

# Fermentation and Electrohydrogenic Approaches to Hydrogen Production



2011 Annual Merit  
Review and  
Peer Evaluation Meeting

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Project ID #: PD038

This presentation does not contain any proprietary, confidential, or otherwise restricted information

# Overview

## Timeline

- Project start date: FY05
- Project not funded in FY06
- Project end date: 10/2011\*
- Percent complete: N/A

## Budget

- Total project funding: \$1,910K (include \$290K subcontract)
- FY10: \$230K (include \$60K subcontract)
- Funding allocated for FY11: \$400K (include \$60K subcontract)

## Barriers

- Production barriers addressed
  - H<sub>2</sub> molar yield (AR)
  - Waste acid accumulation (AS)
  - Feedstock cost (AT)

## Partners

- Dr. Bruce Logan  
Pennsylvania State University
- Drs. David Levin and Richard Sparling  
University of Manitoba, Canada  
(Genome Canada Program)

\*Project continuation and direction determined annually by DOE

# Relevance

- **Objective:** Develop direct fermentation technologies to convert renewable lignocellulosic biomass resources to H<sub>2</sub>.
  - Optimize fed-batch bioreactor (hydraulic and solid retention time) (Task 1).
  - Develop genetic tools to improve H<sub>2</sub> molar yield (Task 2).
  - Design and build a tubular type MEC and conduct performance evaluation (Task 3).
- **Relevance:** Address directly feedstock cost and H<sub>2</sub> molar yield barriers to improve technoeconomic feasibility.

Characteristics	Units	2013 Target	2011 Status
Yield of H <sub>2</sub> from glucose	Mole H <sub>2</sub> /mol glucose	4	3.2
Feedstock cost*	Cents/lb glucose	10	12

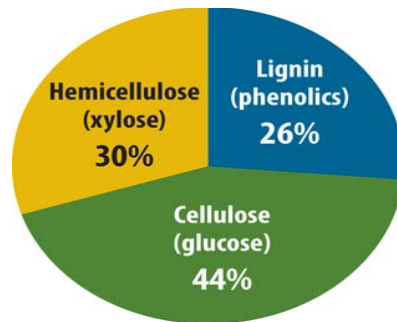
\* DOE Office of Biomass Program status and target

# Objectives/Approach/Milestone

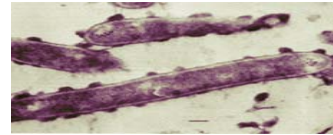
## Task 1: Bioreactor Performance

- **Objective:** Address feedstock cost and improve the performance of bioreactors for H<sub>2</sub> via fermentation of lignocellulose.
- **Approach:** Optimize bioreactor in fed-batch mode by testing parameters such as amount and frequency of cellulose feedings and acclimation of the cellulose-degrading bacterium *Clostridium thermocellum*.

### Lignocellulosic Biomass

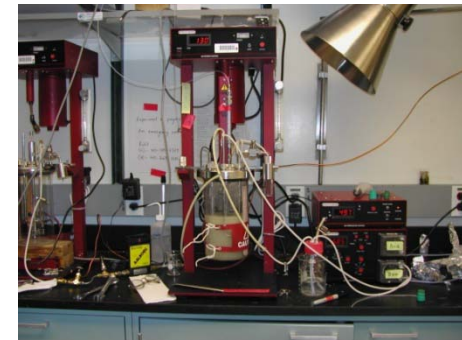


### Fermentation



*Clostridium thermocellum*

### Bioreactor Performance



	Milestone	Completion Date	Status
3.2.1.2	Determine the solid and hydraulic retention time on rates and yield of H <sub>2</sub> in <b>fed-batch</b> reactor	4/11	On track



# Task 1 – Technical Accomplishments

## Fed-Batch Bioreactor Setup

- Work in FY10 determined the effect of substrate loadings on rates and yields of H<sub>2</sub>, which guides the development of fed-batch fermentation.
- In a one-liter working volume bioreactor, we drained and replenished daily (hydraulic retention time = 24 h) with 500 mL fresh medium containing 2.5 g/L cellulose.



Acclimated microbes turned yellow

Settle, Drain, Feed



Settled microbes in yellow;  
~80% recovery

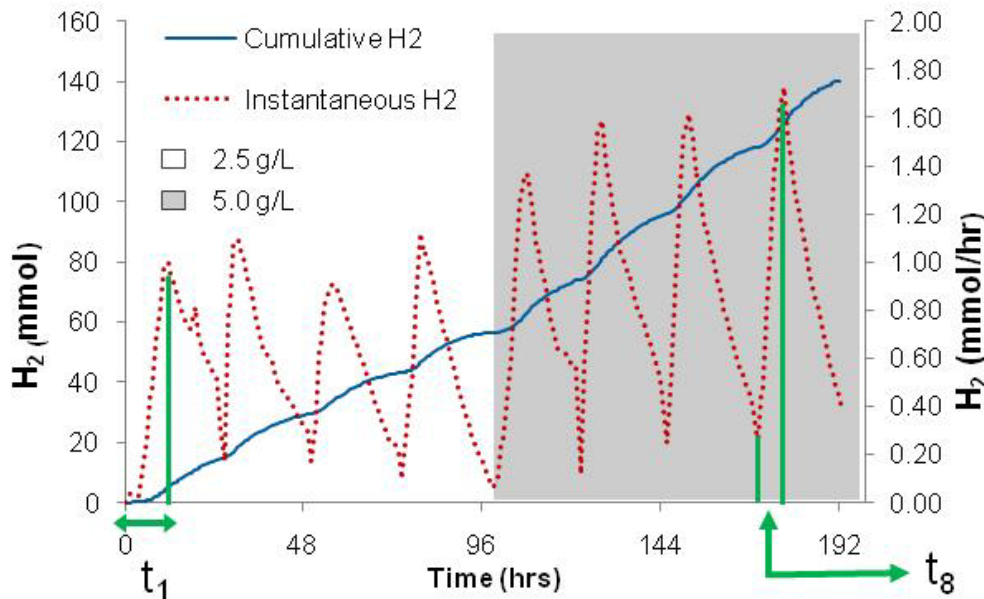
***C. thermocellum* are immobilized on cellulose, allowing the bulk of the growth medium to be replaced without diluting the fully acclimated microbes.**

# Task 1 – Technical Accomplishments

## Hydrogen Production in Fed-Batch Bioreactor

Daily feedings of 2.5 (up to 96 h) and 5.0 g/L of cellulose conducted:

- Fed-batch mode adapted *C. thermocellum* to degrade cellulose, shown by a decrease in “time to peak H<sub>2</sub> production” (faster acclimation; t<sub>1</sub> to t<sub>8</sub>).
- With faster rate of H<sub>2</sub> (> 53% increase), a smaller bioreactor can be built to reduce cost.
- Higher bacterial cell mass was observed; this will lead to higher H<sub>2</sub> output.



Batch	Cellulose Concentration	Time to Peak H <sub>2</sub> Production	Amount of H <sub>2</sub> Produced	Average H <sub>2</sub> Production Rate
	(g/L)	(t, h)	(mmoles)	(mmol L <sup>-1</sup> h <sup>-1</sup> )
1	2.5	<b>18:43</b>	14.92	0.60
2-4		4:14	13.85	
5	5.0	8:09	17.57	0.92
6-8		5:27	22.11	

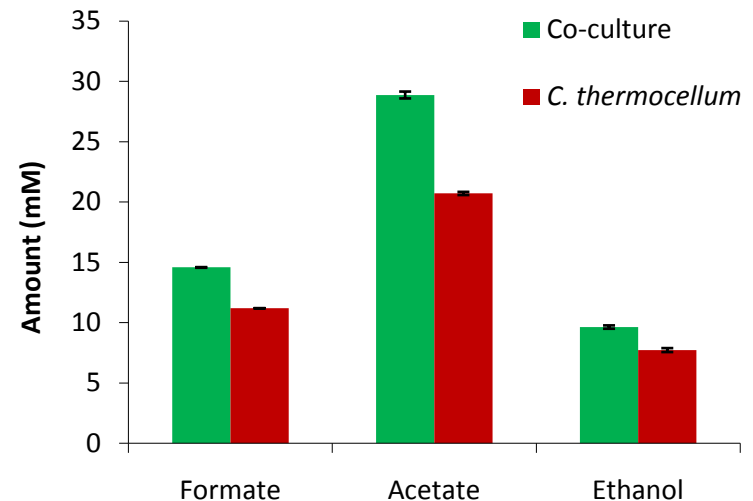
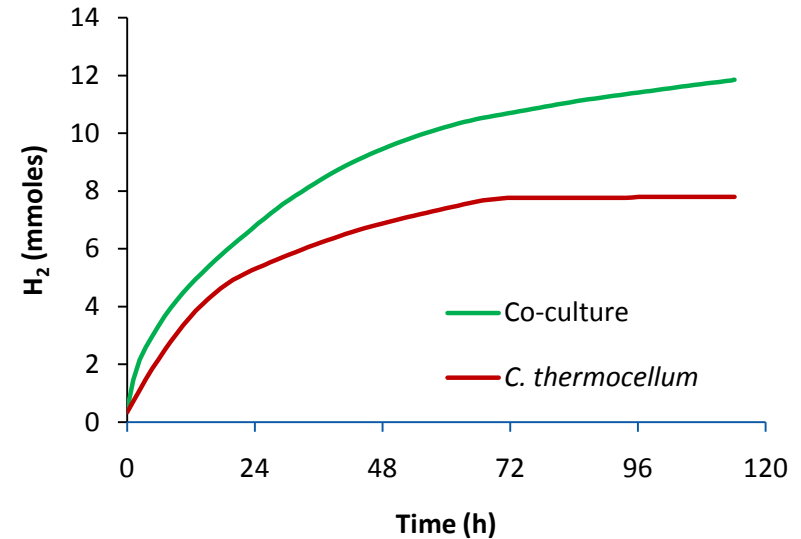
**On track to complete Milestone “Determine the solid and hydraulic retention time on rates and yield of H<sub>2</sub> in fed-batch reactor (4/11).”**

# Task 1 – Technical Accomplishments

## H<sub>2</sub> From “*Untreated*” Corn Stover Using a Co-Culture

- We scaled up fermentation of untreated corn stover using a co-culture of *Clostridium thermocellum* and a *Clostridium* consortium
- *C. thermocellum* hydrolyzed both cellulose and hemicellulose, with the latter utilized by the consortium.
- The co-culture produced **64%** more H<sub>2</sub> than *C. thermocellum* alone, suggesting better substrate utilization.
- The metabolite profiles also corroborate the synergy of the co-culture.

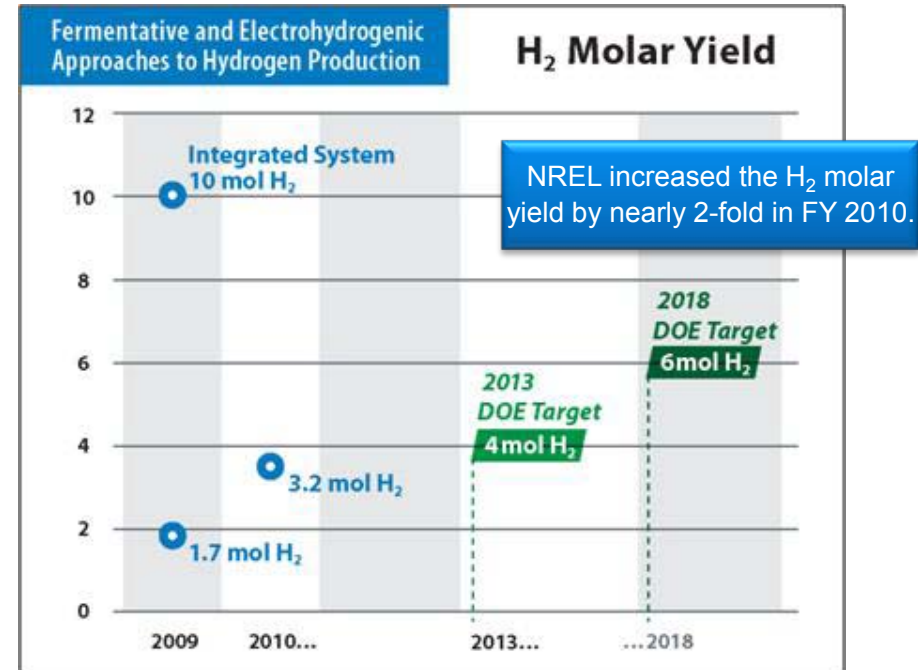
**The utilization of untreated biomass lowers feedstock cost.**



# Objectives/Approach/Milestone

## Task 2 – Develop Genetic Methods for Metabolic Engineering

- **Objective:** Improve H<sub>2</sub> molar yield (mol H<sub>2</sub>/mol hexose) via fermentation.
- **Approach:** Redirect metabolic pathways to maximize H<sub>2</sub> production via the development of genetic methods.
  - Design plasmids and optimize transformation protocols.
  - Create mutant host suitable for targeted mutagenesis.



	Milestone	Completion Date	Status
3.2.2	Produce one genetic transformant in <i>C. thermocellum</i> (FY10)	9/10	Completed
3.2.2	Obtain one mutant of <i>C. thermocellum</i> lacking the <i>pyrF</i> gene as the platform host for targeted mutagenesis (FY11)	9/11	On track



# Task 2 – Technical Accomplishments

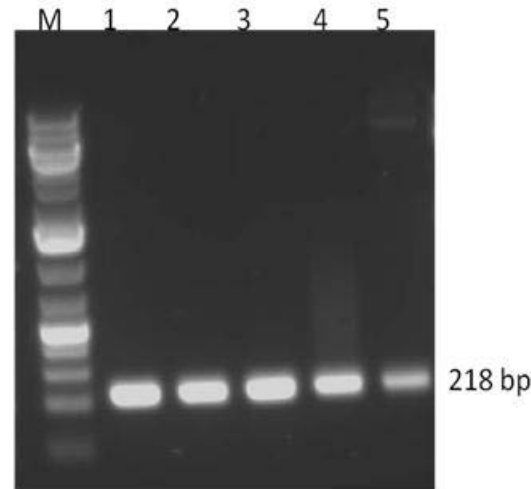
## Develop Tools for Genetic Transformation

- NREL co-developed genetic tools (with the University of Manitoba): proprietary plasmid and transformation protocols, and obtained two mutants of *C. thermocellum*.
- Transformation was verified by (1) growth in antibiotic (chloramphenicol, 100 µg/mL); (2) PCR of the antibiotic gene; and (3) retransformation in *E. coli*.
- This will lead to blocking competing pathways to improve H<sub>2</sub> molar yield.



### Cell growth:

1. Transformant in antibiotic
2. Transformant without antibiotic
3. Control cells in antibiotic
4. Control cells without antibiotic.



### PCR of the antibiotic gene:

- Lanes 1-2: transformant #1;
- Lanes 3-4: transformant #2;
- Lane 5: plasmid control.

**Completed Milestone “Produce one genetic transformant in *Clostridium thermocellum*” (8/10).**

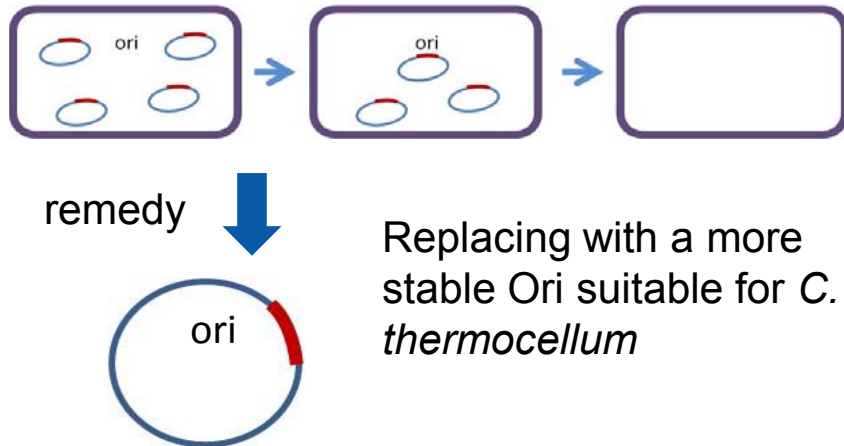
# Task 2 – Technical Accomplishment

## Develop Targeted Pathway Mutant

### Improve plasmid stability

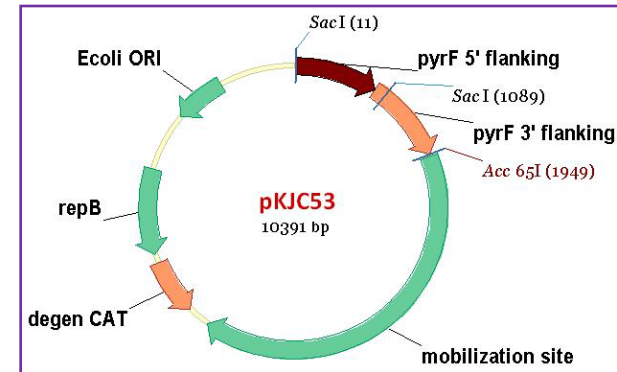
- We discovered that the proprietary plasmid is not stable in the transformant. Work is underway to replace its origin of replication (ori) to improve its long-term stability.

Subculturing



### Block pyruvate-to-formate pathway

- Blocking the pyruvate-to-formate pathway (with hypophosphite) increased  $H_2$  production by 81% (FY09) – proof of concept.
- Work is ongoing to create a  $\Delta pyrF$  mutant host to generate the targeted pathway mutant via an effective suicide method.



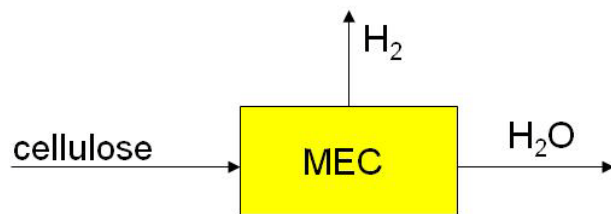
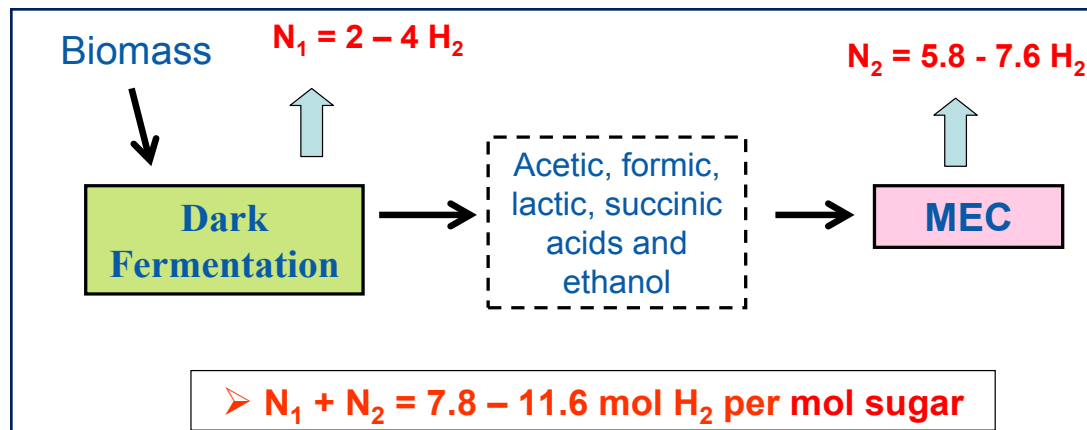
**On track to complete Milestone “Obtain one mutant of *C. thermocellum* lacking the *pyrF* gene as the platform host for targeted mutagenesis” (9/11).**

# Objectives/Relevance

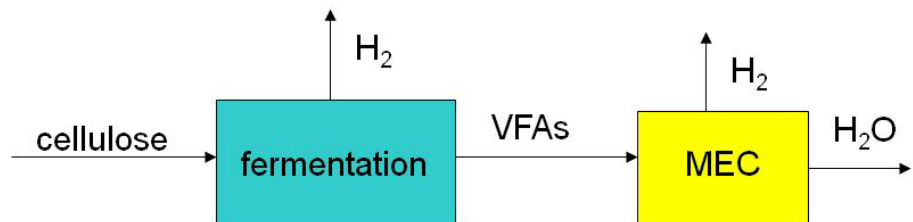
## Task 3 – Electrochemically Assisted Microbial Fermentation



**Objective:** Improve  $H_2$  molar yield (mol  $H_2$ /mol hexose) by integrating dark fermentation with microbial electrolysis cell (MEC) reactor to convert waste biomass to additional  $H_2$ .



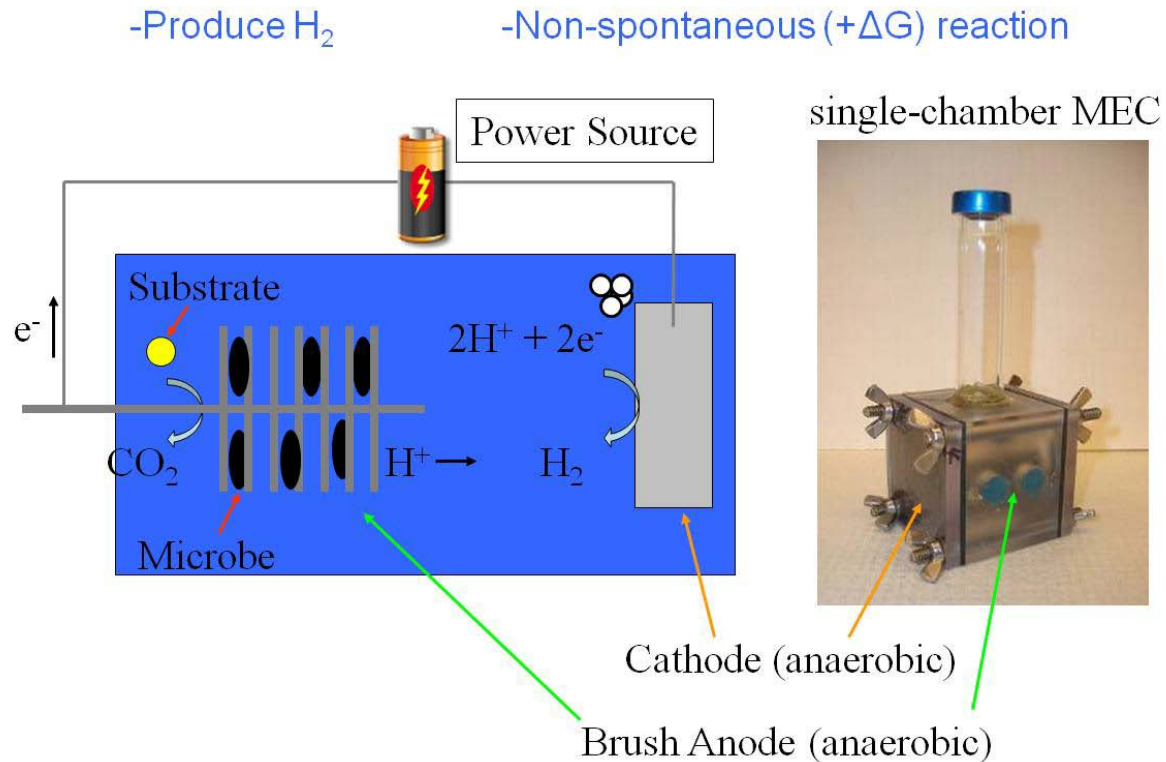
One-stage process: slow



Two-stage process: fast

# Approach/Milestone

## Subtask 3: Electrochemically Assisted Microbial Fermentation

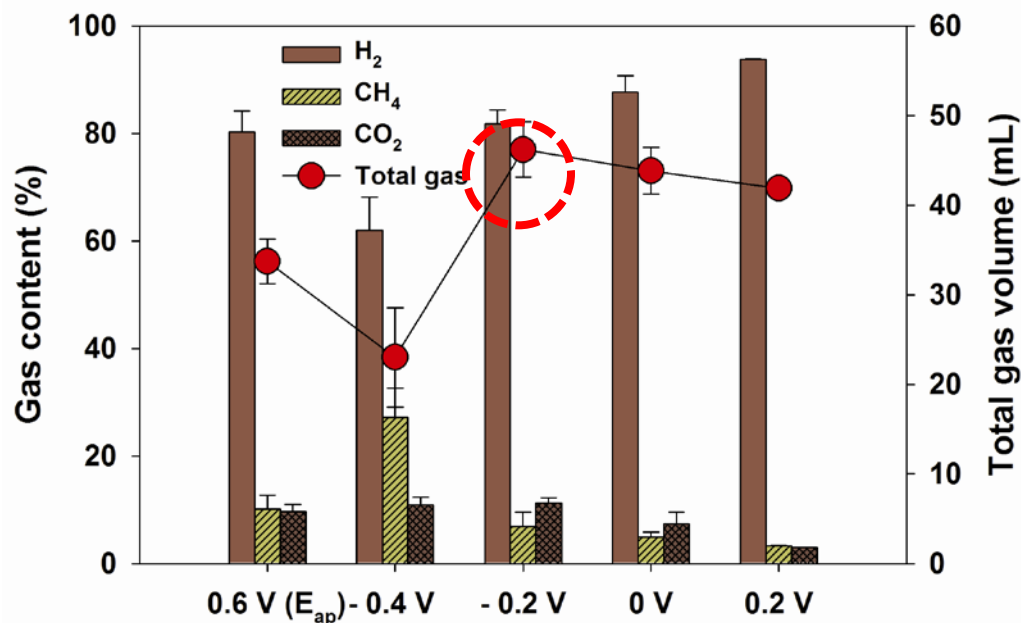


	Milestone	Completion Date	Status
3.2.3	Prototype reactor operational (FY10)	9/10	Completed
3.2.3	Correlate removal of the subcomponents of the NREL fermentation effluent with current density and H <sub>2</sub> production (FY11)	9/11	On track

# Task 3 – Technical Accomplishments

## Methane Reduction by Setting Anode Potential

- Compare “boosting voltage ( $E_{ap}$ )” with a power source (typically adding 0.6 V) with a set anode potential
- When adding 0.6 V, the anode is typically at -0.4 V, so we chose this and more positive anode potentials
- Highest gas production at  $E_{an} = -0.2$  V



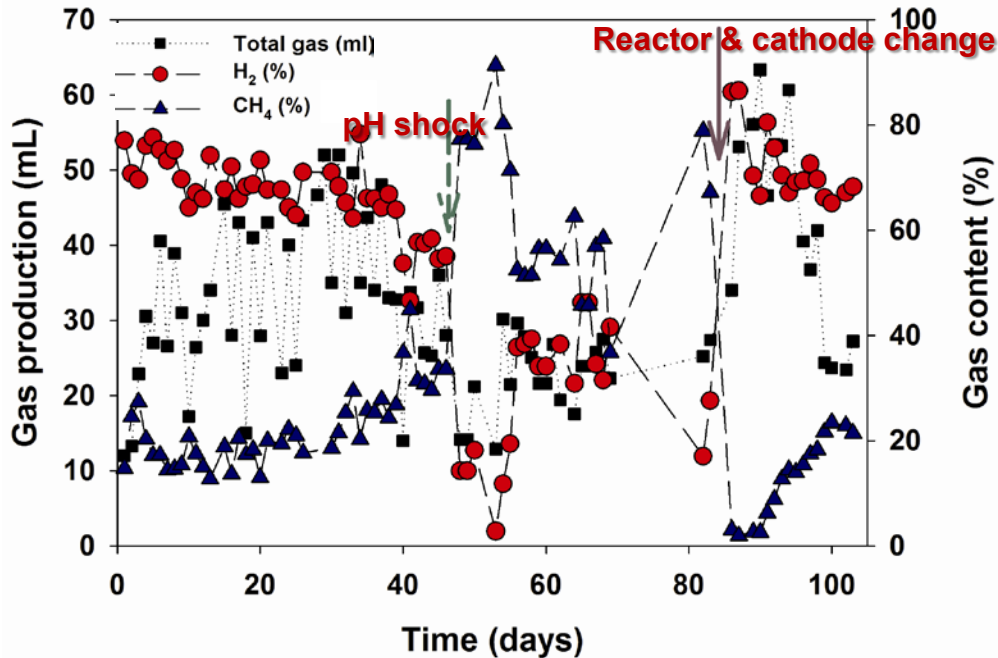
V	Energy Input (kWh/m <sup>3</sup> <sub>reactor</sub> )	Energy Input (kWh/m <sup>3</sup> <sub>H<sub>2</sub></sub> )	$\eta_{E+S}$ (%)	$\eta_E$ (%)	$\eta_S$ (%)	Cycle Time (h)
0.6 ( $E_{ap}$ )	1.7	1.7	57	187	81	26
-0.4	1.2	2.9	27	114	34	40
-0.2	<b>3.0</b>	<b>2.3</b>	<b>59</b>	<b>143</b>	<b>97</b>	<b>16</b>
0	3.9	2.9	55	113	102	10
0.2	5.5	4.7	40	71	90	8





# Task 3 – Technical Accomplishments

## Methane Reduction by Setting Anode Potential



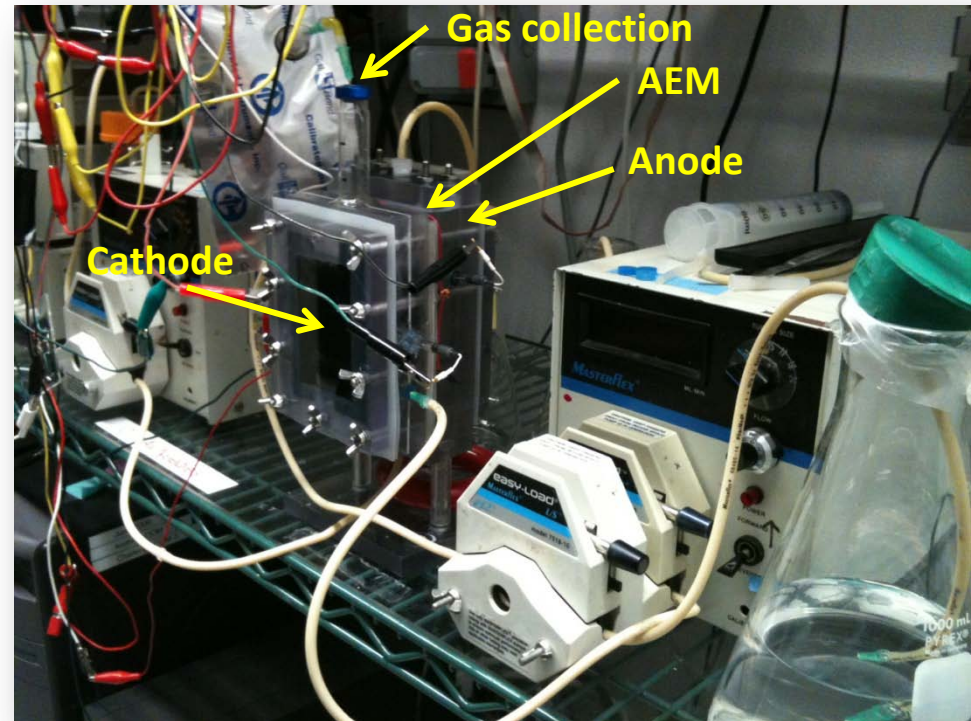
- Continuous operation at  $E_{An} = -0.2 \text{ V}$
- Day 39, 68% H<sub>2</sub>, 21% CH<sub>4</sub>  
(average daily gas production 34 mL)
- → CH<sub>4</sub> increased up to 34% by Day 46.
- pH shock tried... didn't work.
- What did work?  
Placing the brush anode into new clean reactor (Day 85) → 86% H<sub>2</sub>, 3% CH<sub>4</sub>  
→ Methanogens were not on anode, but in reactor assembly

**Conclusion:** Setting anode potential can improve cell performance, but it was difficult to eliminate CH<sub>4</sub> completely in single-chamber MECs.



# Task 3 – Technical Accomplishments

## Design New Tubular MECs

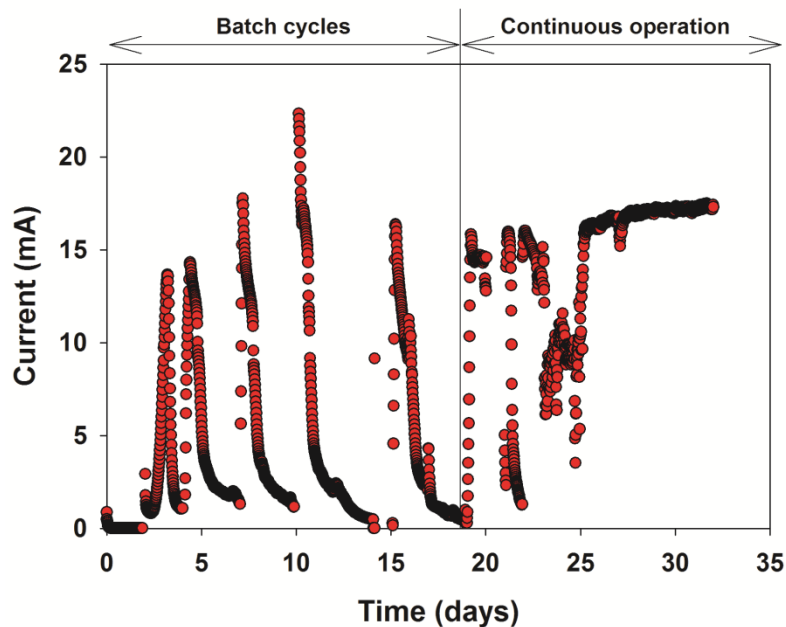


- A two chambered MEC with AEM , 0.9 V applied
- Anode chamber (135 mL) → 3 brush anodes, 1.5 g sodium acetate/L, HRT = 1 day
- Cathode chamber (147 mL) → Pt coated stainless steel, 50 mM PBS, HRT = 1 day

Completed Milestone “*Design tubular MECs to reduce methane generation*” (9/10)

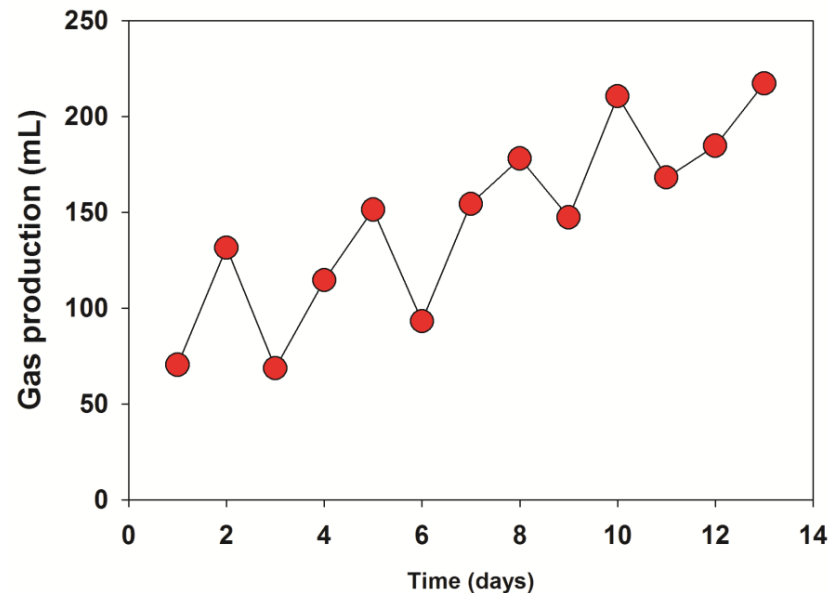
# Task 3 – Technical Accomplishments

## Tubular MEC Performance



Stable current generation in a continuous mode → 17 mA, current density 60 A/m<sup>3</sup>.

<Continuous gas generation>



→ Almost pure H<sub>2</sub> (no CH<sub>4</sub>, CO<sub>2</sub>) with stable current

→ CE 79-84%, r<sub>cat</sub> = 88-107%, COD<sub>rem</sub> > 90%  
H<sub>2</sub> production rate 0.8 H<sub>2</sub> m<sup>3</sup>/m<sup>3</sup>-d (Day 13)

**Completed “Test performance of the new system” (12/10).**

**On track to complete Milestone “Correlate removal of the subcomponents of the NREL fermentation effluent with current density and H<sub>2</sub> production” (8/11).**

# Collaborations

- **Task 1 (Bioreactor):**

Dr. Ali Mohagheghi, National Bioenergy Center at NREL (Biomass pretreatment and characterization)

- **Task 2 (Genetic Methods):**

- Dr. Mike Himmel at NREL (funded by the DOE BER Program)
- Drs. David Levin and Richard Sparling at the University of Manitoba, Canada (funded by the Genome Canada Program). NREL is an international collaborator in the Genome Canada Grant award to co-develop genetic tools for pathway engineering in *C. thermocellum*.

- **Task 3 (MEC):**

Dr. Bruce Logan, Penn State University (microbial electrolysis cells to improve H<sub>2</sub> molar yield)



# Proposed Work



## Task 1:

- Repeat fed-batch experiments (2.5, 5, and 10 g/L cellulose) in one-liter bioreactor for cellulose consumption, carbon balance, rates and yield of H<sub>2</sub> (FY11).
- Scale up fed-batch experiment in 5-L bioreactor as above, testing hydraulic retention time and feeding strategy (FY11/12).
- Collect, analyze, and send fermentation effluent to PSU to generate H<sub>2</sub> via MEC integration (FY11/12).
- Type microbial community of the consortium if supported by the Program (FY12).

## Task 2:

- Modify the proprietary plasmid to improve its long-term stability (FY11).
- Construct a *C. thermocellum* mutant host (*pyrF* knockout) and start deleting pyruvate-to-formate pathway (FY11).
- Test the above mutant for H<sub>2</sub> rates, yield, and carbon balance (FY11/12).
- Target additional competing pathways to improve H<sub>2</sub> molar yield (FY12).

## Task 3:

- Complete analysis of fermentation effluent (FY11).
- Test performance of the new system with fermentation effluent (FY11).
- Conduct tests on performance with respect to hydrogen yields, hydrogen production rates, and gas composition for the fermentation effluent (FY11/12).



# Summary

## Task 1:

- Operated fed-batch reactor with substrate concentrations of 2.5 and 5.0 g/L and hydraulic retention time of 24 hr, and observed shorter acclimation time, improved rates of H<sub>2</sub>, and elevated total cell biomass concentrations.
- Demonstrated that a co-culture (*C. thermocellum* and a *Clostridium* consortium) can ferment untreated corn stover and yielded 64% more H<sub>2</sub> than *C. thermocellum* alone. The outcomes work toward reducing feedstock cost.

## Task 2:

- Developed genetic tools via collaboration and generated *C. thermocellum* mutants harboring the proprietary plasmid.
- Continue to improve stability of the plasmid to allow plasmid-based expression.
- Work is ongoing to generate a *C. thermocellum* mutant host for deleting competing pathway.

## Task 3:

- Setting anode potentials in single chamber MECs could improve reactor performance, but did not completely eliminate methane generation.
- Higher gas production and H<sub>2</sub> composition were obtained at  $E_{An} = -0.2$  V compared to adding voltage of  $E_{ap} = 0.6$  V.
- By designing and operating a tubular MEC composed of two chambers, pure H<sub>2</sub> was obtained. Achieved up to 0.8 H<sub>2</sub> m<sup>3</sup>/m<sup>3</sup>-d.