

# DOE Bioenergy Technologies Office (BETO) 2019 Project Peer Review

## Engineering Thermophiles to Produce Drop-in Fuels from Syngas

**March 4-8, 2019  
Bioconversion Technologies**

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# Project Goals, Outcomes, Relevance

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**GOAL:** Develop recombinant thermophilic microbial strains capable of metabolizing syngas ( $\text{CO}$ ,  $\text{CO}_2$ ,  $\text{H}_2$ ) into monoterpenes to enable industrial production of fuels and solvents

## OUTCOMES:

- Identified and characterized multiple terpene synthase enzymes
- Developed analytical methods for monoterpene products
- Identified new thermophile chassis strain and verified  $\text{CO}$  consumption
- Developed vectors and achieved transformation of thermophilic carboxydophilic bacteria with heterologous genes
- Demonstrated monoterpene synthesis from syngas in a mesophilic host

## RELEVANCE:

- Syngas feedstocks are relatively inexpensive and diverse, thermophiles have several unique advantages, and end products have value as fuel and solvent applications among others.

# Quad Chart Overview

**Start:** October 1, 2015

**End:** September 30, 2017

**Completion:** 86%

**Technical Target:** Conversion

R&D

|                    | Total costs | FY16       | FY17 to End |
|--------------------|-------------|------------|-------------|
| DOE Funding        |             | 815,138    | 784,184     |
| Project cost share |             | 208,310    | 191,522     |
| Kiverdi<br>NREL    |             | 69%<br>31% |             |

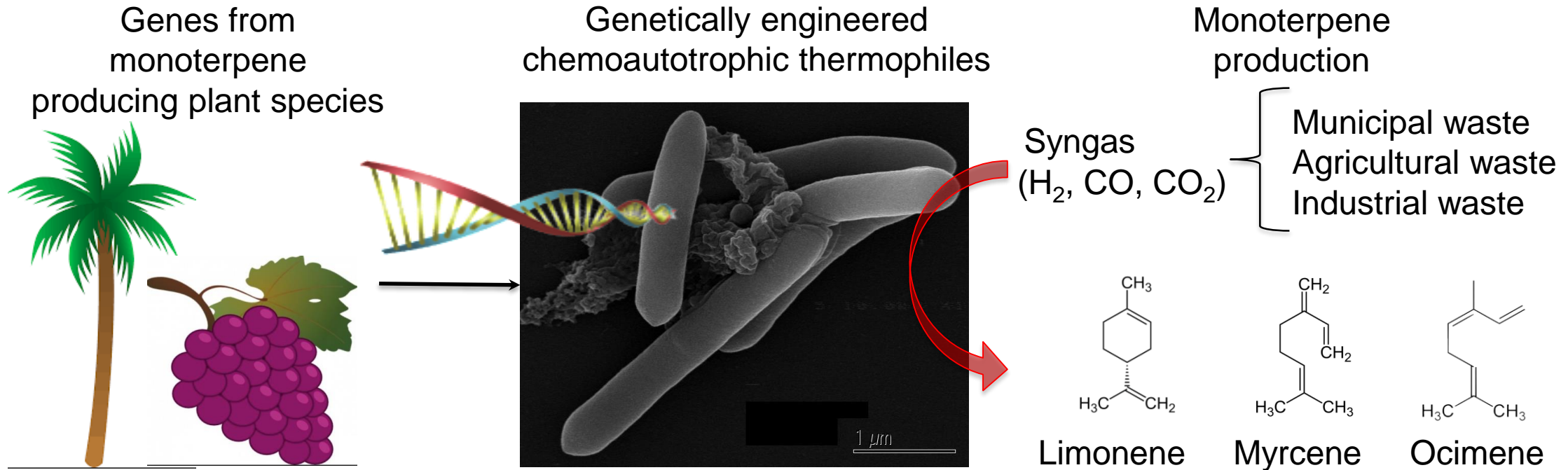
## Barriers Addressed

- Heterologous genetic transformation of a chemoautotrophic thermophile
- Thermostability of monoterpene synthases
- Syngas fermentation by engineered thermophilic strains

**Objective:** Develop microbial biocatalysts for industrial syngas fermentation to fuels and chemicals

**End of project goal:** A thermophilic chemoautotroph using syngas engineered to produce a monoterpene

# Project Overview



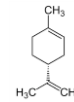
# Project Relevance

## Inexpensive and flexible feedstocks

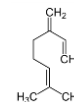


### Coupled Syngas and Bioprocess

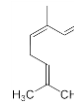
Limonene



Myrcene



Ocimene



## Applications for Monoterpenes

### Industrial Solvents

- Diluent
- Oil sand solvent
- Refining agent for crude petroleum
- Asphalt grading
- Industrial cleaner (e.g., marine vessels, concrete cleaners, parts washer)

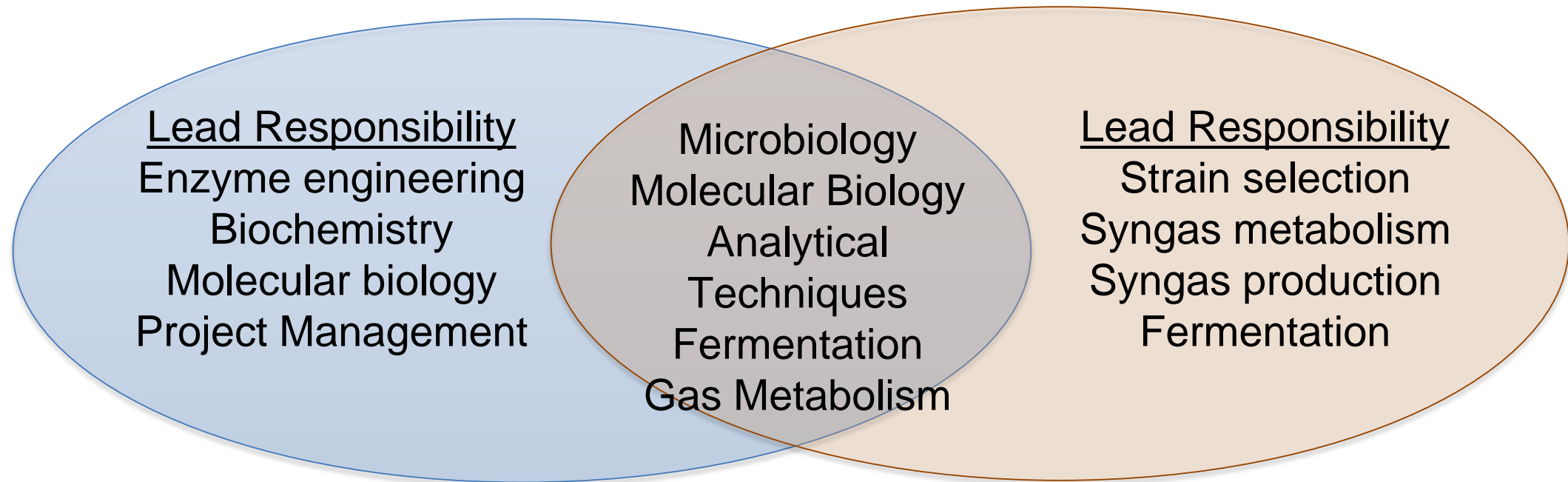
### Fuels

- **JP-10 jet fuel substitute (dimer)**
- **Bio-gasoline**
- **Gasoline or diesel additive**
- **P-menthane (component of JP-4 / Jet B)**

## Additional Advantages for Bio-derived Monoterpenes:

- Non-carcinogenic and non-toxic hydrocarbon
- High energy density (37.8 MJ/L, limonene)
- Bio-degradable
- Multiple products with the same process and equipment
- Consumer and home product applications (cleaners and insecticides)

# Project Management



Highly cooperative team science approach. Frequent meetings and offline communication.

# Technical Approach

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## **Integration of five objectives:**

1. Identification and validation of chemoautotrophic host strains
2. Development of genetic tools and molecular biology approaches
3. Identification and recombinant expression of thermostable monoterpene synthesis enzymes
4. Development of reliable analytic techniques to quantify monenterpene molecules
5. Fermentation methods and solvent overlays optimized for monoterpene production



# Technical Tools

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## **Analytical methods**

- Detection and quantitation of monoterpene
- Detection of substrate molecules and side products

## **Microbiology**

- Identification of new chassis strain (CO consuming thermophile)
- Routine manipulation of chassis strain(s)
- Liquid culture, cryogenics, plating, antibiotic selection

## **Biochemistry**

- Biochemical assays to test enzyme activity, solubility, and performance
- Biochemical methods to detect enzyme substrates and precursor enzymes

## **Molecular biology**

- Development of vectors to manipulate thermophilic chassis strain(s)
- Identified promoter to drive expression and thermo-tolerant selection method



# Technical Achievements

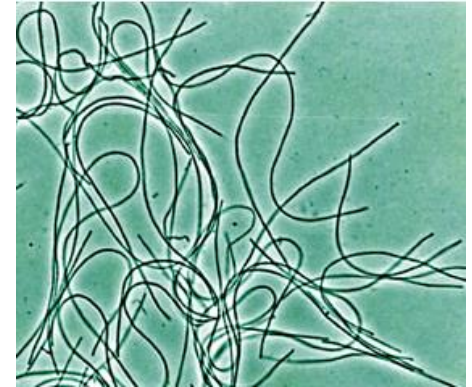
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- Research and management teams demonstrated the feasibility of the syngas bioprocess
- 90% of the work developing the recombinant chemoautotrophic thermophile was completed; monoterpene synthesis from CO was not achieved
- Monoterpene production from syngas was demonstrated in a mesophilic chemoautotroph; titers reached 25% of original target
- Genetic tools and analytical methods were developed that support ongoing biocatalyst R&D
- Data were recorded for future technoeconomic process analysis

# Chassis Strain Selection

## Originally Proposed Strain: *Chlororflexus aurantiacus*

- Thermophilic green non-sulfur bacteria (40-50°C)
- Photosynthetic (anaerobically), but capable of aerobic growth in the dark
- Grows autotrophically on H<sub>2</sub>/CO<sub>2</sub>, and heterotrophically (organic acids, sugars)
- Putative CO dehydrogenase identified
- High natural terpenoid pathway flux (photosynthetic pigments)



**Early evaluation revealed *Chlororflexus aurantiacus* to be unsuitable host for project goals.**

- **No/poor growth under dark aerobic conditions regardless of carbon source**
- **No evidence of any CO/Syngas uptake under any growth condition**

- Effort was targeted to identification of a new host strain candidate
- This was not part of the original scope but clearly essential for project goals

# Candidate Strain Screening

12+ potential strains identified, 9 strains underwent extensive testing on criteria.

## Strain evaluation criteria:

- 1) CO/Syngas consumption
- 2) Thermal range of growth
- 3) Available genome sequence
- 4) Genetic tractability
- 5) Metabolic properties (e.g. terpene flux)

| Microbe                                                     | CO uptake | Aerobic/ Anaerobic | Autotrophic growth on syngas components (with O <sub>2</sub> for aerobic strains) |                                 |                                     |                    | Genetic tractability | Genome Sequence | High flux to terpenes | Growth Temp (°C): Optimum (Min) |
|-------------------------------------------------------------|-----------|--------------------|-----------------------------------------------------------------------------------|---------------------------------|-------------------------------------|--------------------|----------------------|-----------------|-----------------------|---------------------------------|
|                                                             |           |                    | CO                                                                                | H <sub>2</sub> /CO <sub>2</sub> | H <sub>2</sub> /CO <sub>2</sub> /CO | H <sub>2</sub> /CO |                      |                 |                       |                                 |
| <i>Chloroflexus aurantiacus</i>                             | -         | Both               | -                                                                                 | +                               | ND                                  | ND                 | ?                    | +               | +                     | 48 (40)                         |
| <i>Rubrivivax gelatinosus</i> CBS                           | +         | Both               | ++<br>+                                                                           | +++                             | +++                                 | +++                | +                    | +               | +                     | 35                              |
| <i>Cupriavidus necator</i>                                  | -         | Both               | -                                                                                 | +++                             | ND                                  | ND                 | +                    | +               | -                     | 30                              |
| <i>Cupriavidus metallidurans</i>                            | -         | Both               | -                                                                                 | +++                             | ND                                  | ND                 | (+)                  | +               | -                     | 30                              |
| <i>Thermomicrobium roseum</i>                               | +         | Aerobic            | -                                                                                 | -                               | -                                   | ND                 | ?                    | +               | +                     | 70 (55)                         |
| <i>Moorella thermoacetica</i>                               | +         | Anaerobic          | ND                                                                                | ND                              | +                                   | ND                 | (+)                  | +               | +*                    | 55 (45)                         |
| <i>Hydrogenobacter thermophilus</i>                         | ND        | Aerobic            | ND                                                                                | +                               | ND                                  | ND                 | ?                    | +               | ?                     | 70                              |
| <i>Carboxydotherrmus hydrogenoformans</i>                   | (+)       | Anaerobic          | ND                                                                                | ND                              | ND                                  | ND                 | ?                    | +               | +*                    | 65 (40)                         |
| <i>Carboxydocella thermautotrophica</i>                     | (+)       | Anaerobic          | (+)                                                                               | ND                              | ND                                  | ND                 | ?                    | -               | ND                    | 58 (40)                         |
| <i>Caldanaerobacter subterraneus</i> sp. <i>yonseiensis</i> | +         | Anaerobic          | ND                                                                                | ND                              | ND                                  | +                  | ?                    | +               | +*                    | 75 (50)                         |
| <i>Hydrogenibacillus schlegelii</i>                         | +         | Aerobic            | ND                                                                                | ND                              | +                                   | ND                 | ?                    | +               | -*                    | 65 (42)                         |

# Identification of New Chassis Candidates

Variables tested during growth optimization experiments:

- 1) Autotrophic vs. mixotrophic growth
- 2) Media composition
- 3) Uptake of Syngas components (CO, H<sub>2</sub>, CO<sub>2</sub>)
- 4) Culture: headspace ratio (optimized gas mixing)
- 5) Temperature (minimal/maximal, optimal)

35+ growth experiments  
for top 6 candidate strains,  
each lasting 3-5 days

Three candidate thermophilic strains with sequenced genomes were identified as project chassis strains

| Microbe                                                          | Aerobic/<br>Anaerobic | CO<br>uptake | Growth and<br>utilization of<br>synthetic syngas | Genetic<br>tractability | Genome<br>Sequenced | High flux to<br>terpenes | Optimum growth<br>temp °C<br>(Minimum) |
|------------------------------------------------------------------|-----------------------|--------------|--------------------------------------------------|-------------------------|---------------------|--------------------------|----------------------------------------|
| <i>Moorella thermoacetica</i>                                    | Anaerobic             | +            | +                                                | (+)                     | +                   | +*                       | 55 (45)                                |
| <i>Thermomicrobium roseum</i>                                    | Aerobic               | +            | +                                                | ?                       | +                   | +                        | 70 (55)                                |
| <i>Caldanaerobacter subterraneus</i> sp.<br><i>tengcongensis</i> | Anaerobic             | +            | +                                                | (+)                     | +                   | +*                       | 75 (50)                                |

Note: Two mesophilic backup strains also identified as contingency strains for this project.

(+) reported in literature; \* based on annotated pathways in genomes

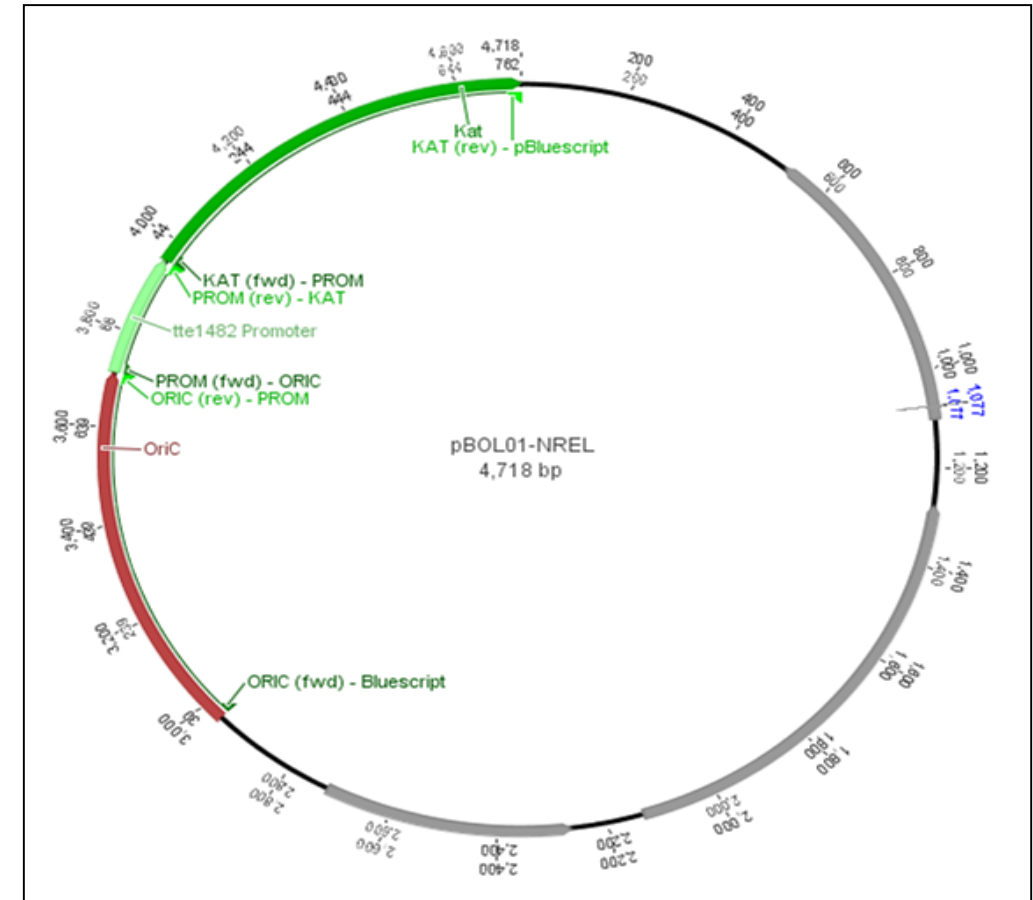
# Thermophile Genetic Tools: *C.s. tengcongensis*

## Expression Vector Development

- *tengcongensis* expression vectors built
- Amylase promoters driving:
  - myrcene synthase (thermostable)
  - GPPS
  - Both MS and GPPS as polycistronic
  - Both MS and GPPS independently
- Phosphate acetyltransferase promoter driving:
  - myrcene synthase
  - limonene synthase
  - GPPS and LS as polycistronic

Constructs were built and used for transformation of *C.s. tengcongensis*

## Novel Vector Constructed (pBOL01-NREL)

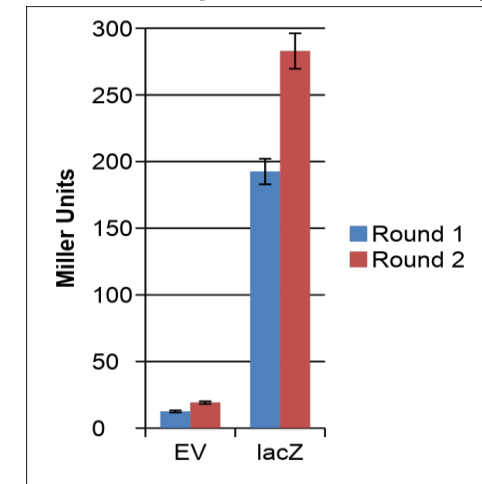


# Thermophile Genetic Tools: *C.s. tengcongensis*

Goal: Identify independently controlled promoters for expression strength and inducibility; Demonstrate expression of at least 2 genes in *C.s. tengcongensis*

| Promoter for Gene                        | Constitutive/Inducible                   | Construct |
|------------------------------------------|------------------------------------------|-----------|
| Phosphate acetyltransferase              | strong constitutive                      | ✓         |
| Glyceraldehyde-3-phosphate dehydrogenase | strong constitutive                      | ✓         |
| Glucoamylase                             | Potentially inducible (starch)           | ✓         |
| Alpha-amylase                            | Potentially inducible (starch)           | ✓         |
| Fructose-specific phosphotransferase     | Potentially inducible (fructose)         | ✓         |
| Maltose-binding protein                  | Potentially inducible (maltose)          | ✓         |
| Phosphoenolpyruvate carboxykinase        | Potentially repressible (mannoheptulose) |           |
| bgaR-PbgaL                               | lactose-inducible promoter               | ✓         |
| pRpls1                                   | strong constitutive                      |           |
| pRpls10                                  | mid constitutive                         |           |
| pRpls20                                  | weak constitutive                        |           |

## LacZ reporter assay

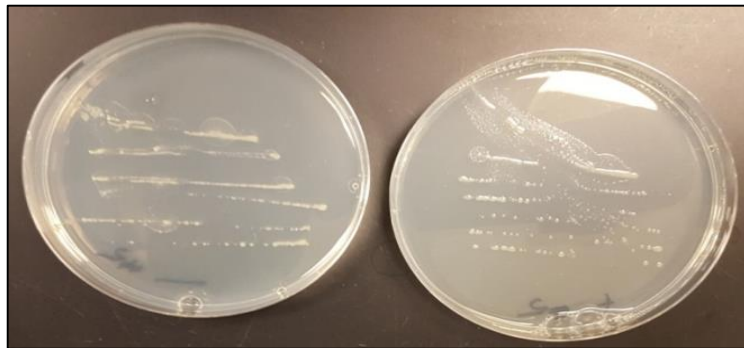


- Three constitutive promoters of varying expression strength were identified
- The range of strengths will be used to balance recombinant gene expression



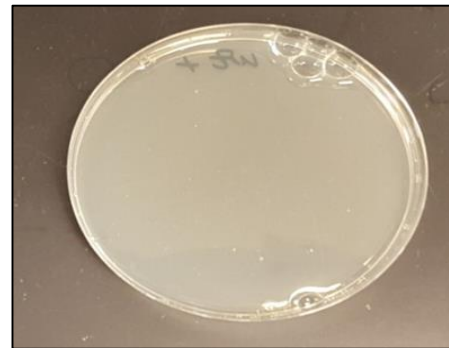
# Transformation Go/NoGo

- *Moorella thermoacetica* and *Thermomicrobium roseum* were untransformable
- **Transformed kanamycin resistance is stable in *C. tengcongensis***
- No-DNA controls remain sensitive to Kan and fail to grow (mock transformations)



Transformant  
- kanamycin

Transformant  
+ kanamycin



No-DNA control (WT)  
+ kanamycin



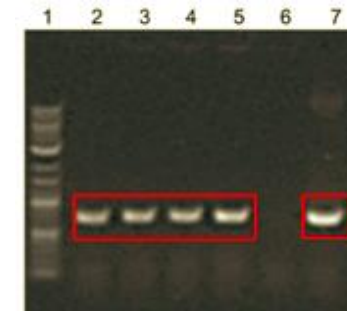
Transformants  
+ kanamycin

WT  
- kan

WT  
+ kan

## Milestones reached

- ✓ “Isolation of 1 or more transformants of the host strain”
- ✓ “functional expression of the antibiotic selection marker”
- ✓ “expression of heterologous beta-galactosidase enzyme”

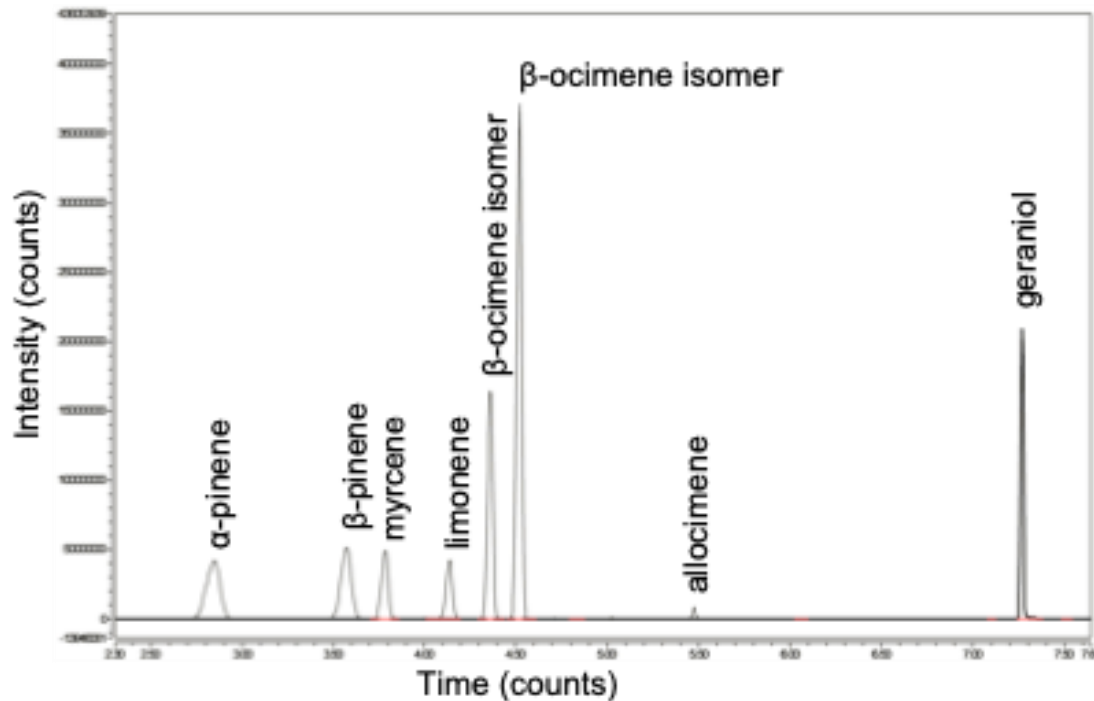


Kanamycin  
resistance  
Transformant



# Analytical Methods Development

Goal: Develop a protocol to extract, simultaneously detect and quantify monoterpenes



- Re-established standards for mercene, limonene, geraniol, pinene etc.)
- Tested overlays, dodecane, isopropyl myristate, heptane on live cultures
- Developed “universal” analytical method for monoterpenes, precursor, and overlays
- Determined IPM is leading overlay for methods and performance
- Detected bio-derived limonene from *E. coli* cultures over time with overlay
- Established reliable LOD and LOQ for all molecules of interest



# Monoterpene Synthase Candidate Screening

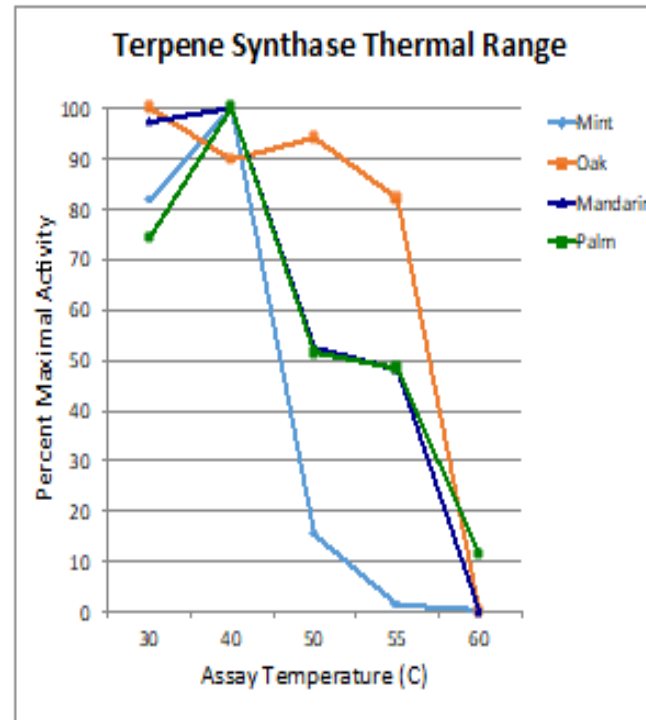
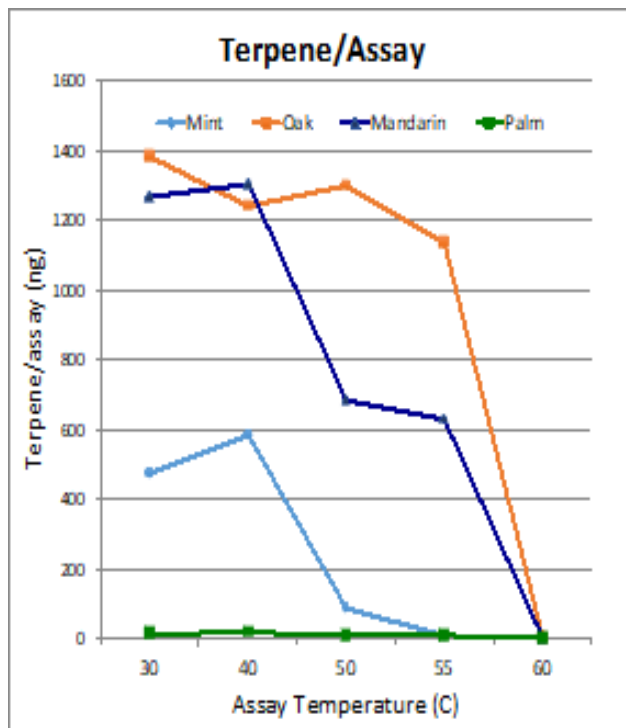
Goal: Identify at least one monoterpene synthase that would function at high temperature

- Candidate genes were screened for expression and solubility in *E. coli*.
- Genes from poplar, mint, date palm, mandarin, grape and holm oak.
- Mint, mandarin, holm oak and date palm expressed sufficiently to be detected by blotting.
- Candidates were further analyzed by *in vitro* assay for conversion of geranyl pyrophosphate (Gpp) to a monoterpene.

| TS Construct | CBB | West. | Soluble | Activity (Insol.) | Activity (solub.) | Other Products    | Product (ng)  |
|--------------|-----|-------|---------|-------------------|-------------------|-------------------|---------------|
| Mint LS*     | No  | Yes   | No      | Yes               | Yes               | -                 | <b>413</b>    |
| Mint LS      | No  | Yes   | No      | Yes               | Yes               | -                 | <b>1093</b>   |
| Date palm LS | No  | Yes   | No      | No                | Yes               | ocimene           | <b>30</b>     |
| Mand. LS     | No  | Yes   | Yes     | Yes               | Yes               | myrcene           | <b>11,173</b> |
| Oak TS       | No  | Yes   | Yes     | Yes               | Yes               | myrcene<br>pinene | <b>1564</b>   |

# Monoterpene Synthase Thermostability

- Four enzymes showing activity in E coli extracts were tested by challenging at successively higher temperature; 30, 40, 50, 55 and 60°C
- Gpp conversion to terpene products was measured in the *in vitro* assay



- Primary candidate enzyme: Holly Oak myrcene synthase
- Highest activity and thermostability
- Second candidate: Mandarin myrcene synthase

# Fermentation

Goal: Demonstrate recombinant organism growth and monoterpene production using syngas

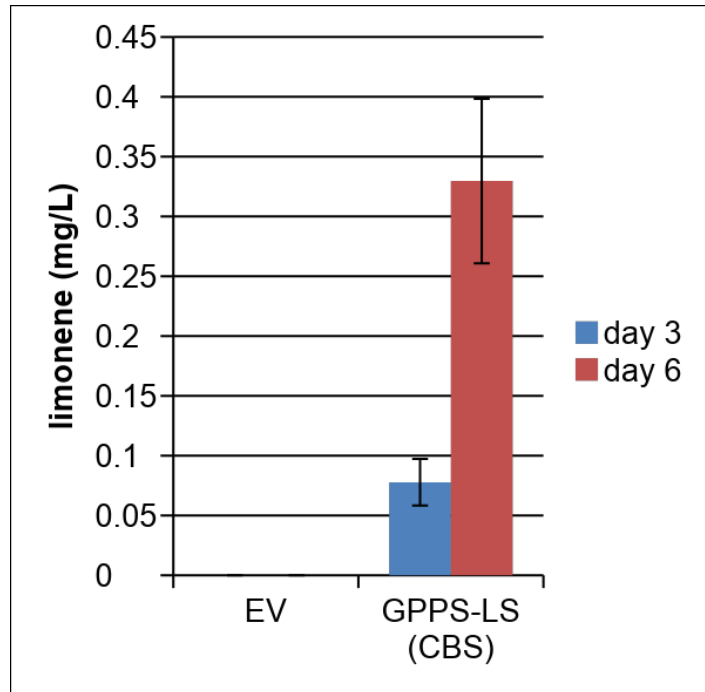
- Developed an air-lift bioreactor system (Appendix)
- Continuous delivery of syngas at controlled temperatures
- Demonstrated growth of *C.c tengcongesis* w/w/o solvent overlay
- Developed an organic solvent culture overlay to continuously extract monoterpene into isopropyl myristate



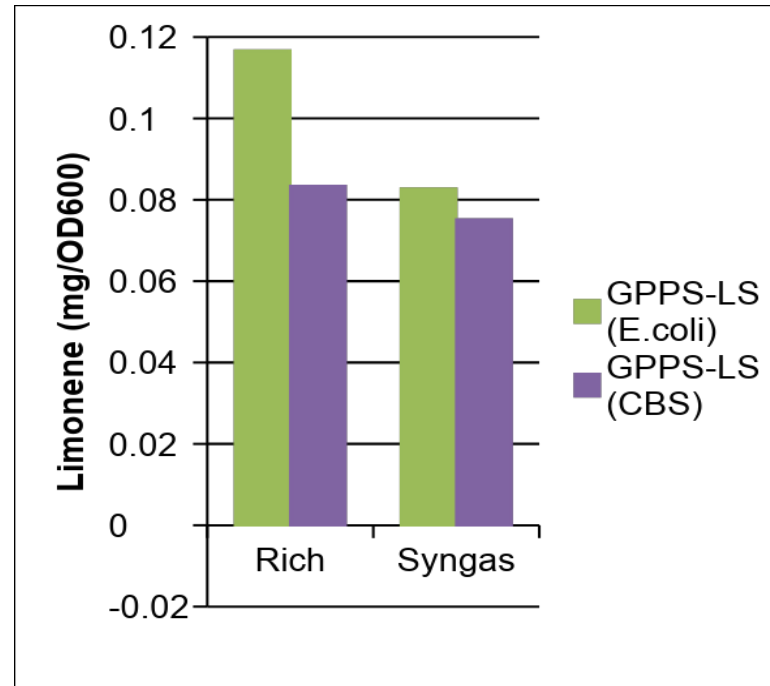
# Terpene production *in vivo*: *R. gelatinosus* CBS

Goal: Demonstrate monoterpene production in a recombinant chemoautotroph

*Rubrivivax gelatinosus* CBS strain expressing mint LS and grand fir GPP-synthase (codon optimized for CBS or *E. coli*); grown with dodecane overlay



Day 3 vs. day 6



Rich vs. minimal media (day 6)



CBS produced limonene

- 1.2mg/L on rich medium + syngas
- 0.2mg/L on min media +syngas



# Summary

| B. Start-Up                                                             |                                    | Notes                                                                                       |
|-------------------------------------------------------------------------|------------------------------------|---------------------------------------------------------------------------------------------|
| B.1 Ordering of equipment/strains/consumables                           | Completed                          |                                                                                             |
| B.2 Project syncup: Visit to NREL                                       | Completed                          |                                                                                             |
| 1.3 Materials and equipment received                                    | Completed                          |                                                                                             |
| <b>1. Host strain characterization and screening</b>                    |                                    |                                                                                             |
| 1.1 Establish base case fermentation process                            | Completed                          |                                                                                             |
| 1.2 Syngas impurity tolerance tests                                     | Completed                          |                                                                                             |
| 1.3 Test strain variant options                                         | Completed                          | Completely new host strains identified                                                      |
| 1.4 Bioinformatics                                                      | Completed                          |                                                                                             |
| 1.5 Metabolic modeling                                                  | Completed                          |                                                                                             |
| 1.6 Chloroflexus aurantiacus growth on CO ( <i>C. tencongensis</i> )    | Completed                          | Tested a dozen strains in searching for new host/platform microbes                          |
| <b>2. Development of a genetic system in host strain</b>                |                                    |                                                                                             |
| 2.1 Analysis of selection/counterselection methods                      | Completed                          |                                                                                             |
| 2.2 Test transformation methods                                         | Completed                          | Completed (Multiple strains)                                                                |
| 2.3 Apply selection protocols for clone isolation                       | Completed                          | Completed -Go/No-Go                                                                         |
| 2.4 Genetic Tool Development                                            | Completed                          |                                                                                             |
| <b>3. Terpene Synthase Development</b>                                  |                                    |                                                                                             |
| 3.1 Identify and test thermostable terpene synthases                    | Completed                          | Completed - Multiple enzymes identified                                                     |
| 3.2 Engineer terpene synthase(s)                                        | Completed                          | Determined to be unnecessary due to native thermostability                                  |
| 3.3 Demonstrate enzyme specificity                                      | Completed                          |                                                                                             |
| 3.4 Express engineered enzyme in host strain                            | Completed                          | Completed - success in <i>E. coli</i> , <i>CBS</i> and <i>C. necator</i> Not in Thermophile |
| <b>4. Biocatalyst process integration, development and optimization</b> |                                    |                                                                                             |
| 4.1 Screen primary production strains in bottles                        | Nearly completed                   | developed fermentation and overlay methods                                                  |
| 4.2 Test recombinant Task 1 strains under optimized conditions          | Fermentation and overlay optimized | final strain not available                                                                  |
| 4.3 Demonstrate capability to produce product from syngas               | Completed for CBS,                 |                                                                                             |
| <b>5. Techno-economic Analysis (TEA)</b>                                |                                    |                                                                                             |
| 5.1 Verify base case TEM assumptions for commercial success             | Began work                         |                                                                                             |
| 5.2 Complete integrated process TEA and process flow diagrams           | Incomplete                         |                                                                                             |

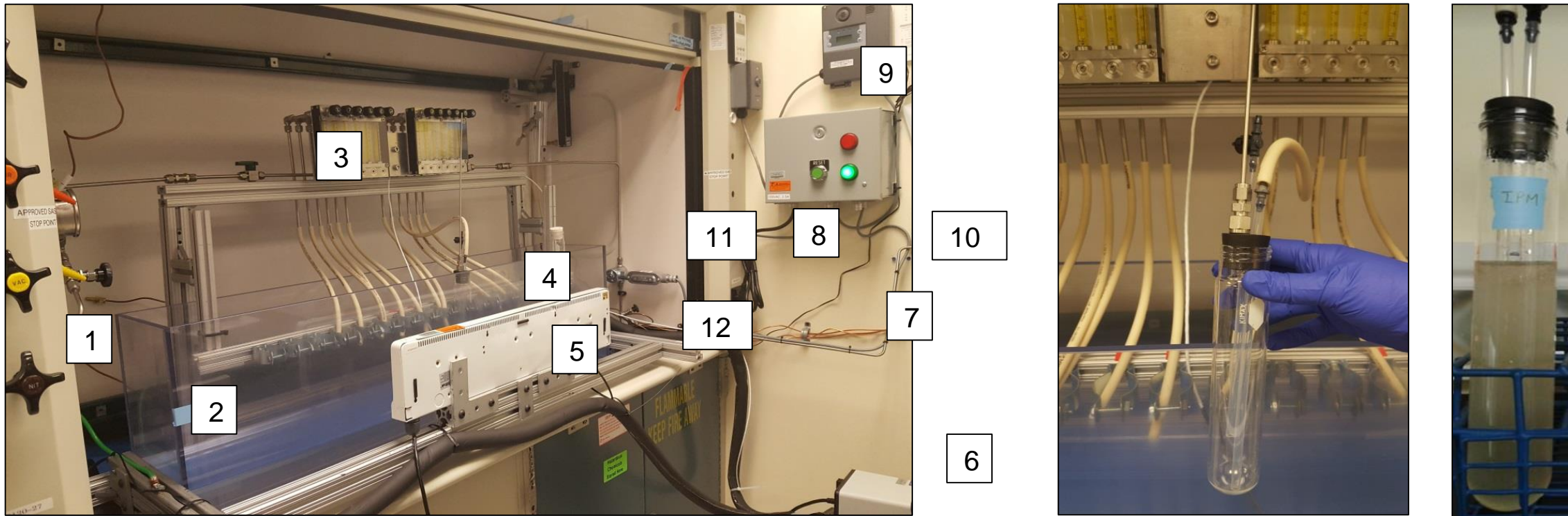
- Research and management teams demonstrated the feasibility of the syngas bioprocess
- 90% of the work developing the recombinant chemoautotrophic thermophile was completed; monoterpene synthesis was not achieved
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- Genetic tools and analytical methods were developed that support ongoing biocatalyst R&D
- Data were recorded for future technoeconomic process analysis

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# Slides Not Shown



# Air-lift Syngas Bioreactor



1) N<sub>2</sub> line (for purging samples), 2) water bath, 3) rotameters, 4) tubing and tube bioreactor, 5) halogen lights (outside hood, sash will be closed), 6) water recirculator for temperature control of water bath, 7) syngas line (stainless steel), 8) hood ventilation monitor (internal), 9) CO and H<sub>2</sub> monitor (outside hood, audible alarm), 10) relay to solenoid valve (connected to CO/H<sub>2</sub> and ventilation monitors), 11) solenoid valve (to shut off syngas if CO/H<sub>2</sub> detected outside hood or ventilation failure), 12) anti-static mat in hood (and equipment grounded/bonded, second anti-static mat will be in front of hood).

Materials chosen for compatibility with flammable gas.

# Responses to Reviewers

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# Publications, Patents

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