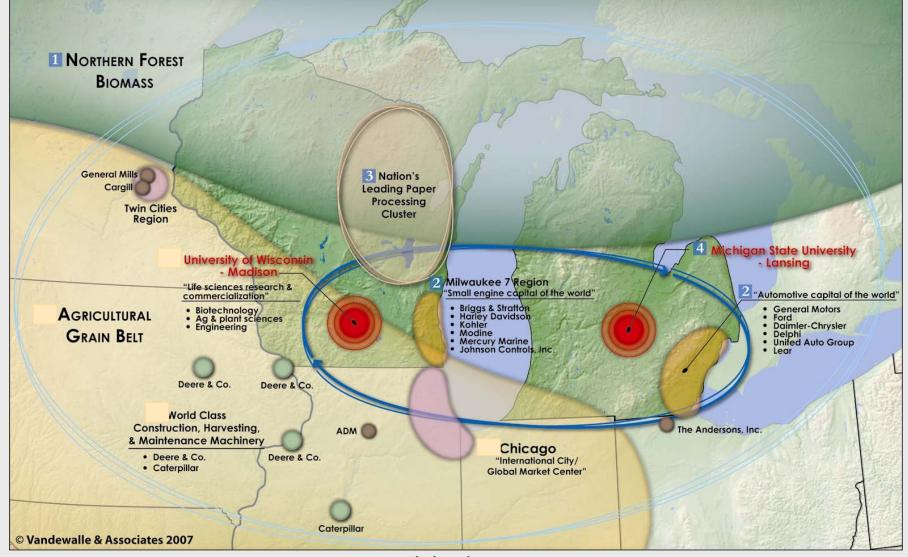


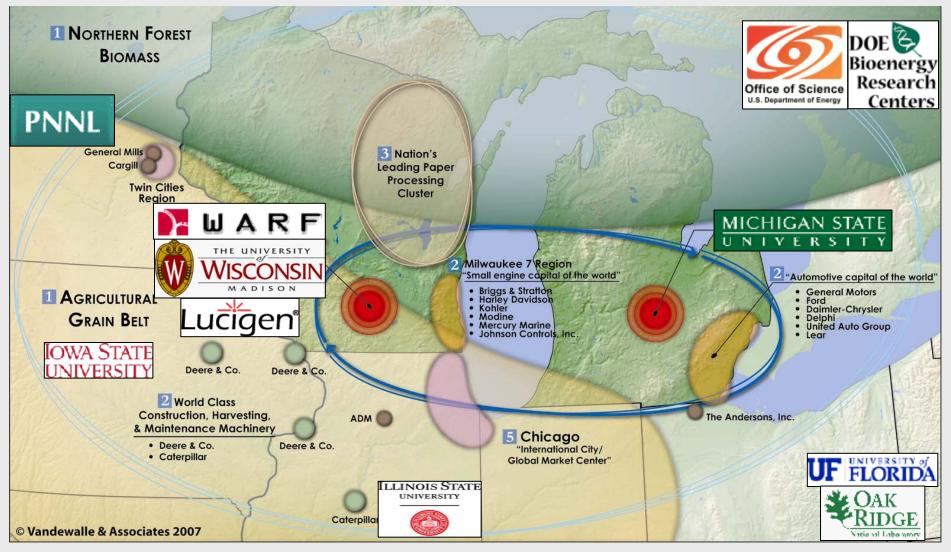
Tim Donohue (UW-Madison) BER Advisory Committee November 29, 2007

http://www.greatlakesbioenergy.org

Vision: Plugging the scientific, agricultural, economic & technological excellence in the Great Lakes Basin into the energy grid



www.greatlakesbioenergy.org



www.greatlakesbioenergy.org

Mission: perform fundamental research aimed at removing bottlenecks in the biomass to bioenergy pipeline

CENTER

RESEARC

PROCESSING

BIOMASS

Integrated research goals

- 1. Improve plant biomass
- 2. Improve biomass processing
- 3. Improve conversion of plant biomass to liquid & other fuels
- Evaluate sustainability of biomass to biofuels pipeline
- 5. Develop & use enabling genome-based technologies

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Education & Outreach
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Vandewalle & Associates 2007

JREAT

HES BIOENERGY

SUSTAINABLE

SYSTEMS

BIOMASS

ENABLING

TECHNOLOGIES

BIOMASS

GLBRC Thrust 1: Improved Plant Biomass

Objectives: obtain mechanistic understanding of biochemical & regulatory pathways needed to divert plant carbon into:

- > More digestible cell wall polymers
- Starch, fructans and other digestible carbohydrates
- Plant-derived oils

>50% of Thrust One budget devoted to creation of energy crops with novel forms of biomass storage

Discovery of cell wall biosynthetic genes & regulators

Approach: expression profiling of developing seeds

Gene Discovery -Goal: > 1 million ESTs from each system (with JGI)

Fenugreek







Psyllium

Nasturtium





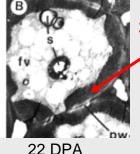
Mucilaginous layer 60% Xylan

Endosperm 90% Mannan





20 DPA

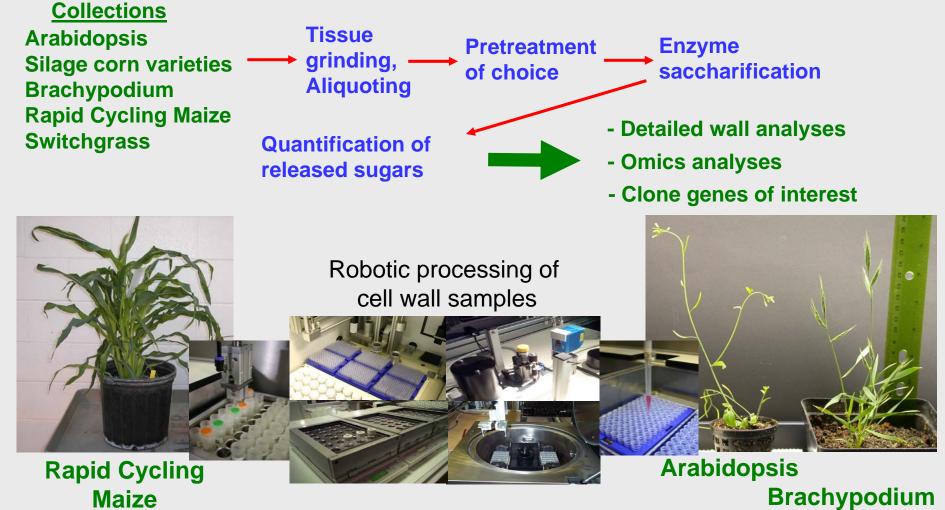


Cotyledon 30% Xyloglucan

Genes affecting cell wall composition & digestibility

Approach: High throughput mutant analysis (with Thrust 2)

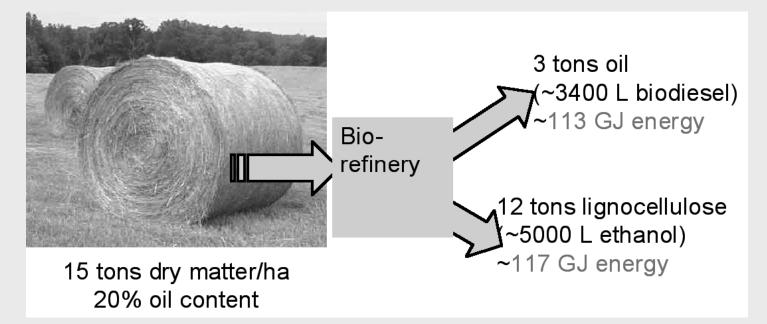
Cell wall analytical platform: Fully automated, 400 samples/day (WT & mutants)



Oil accumulation in biomass plants

Plant oils are the most energy rich, natural form of abundant carbon

Use as energy source does not require fermentation or distillation



Biomass crop with 20% oil content will almost double energy content available for liquid fuel

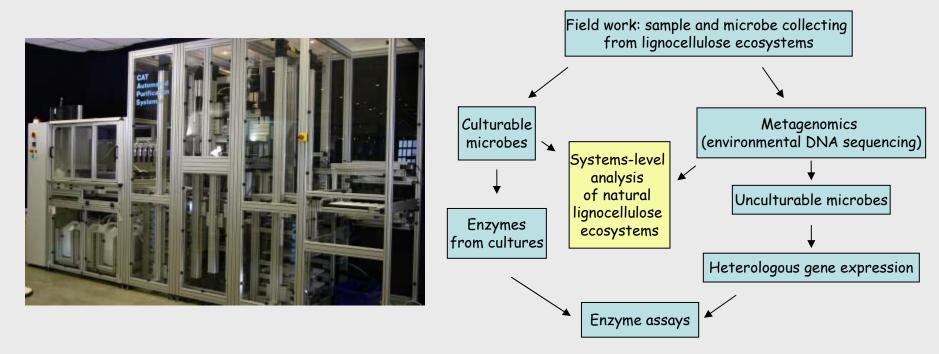
Similar approaches to dissect metabolic and regulatory circuits that control oil biosynthesis

Objective: improve conversion of plant cell walls into fermentable or chemically-convertible materials by

- > analyzing a range of plant material & pretreatment conditions
- > discovery & application of improved enzymes

Enhanced digestibility

Bioprospecting



Goals:

Enhanced biomass conversion potential

Combinatorial, high throughput, screen of candidate materials with various pretreatments for improved cell wall digestion (with Thrust 1)

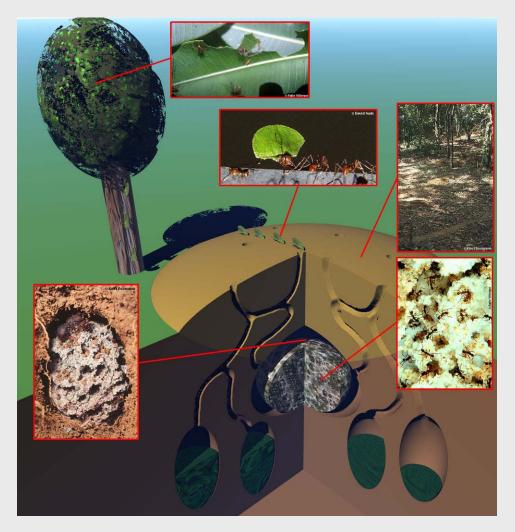
Understand pretreatment chemistries

Identify & quantify relevant small molecules produced by pretreatment chemistries (with Thrust 5)

Screen products for fermentation (microbes) or conversion (chemistry)

- Assess fermentatability (with Thrust 3)
- Assess use in catalytic conversion (with Thrust 3)

Goal: Discover/improve deconstruction enzymes



 Bioprospecting in cellulosedegrading ecosystems (with JGI)

High-throughput screening of genetic material from promising niches

Engineer improved enzymes(with Thrust 5)

Goals:

- hydrolyze biomass
- Develop suites of enzymes tailored for specific pretreated biomass samples (with Thrust 5) \succ

Decrease enzyme cost

- Test expression of new cellulolytic enzymes in plants (with \succ Thrust 1)
- Improve expression of cellulolytic enzymes in maize, alfala and other plants (with Thrust 1) \succ

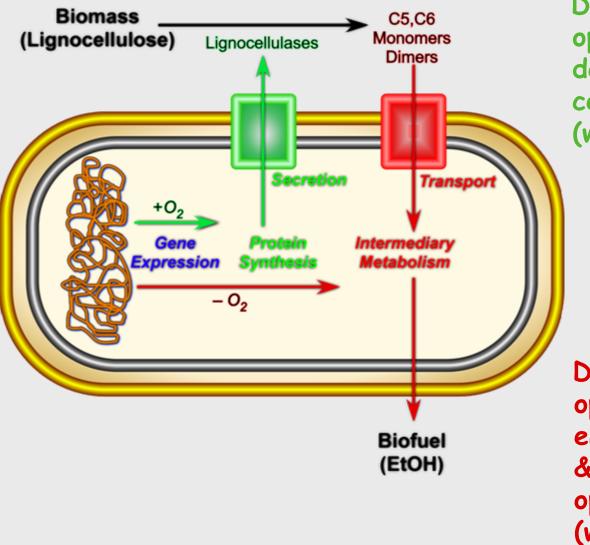
Conversion of biomass into energy products: improve methods for converting plant biomass into materials that can replace fossil fuels

- > ethanol
- > hydrogen
- chemical feedstocks

Polysaccharides Monosaccharides Polyols Engineering Analysis Polyglucan Microbe 1a. Stochastic strain alteration 2. Growth condition variation fructose sorbitol selection Biomass source, pH. Variant creation for detailed glucose hydrolysis hydrogenation Temperature, Solutes, Metagenomics, Targeted gene analysis Small molecule effectors mutagenesis, Global network xylitol xylose polyfructan mutagenesis (gTME) 3. Comprehensive molecular Screen/selection assay ("multi-omics") 4. HTMPS, Conventional selection, 1. Transcriptomics, ChIP-chip, Competitive evolution dehydration aqueous phase reforming 3. Proteomics, Lipidomics, (APR) dehvdration/ Metabolomics, Fluxomics 1b. Rational strain alteration hydrogenation H₂ and CO₂ (APD/H) Metabolic engineering 4. Data-driven computational $C_3 - C_6$ alkanes 2. Expression engineering models of microbial state Target catalytic Genome optimization Gene expression networks selected for gasification hydrogenation 5. Comparative genomics Metabolic flux models hydrodeoxygenation aldol condensation reengineering Protein engineering APD/H Membrane transport models H₂ and CO Protein interaction predictions 6. 1c. Microbial experimental evolution FT synthesis Liquid Alkanes P-series Fuel

Biological & chemical conversion platforms

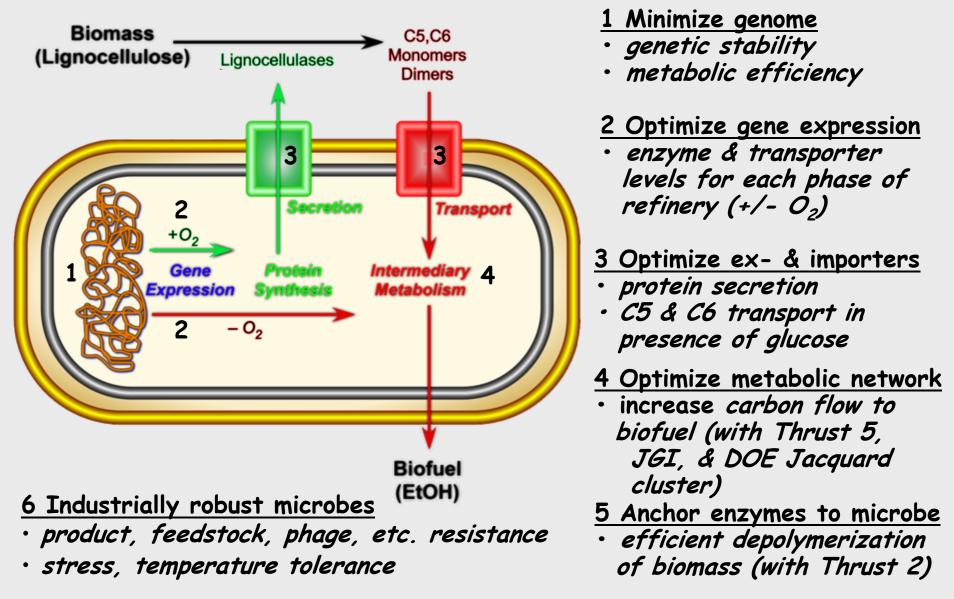
Long-Term Strategy for Consolidated Ethanologens



During aerobic growth, optimize production of desired extracellular cellulases & hemicellulases (with Thrusts 2 & 5)

During anaerobic phase, optimize production of enzymes, transporters, & pathways to optimize ethanologenesis (with Thrust 5)

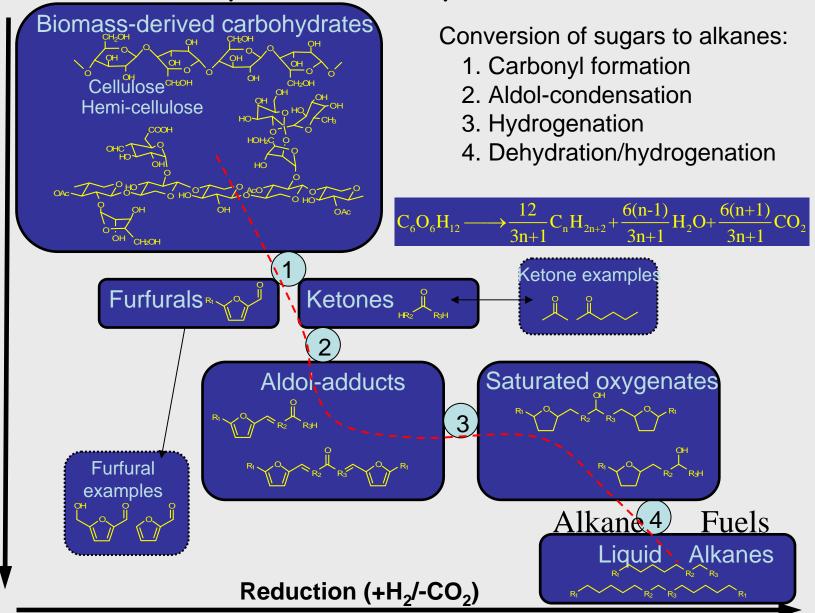
GLBRC Targets for Removal of Bottlenecks in Ethanologenesis



Reiterative Directed Microbial Evolution (REDIME) to Obtain Improved Bioenergy Microbes (apply to EtOH, H₂, CO₂ sequestration, other fuels & feedstocks)

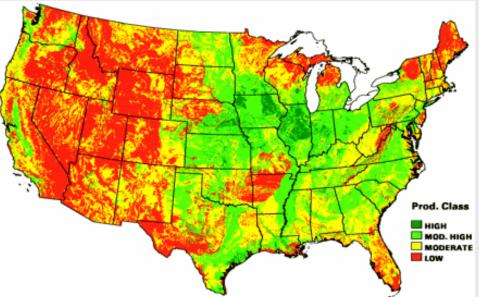
Engineering	\longrightarrow	Analysis
1a. Stochastic strain alteration	Microbe selection for detailed analysis	2. Growth condition variation
 Variant creation Metagenomics, Targeted gene mutagenesis, Global network mutagenesis (gTME) Screen/selection HTMPS, Conventional selection, Competitive evolution 		Biomass source, pH, Temperature, Solutes, Small molecule effectors
		3. Comprehensive molecular assay ("multi-omics")
		Transcriptomics, ChIP-chip, Proteomics, Lipidomics,
1b. Rational strain alteration		Metabolomics, Fluxomics
Metabolic engineering Expression engineering Genome optimization Comparative genomics Protein engineering	Target	4. Data-driven computational models of microbial state
	selected for reengineering	Gene expression networks Metabolic flux models Membrane transport models
1c. Microbial experimental evolution		Protein interaction predictions

Process & Discovery Chemical Catalysis for Biofuels Production



Dehydration (-H₂O)

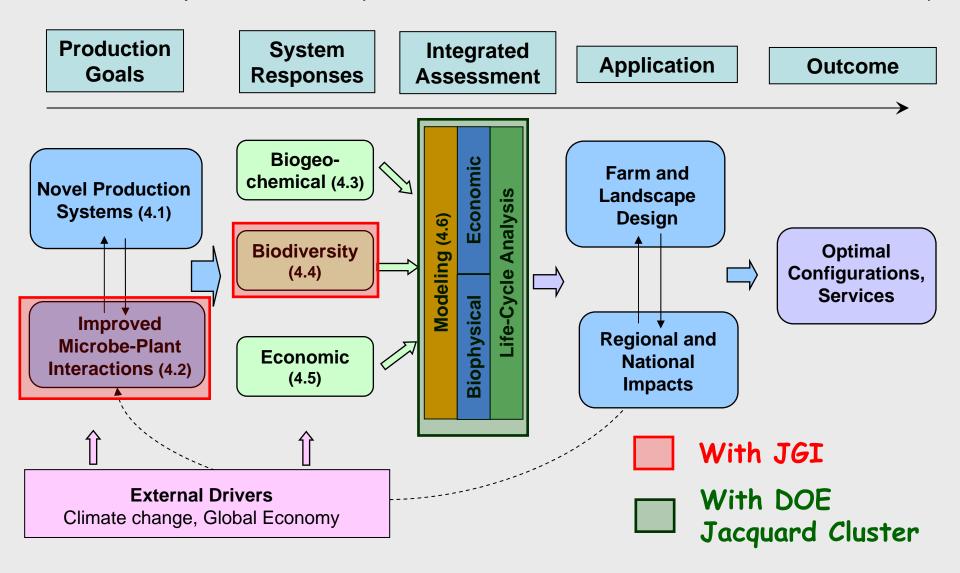
Development of a sustainable bioenergy economy: support the biomass-to-bioenergy pipeline by developing ecological, agricultural & life cycle practices that are economically viable & environmentally responsive



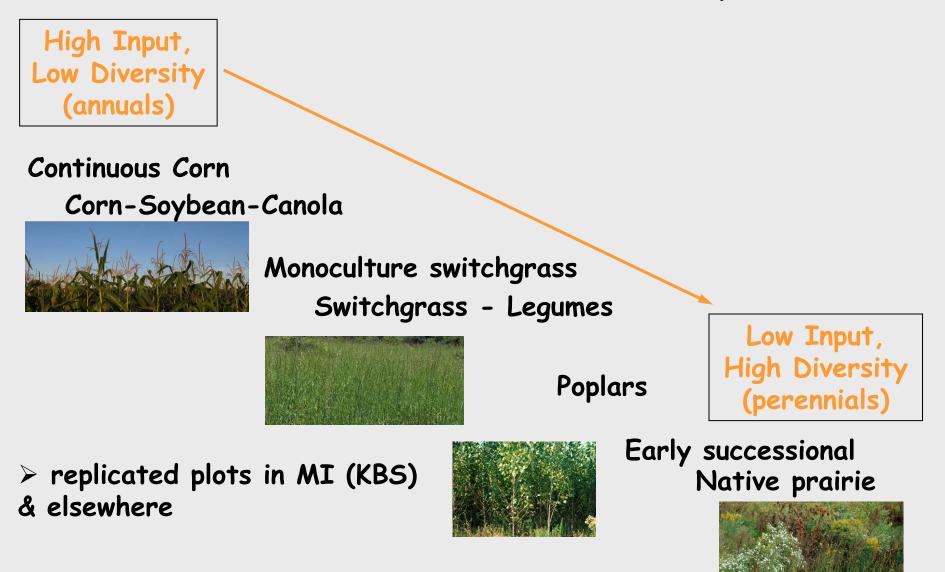
Overcome bottlenecks in agricultural, industrial, & behavioral systems to improve

- carbon neutrality and net greenhouse gas mitigation across the entire biofuel life cycle at multiple scales
- ecosystem services in biofuel landscapes (e.g. water quality, biodiversity, pest suppression)

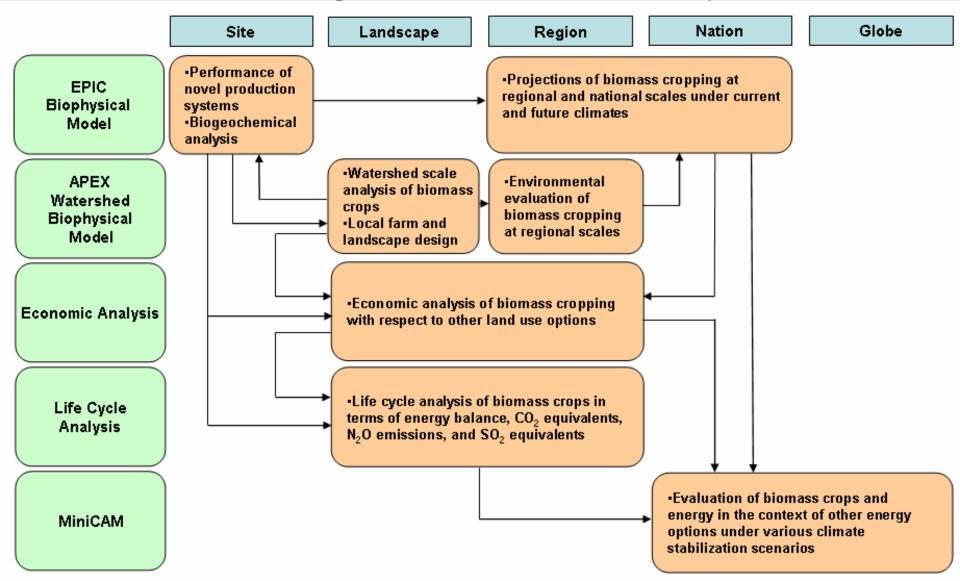
Objective: Determine elements of **integrated** biofuel production systems that can be **optimized** to improve **environmental & economic** sustainability



Goal: Predict Behavior of Novel Production Systems



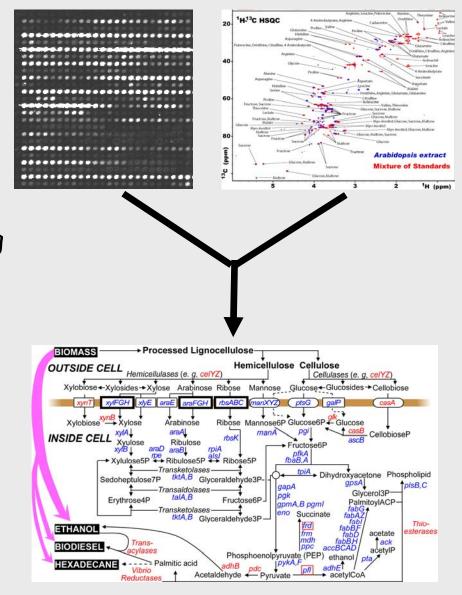
Multi-Scale, Modeling Framework (with DOE Jacquard cluster)



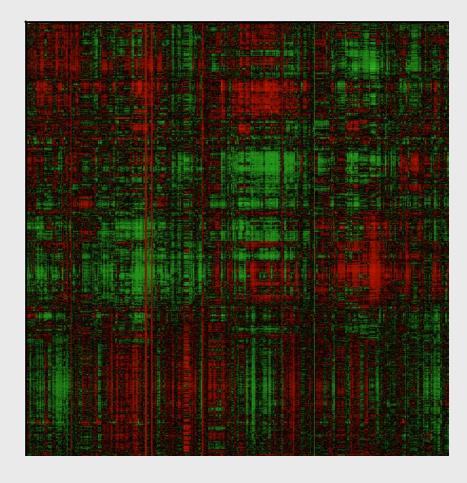
Objective: provide cutting edge genome-based technologies that enable the innovation, discoveries & creative solutions needed to remove biofuels bottlenecks

- high-throughput screens
- global analyses
- metabolic flux analysis
- protein & metabolic engineering
- computational modeling

Genome-enabled analyses



Mapping Biofuels Regulatory Networks



Cluster sets of co-regulated genes to identify network structure

High resolution analysis of biofuels transcription factor binding sites (with JGI)

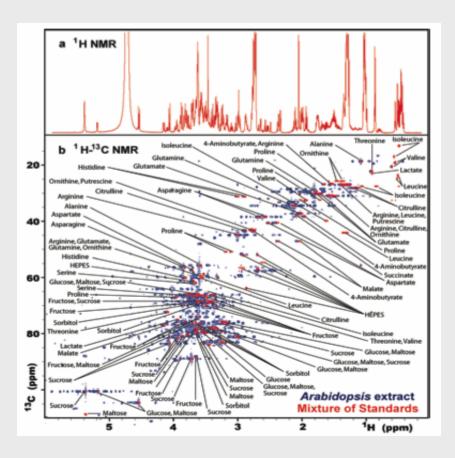
Predictive models for bioenergy regulators and regulatees (with DOE Jacquard cluster)

Bioenergy Protein Blueprints



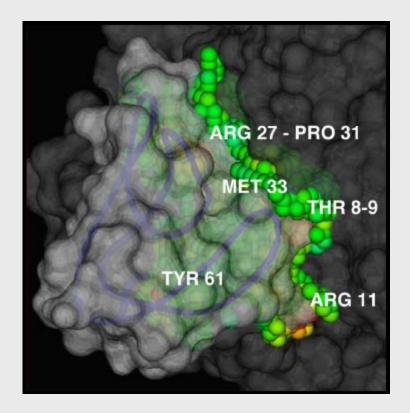
- Isotope-assisted protein abundance measurements
- Localization of proteins in subor extra-cellular fractions
- Monitor and map sites of covalent modifications that impact biomass or fuel production

Bioenergy Metabolites



- MS & NMR pipeline to quantify critical biofuel or biomass intermediates
- Discovery platform for new biofuel metabolites
- Isotope assisted flux balance analysis
- Predictive computational models for diverting carbon skeletons into biomass or biofuels pathways (with DOE Jacquard cluster)

Protein Expression and Engineering



- High throughput pipeline for protein expression and functional analysis
- Multiplex screening for new and improved biofuel enzymes
- Computational predictions of active sites, protein-protein interfaces, or protein stability to engineer new biomass pathways or improve biofuel enzymes (with BACTER & DOE Jacquard cluster)

Objectives: Develop a coordinated bioenergy education and outreach program

".....solve today's bioenergy bottlenecks while training the bioenergy leaders of tomorrow...."

GLBRC Education and Outreach

Goal: Coordinated bioenergy education and outreach program

> Workshops & educational modules for K - 12, teachers or public on carbon chemistry, sustainability, biodiversity

Public talks/workshops/communications program

> Develop materials to inform farmers, municipalities and other **members** of the community about bioenergy

> Exhibits on biomass & bioenergy (aka- "Bioenergy Discovery Center")

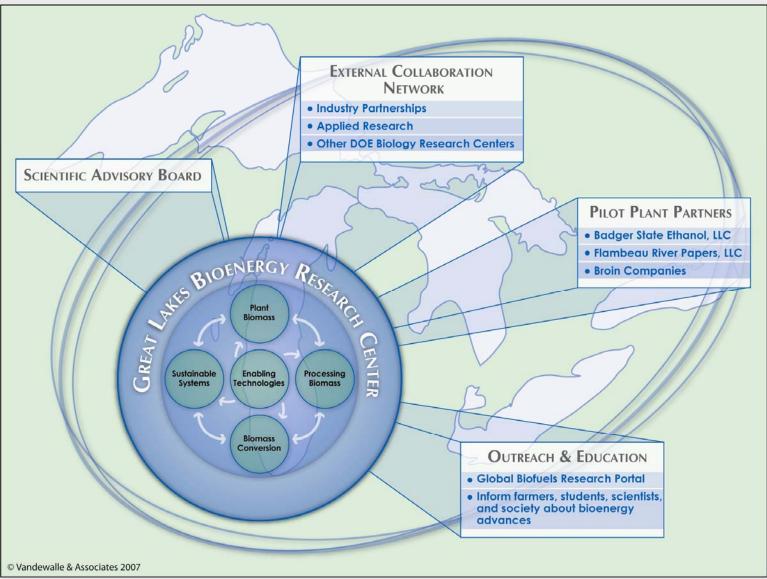
Bioenergy seminar programs and topics in biology, engineering & computational courses and labs (partnering with BACTER and others)

Summer research programs for undergraduates from other campuses; including major URM institutions

> Attract graduate students from highly rated programs

Integration of GLBRC Functions

Public, private & institutional partnerships to function as a worldwide research Center of Excellence



Plugging the scientific, agricultural, economic & technological excellence in the Great Lakes Basin into the energy grid

