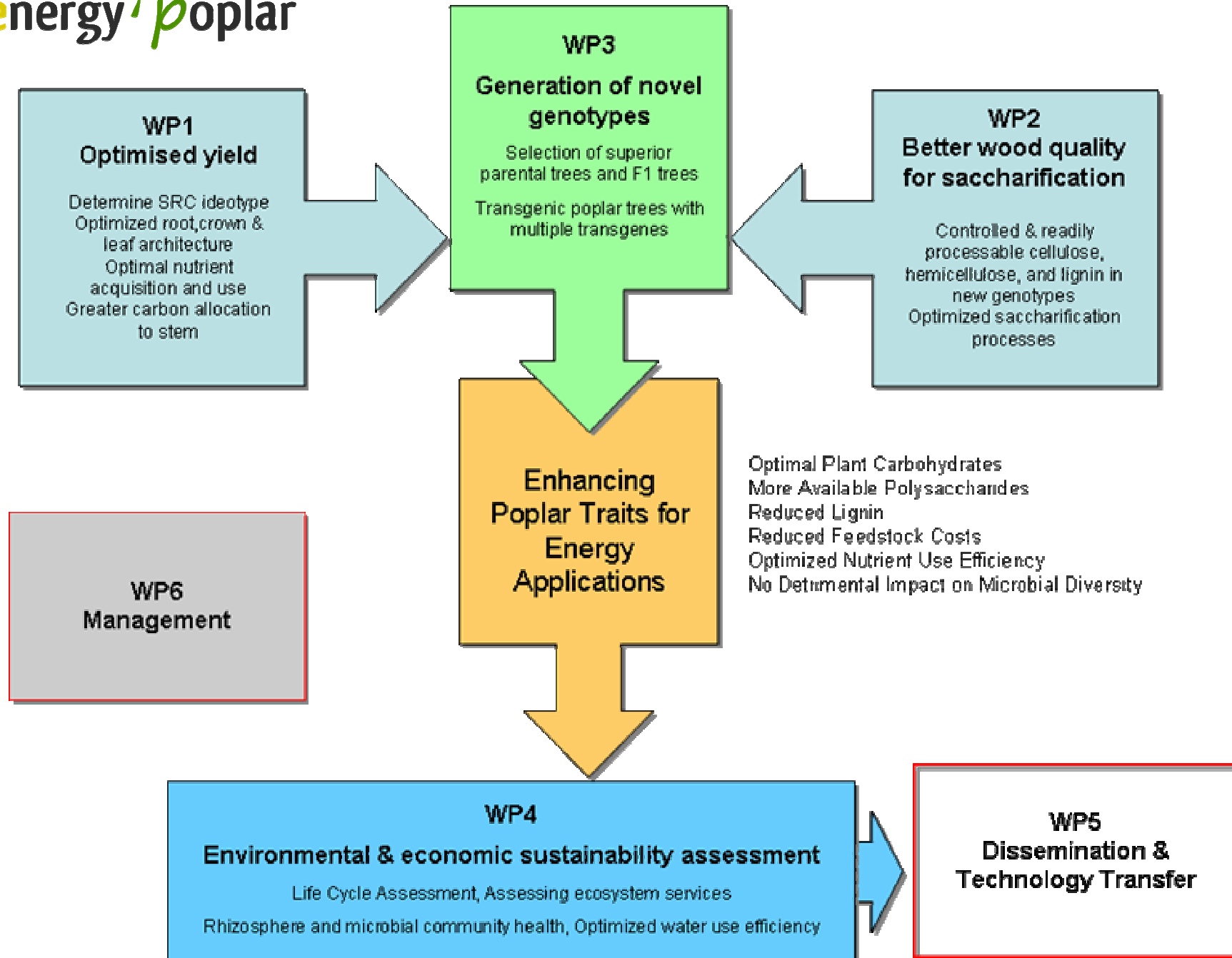
Genomic and post-genomic tools

Aims of ENERGYPOPLAR

- The aim of the project is to improve feedstock characteristics of poplar for bioethanol production
- Improved yield
- Improved saccharification
- Improved sustainability and water use efficiency
- Optimized LCA



- INRA, France- (F Martin) -mapping pop, association, mycorrhizal associations, genomics, ESTs, EVOLTREE, GM low lignin trees
- UPSC, Sweden (R Bhalerao) - GM lines, HT phenotyping cell walls, transcriptomics, metabolomics, gene discovery platform
- Southampton, UK (G Taylor)- mapping pop, association, Carbon isotopes, QTL elucidation, saccharification
- Udine, Italy (M Morgante) – Genotyping, HT SNPs discovery,
- Ghent, Belgium (W Boerjan) – GM low lignin trees, mapping pop
- Gottingen, Germany (A Polle) – GM trial, carbon isotopes, HT
- Imperial College, UK (R Murphy)- LCA , HT saccharification assay
- ALU, Germany (K Palme) - Gene discovery, Systems Biology
- SweeTreeTechnologies (M Hertzberg)- commercialisation

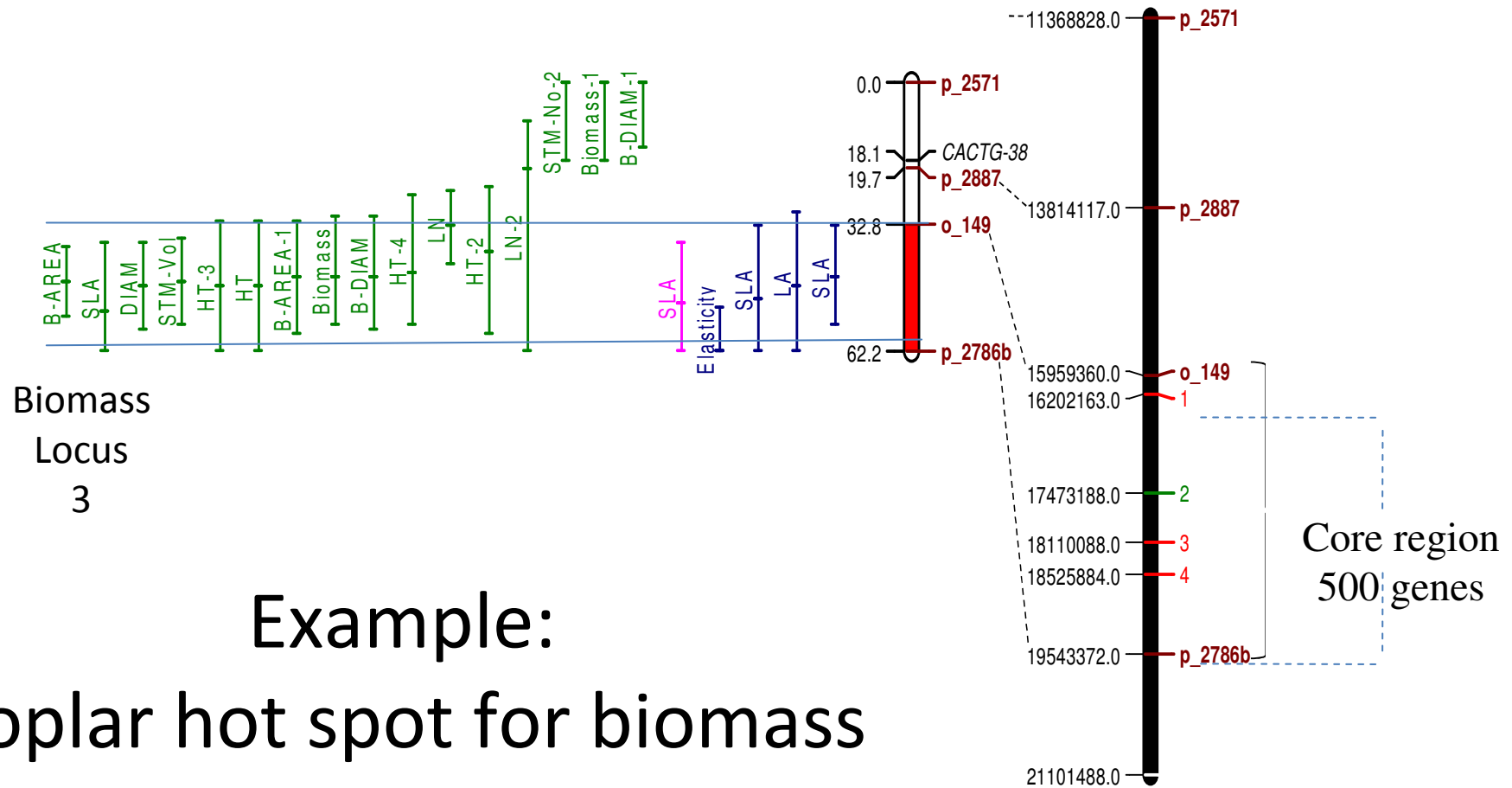


Workpackage 1

- Improve yield through genomics and genetical genomics
- Optimise shoot function using GM technologies
- Optimise root function using GM technologies



Five 'Poplar BIOMASS LOCI' identified in *Populus* following 10 years of research

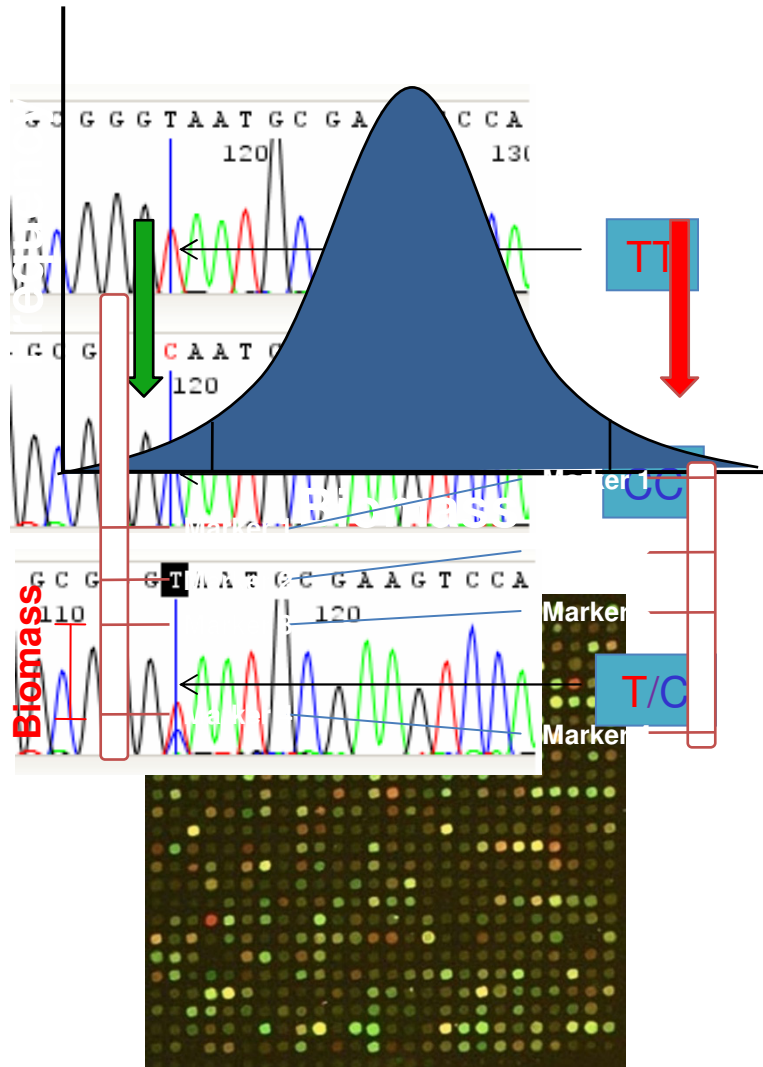


Biomass
Locus
3

Example:
Poplar hot spot for biomass
traits

Core region
500 genes

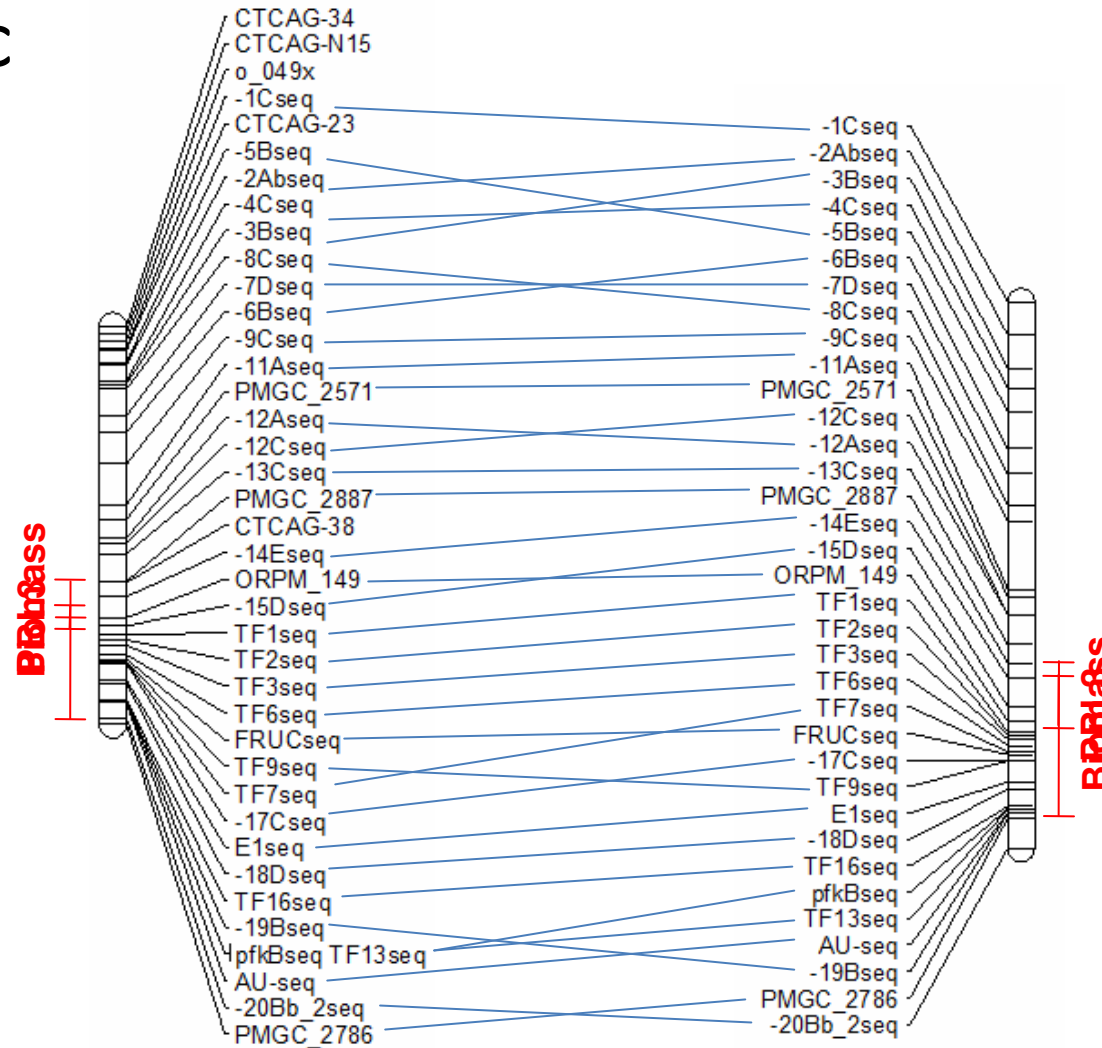
Genetical genomics approach



- Improve density of markers on the key biomass LG and in PBL3 using SNPs.
- Conduct microarray experiment using 'biomass extremes'.
- Aim to identify key genes in the core regions of the QTLs:
 - Transcription factors potentially controlling biomass
 - Biomass related genes e.g. cell wall synthesis, sugar synthesis, cell cycle etc

Genetic

Physical



energy poplar Workpackage 2

- Establish GM field trials of current and new material e.g. low lignin trees
- Establish high throughput phenotyping for lignin and cellulose traits
- Establish association genetics populations
- Undertake saccharification assay on 500 plus genotypes of *Populus* – SRC, GM, others



What is Saccharification?

- “The process of breaking down a complex carbohydrate into monosaccharide components”
- Breaking down ligno-cellulosic mass into glucose.
- This glucose can then be fermented to produce ethanol.

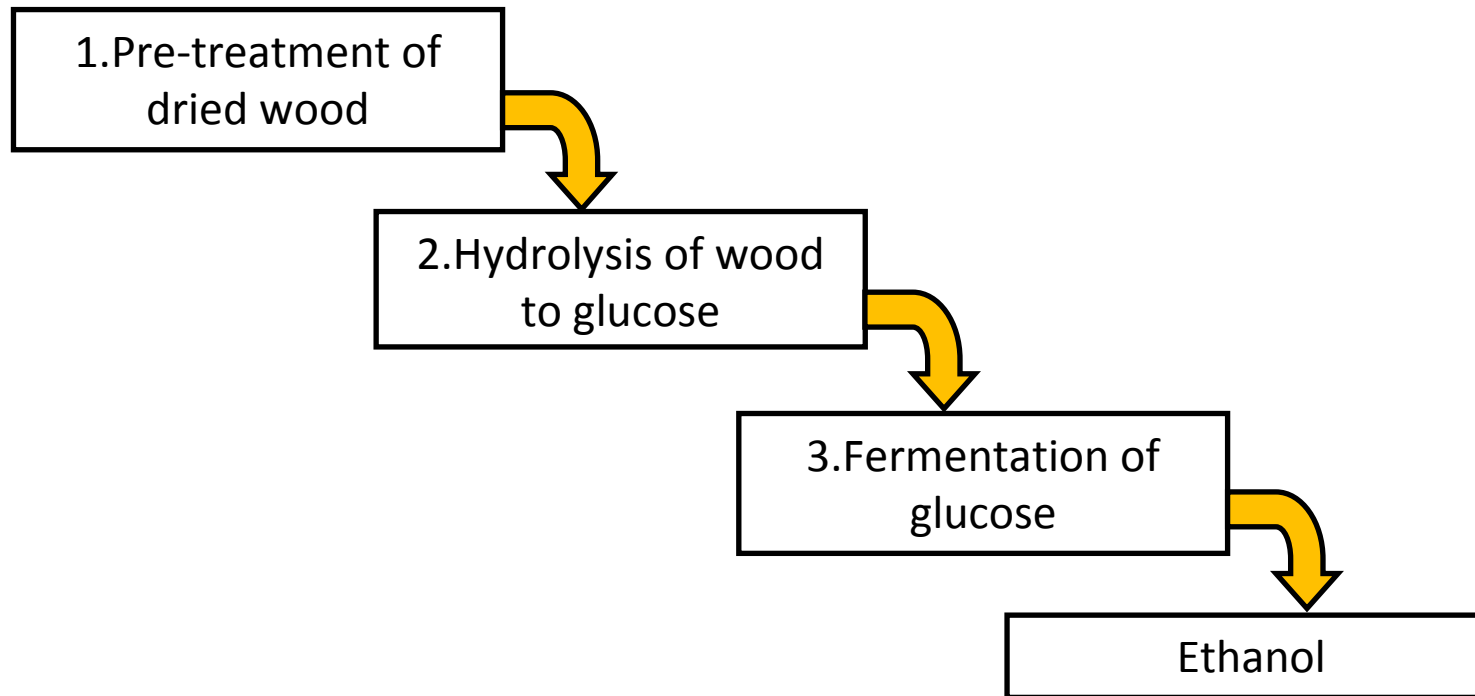


Saccharification potential of *Populus*?

- Identify natural genetic variation in glucose release across Family 331 genotypes (F2 mapping population) using a high throughput assay.
- Combine data with existing F331 markers to produce QTL and identify genes of interest.

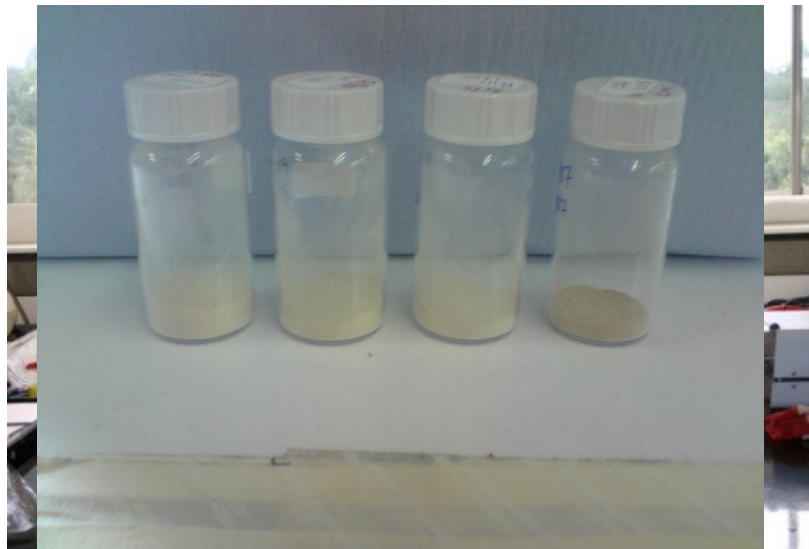


Overview of Process



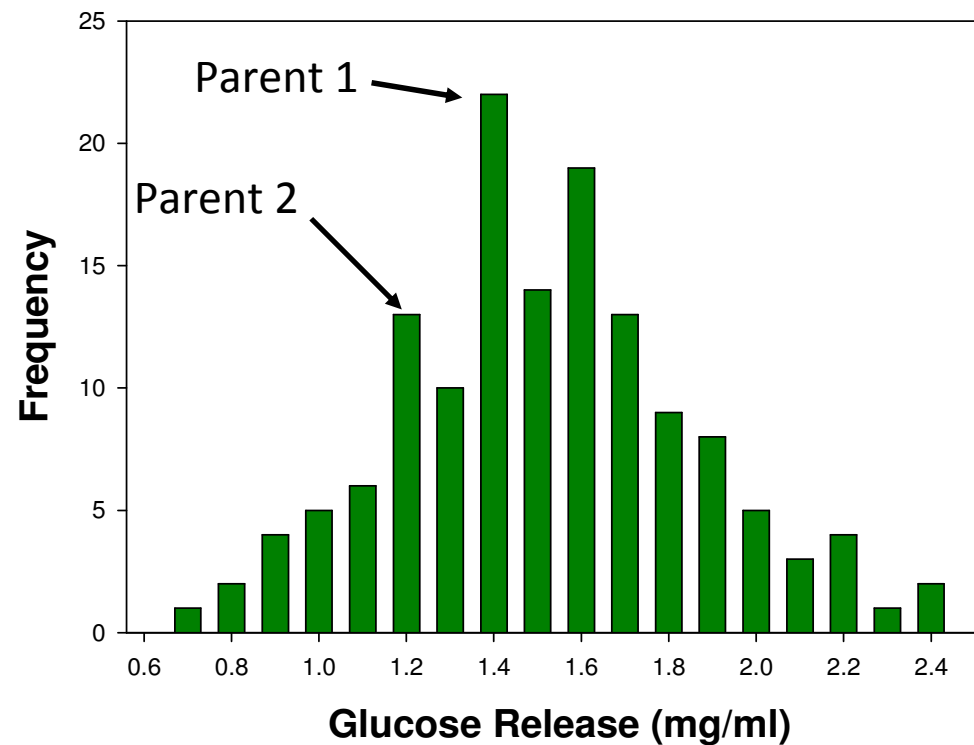
Wood Pre-treatment

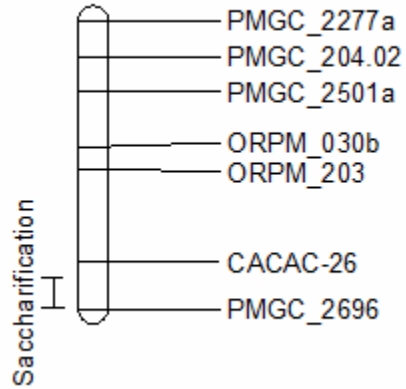
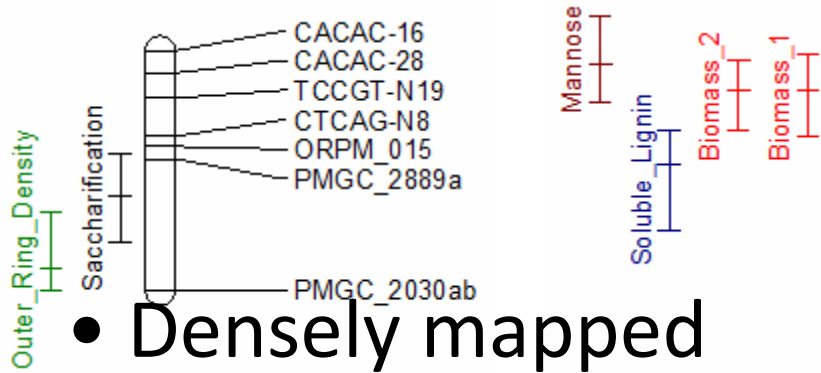
- Wood ground to fine powder using 'Retsch 300MM Mixer miller'.



Results

Glucose release across F331 genotypes after 48 hours

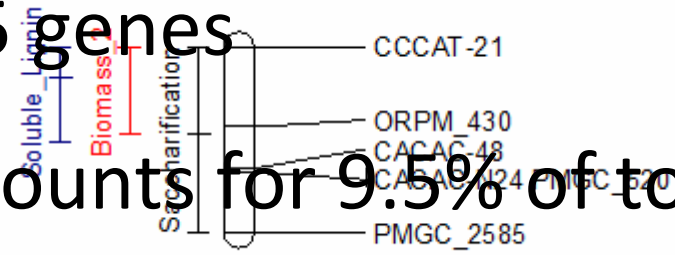




- Densely mapped

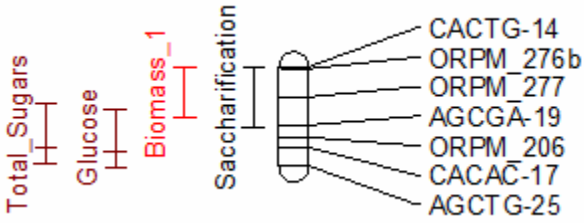
- Comparatively small - consists of 75 genes

- Accounts for 9.5% of total variation

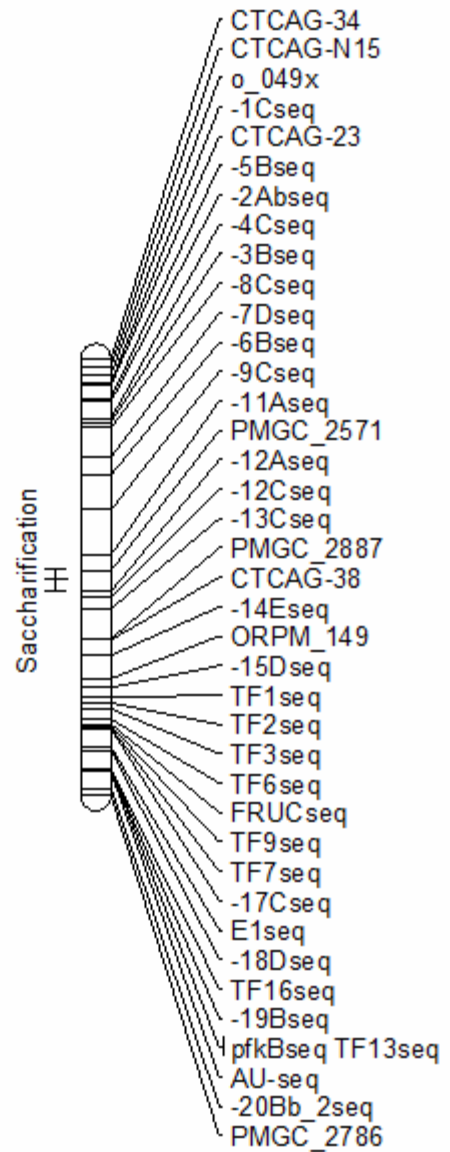


Biomass_2

Biomass_1



- In Total account for 35% of Variation



energy poplar Workpackage 3

- Genomics and breeding
- SNPs discovery and HT genotyping based on SNPs – areas of the genome and candidate genes of interest
- Select new trees for crossing with superior alleles



Workpackage 4

- Life Cycle Analysis
- Sustainability and ecosystem services
- Water use efficiency



Summary



- Poplar has potential, globally as a bioenergy feedstock crop
- Genomic approaches are beginning to unravel complex traits such as biomass, response to drought
- New Bioenergy Centres in the USA will invest massively in *Populus*. Europe should aim to remain competitive