

# Sweet Hydrogen: High-Yield Production of Hydrogen from Biomass Sugars Catalyzed by *in vitro* Synthetic Biosystems

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**6/7/2017 (2:45:00 PM)**

**Project ID PD127**

# Overview

## Timeline

- Project Start Date: 06/15/2015
- Project End Date: 12/31/2017 (2.5 years), including a half-year no-cost extension from June 30, 2017 to Dec. 31 2017.

## Budget

- Total Project Budget: \$937,602 (two years)
  - Total Recipient Cost-Sharing: \$187,602 (97,882 VT part)
  - Total Federal Share: \$750,000
  - Total DOE Funds Spent\*: \$611,386\* As of 3/31/2017

## Barriers

### Hydrogen production from biomass

AX. Hydrogen Molar Yield

AY. Feedstock Cost

AZ. Systems Engineering

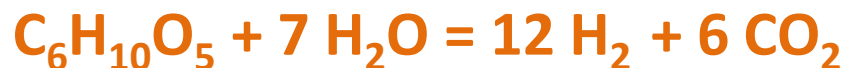
## Collaborators

- Virginia Tech (lead)
- University of Georgia

# Relevance

**Fuel Cell Technologies Office Objective** -- Develop advanced biological generation technologies to produce hydrogen with a projected cost of \$10.00/gge at the plant gate by 2020.

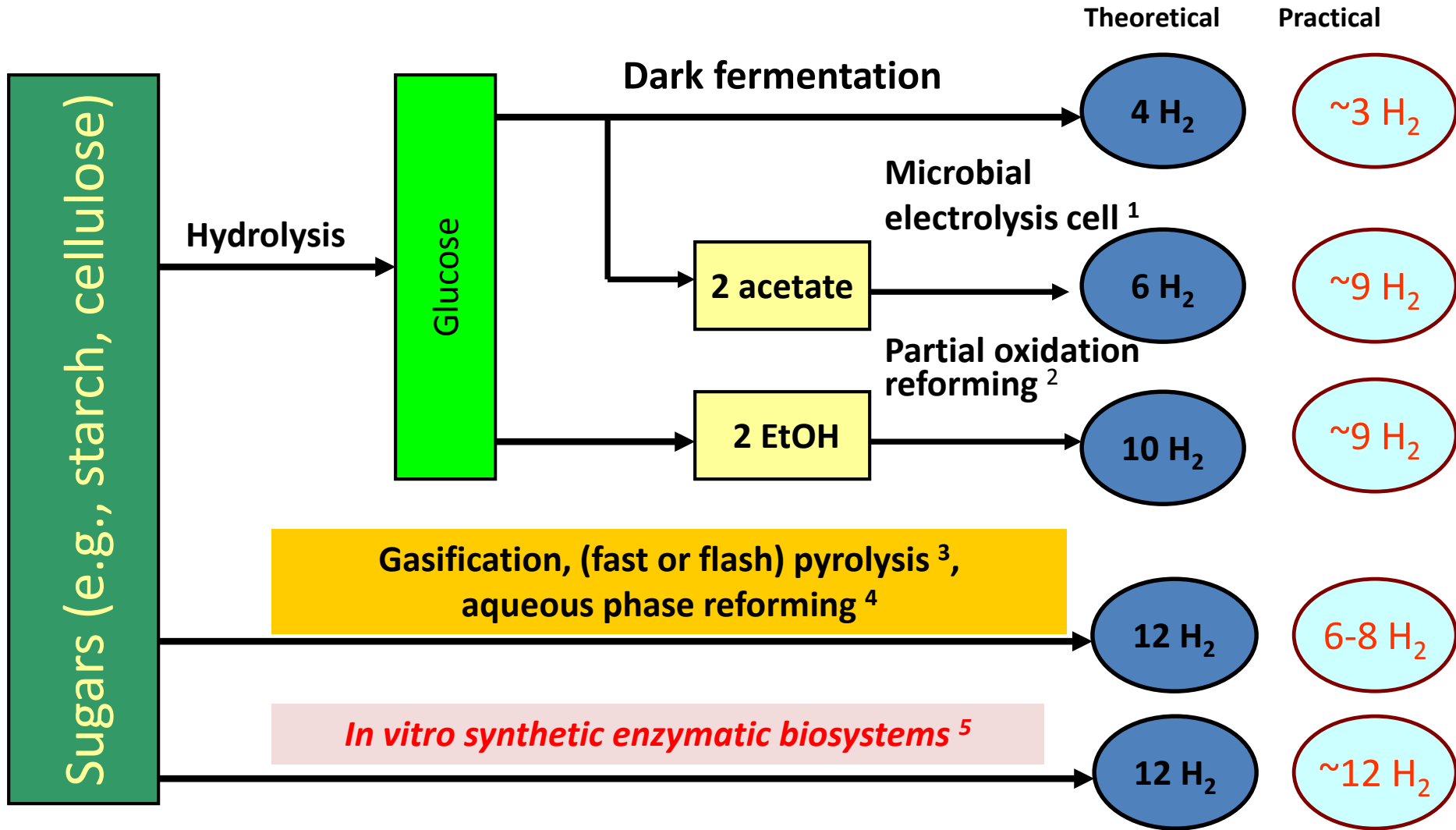
**Our novel approach – use renewable sugars (e.g., biomass sugars or starch) to split water to produce H<sub>2</sub> catalyzed by enzyme cocktails**



- Low-carbon production in terms of entire life cycle
- High-purity hydrogen generated (no CO)
- Mild reaction conditions (1 atm, 20-90°C, pH 7.5, aqueous phase)
- Local resources for distributed hydrogen small-size production (e.g., 1,500 kg/day)
- Highest biological hydrogen generation rates (e.g., **550 mmole H<sub>2</sub>/L/h**, now)

| Targets         | Units                     | June 2017 Target | Dec. 2017 Target (estimated) | Year 2020 Target (Plant gate)    |
|-----------------|---------------------------|------------------|------------------------------|----------------------------------|
| Production cost | \$/kg H <sub>2</sub>      | 1000             | 10                           | 10.00 (year 2020)                |
| Productivity    | mmole H <sub>2</sub> /L/h | 750              | 750                          | 2,000                            |
| Reactor volume  | L of reactor              | 1                | 1                            | 65,000*                          |
|                 |                           |                  |                              | *1,500 kg H <sub>2</sub> per day |

# Production of H<sub>2</sub> from sugars (overview)



1. Logan et al. 2007. PNAS 104:18871.

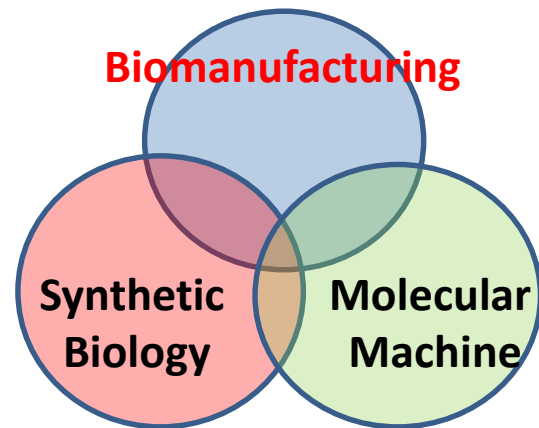
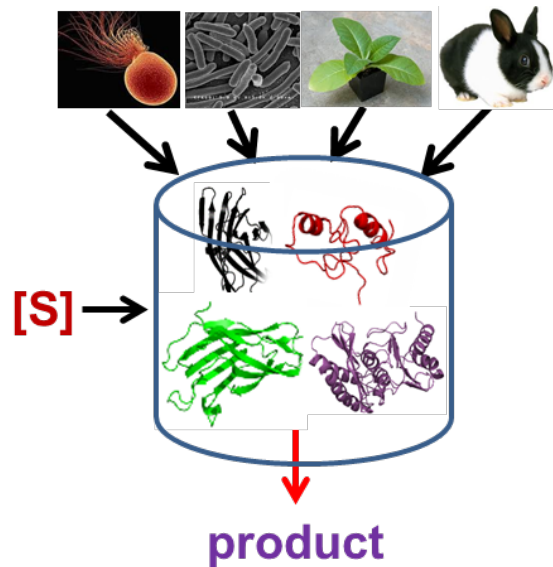
2. Schmidt. 2004. Science. 303:993

3. Schmidt. 2007. Agnew Chem. 46: 586

4. Dumesic et al. 2002. Nature 418:964

5. Zhang et al. PLoS One 2007; ChemSusChem 2009; Angew Chem 2013; Metab Eng 2014; PNAS 2015.

## Advanced biotransformation catalyzed by *in vitro* synthetic enzymatic biosystems



### Unique features

**A disruptive platform, against the dominant biotech paradigm.**

- High product yield
- High energy efficiency (oxygen-free biotransformation)
- Fast volumetric productivity
- Broad (mild) reaction conditions
- Easy product separation
- High product titer
- Tolerance of toxic compounds (e.g., substrate, product, solvent)
- Great engineering flexibility
- Easy process control and scale-up
- Low capital investment (CapEx)
- Accomplish non-natural bio-reactions

# Approaches/Milestones

(\*As of 3/31/2017)



|                                                                                 |           |      |
|---------------------------------------------------------------------------------|-----------|------|
| Task 1.1. Co-expression of multiple enzymes in one host                         | (FY17 Q1) | 100% |
| Task 1.2. Two redox enzymes on biomimics at 1 U/mg                              | (FY17 Q3) | 80%  |
| Task 1.4. H <sub>2</sub> production cost of \$10/kg H <sub>2</sub> by H2A model | (FY17 Q3) | 75%  |
| Task 2.2. Data fitting and validating of rate-limiting steps                    | (FY16 Q4) | 100% |
| Task 2.3. Construction of artificial electron transport chains                  | (FY17 Q3) | 100% |
| Task 2.4. Construction of five synthetic enzyme complexes                       | (FY17 Q3) | 100% |
| Task 3.1. High-density of protein expression in <i>E. coli</i>                  | (FY17 Q3) | 100% |
| Task 3.2. Mass production of hydrogenase (SH1) (UGa)                            | (FY17 Q3) | 70%  |
| Task 3.3. 1000-mL level demonstration                                           | (FY17 Q3) | 90%  |

**This reporting period (Dec 2017, a final deliverable)**

- ❑ **1,000-fold volume scale-up (1 L reactor) with 5-fold increase in H<sub>2</sub> peak production rate (i.e., 750 mmol H<sub>2</sub>/L/h or 1.5 g H<sub>2</sub>/L/h)**

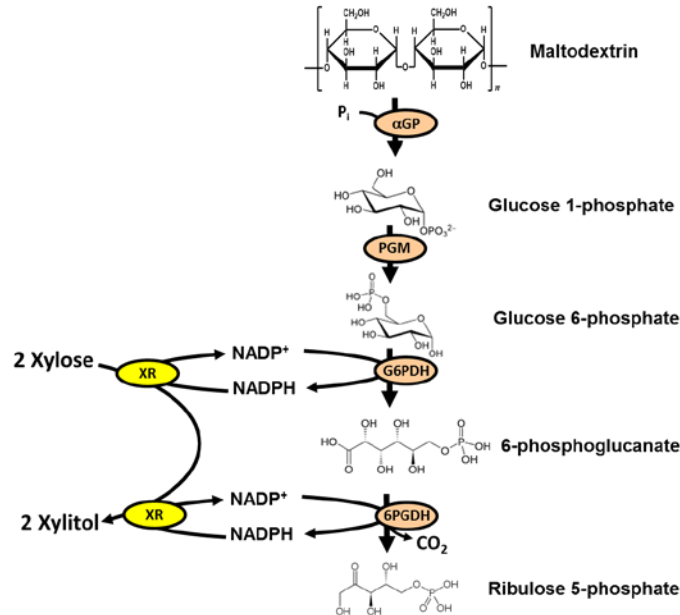
# Accomplishments and Progress:

## Responses to Previous Year Reviewers' Comments

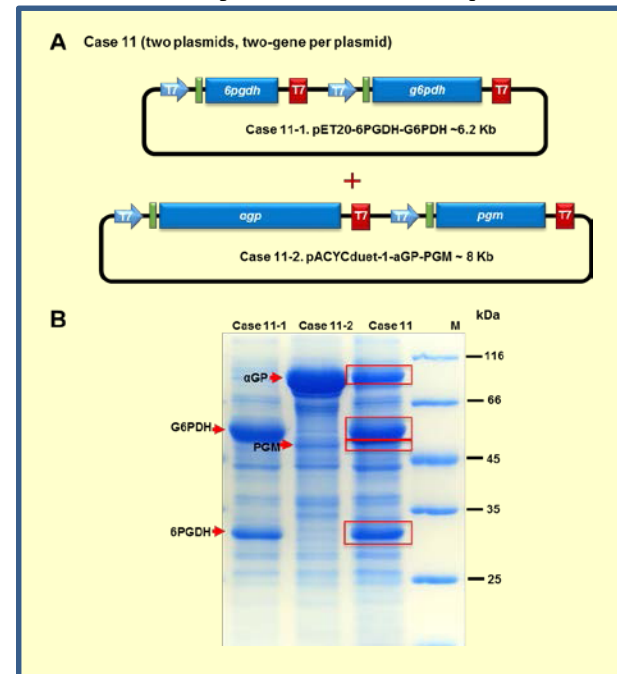
| FY16 reviewers' comments                                                                                                                                                                                                                                           | FY17 response to comments                                                                                                                                                                                                                                                                                                                                                              |
|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <b>Substrate cost.</b> <i>“This in vitro approach will likely be effective only for very clean sugars/starches and so will have somewhat limited greenhouse gas benefits when compared to a lignocellulosic system.”</i>                                           | Starch (\$0.30/kg) is the cheapest sugar. Glucose made from starch is cheaper than that made from biomass. It can use biomass sugars in the presence of toxic compounds (PNAS 2015). Also, we will make artificial starch from biomass (PNAS 2013) and via artificial photosynthesis (Energy Sci. Eng., 2013). <u>Starch may be the best solar fuel and hydrogen storage compound.</u> |
| <b>Enzyme production cost.</b> <i>“With the <u>synthesis of the enzymes</u> and <u>identification of challenges/solutions with the cofactors</u>, the project appears to be off to a promising start in terms of meeting the H<sub>2</sub> volume objectives.”</i> | Current production costs of recombinant thermophilic enzymes in <i>E. coli</i> are \$50/kg dry enzyme or lower (JIMB 2017). It may be decreased to \$10-20/kg, like amylase, protease (JIMB 2017). This first industrial biomanufacturing example with in vitro biosystem has been established in China (B&B 2017).                                                                    |
| <b>Coenzyme replacement and costs</b> (ibid)                                                                                                                                                                                                                       | Replacement of NADP with NMN and NAD-conjugate could decrease H <sub>2</sub> production costs to less than \$10/kg H <sub>2</sub> , when enzyme costs and stability are addressed.                                                                                                                                                                                                     |
| <b>H<sub>2</sub> production rates.</b> <i>“how long certain hydrogen productivity rates could be achieved”</i>                                                                                                                                                     | We demonstrate up to 3-h high-speed H <sub>2</sub> production (i.e., 1 g H <sub>2</sub> /L/h) in a batch mode (see following PPT).                                                                                                                                                                                                                                                     |

# Accomplishments: Four-enzyme coexpression

Four enzymes:  $\alpha$ GP, PGM, G6PDH & 6PGDH



Four enzymes in two plasmids



Chen et al. 2017.  
Appl. Microbiol.  
Biotechnol.:Doi:  
10.1007/s00253-  
017-8206-8.

Table 3 Comparison of four-enzyme co-expression cases in *E. coli* and the activity based on xylitol generation

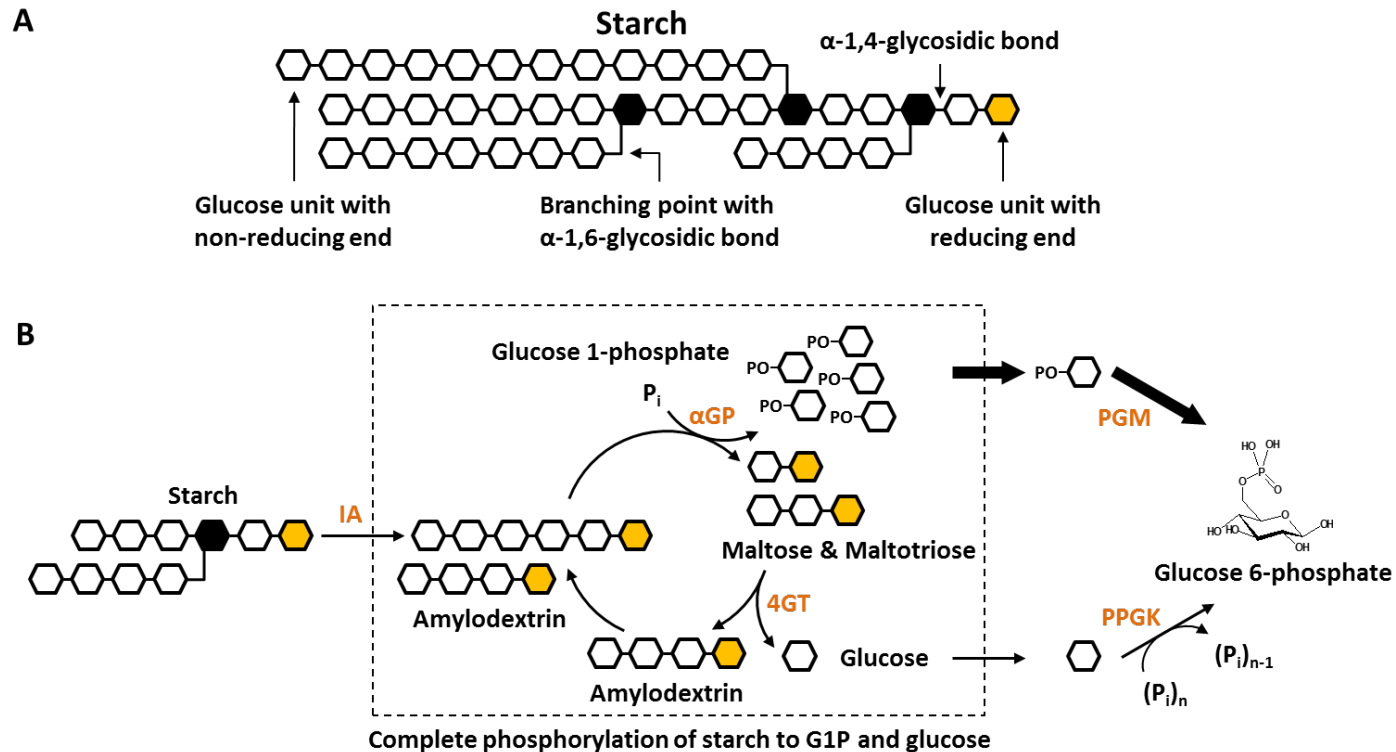
| Case name | Protein expression percentage (%) |      |       |       |     | Apparent enzyme activity (U/mL) <sup>a</sup> |     |       |       | Overall activity (mM) <sup>b</sup> |
|-----------|-----------------------------------|------|-------|-------|-----|----------------------------------------------|-----|-------|-------|------------------------------------|
|           | $\alpha$ GP                       | PGM  | G6PDH | 6PGDH | Sum | $\alpha$ GP                                  | PGM | G6PDH | 6PGDH |                                    |
| Case 8    | 30                                | 6.2  | 4.2   | 1.6   | 42  | 2.8                                          | 16  | 0.29  | 0.10  | 15.48                              |
| Case 9    | 28                                | 6.9  | 3.9   | 1.3   | 40  | 2.6                                          | 18  | 0.27  | 0.083 | 15.27                              |
| Case 10   | 16                                | 5.4  | 3.6   | 17    | 42  | 1.5                                          | 14  | 0.25  | 1.1   | 16.67                              |
| Case 11   | 17                                | 0.75 | 28    | 20    | 66  | 1.6                                          | 1.9 | 1.9   | 1.3   | 24.58                              |

We found out that the best strategy (two different strength plasmids, each has two genes encoded) to precisely control four enzyme expression levels in *E. coli* BL21(DE3).



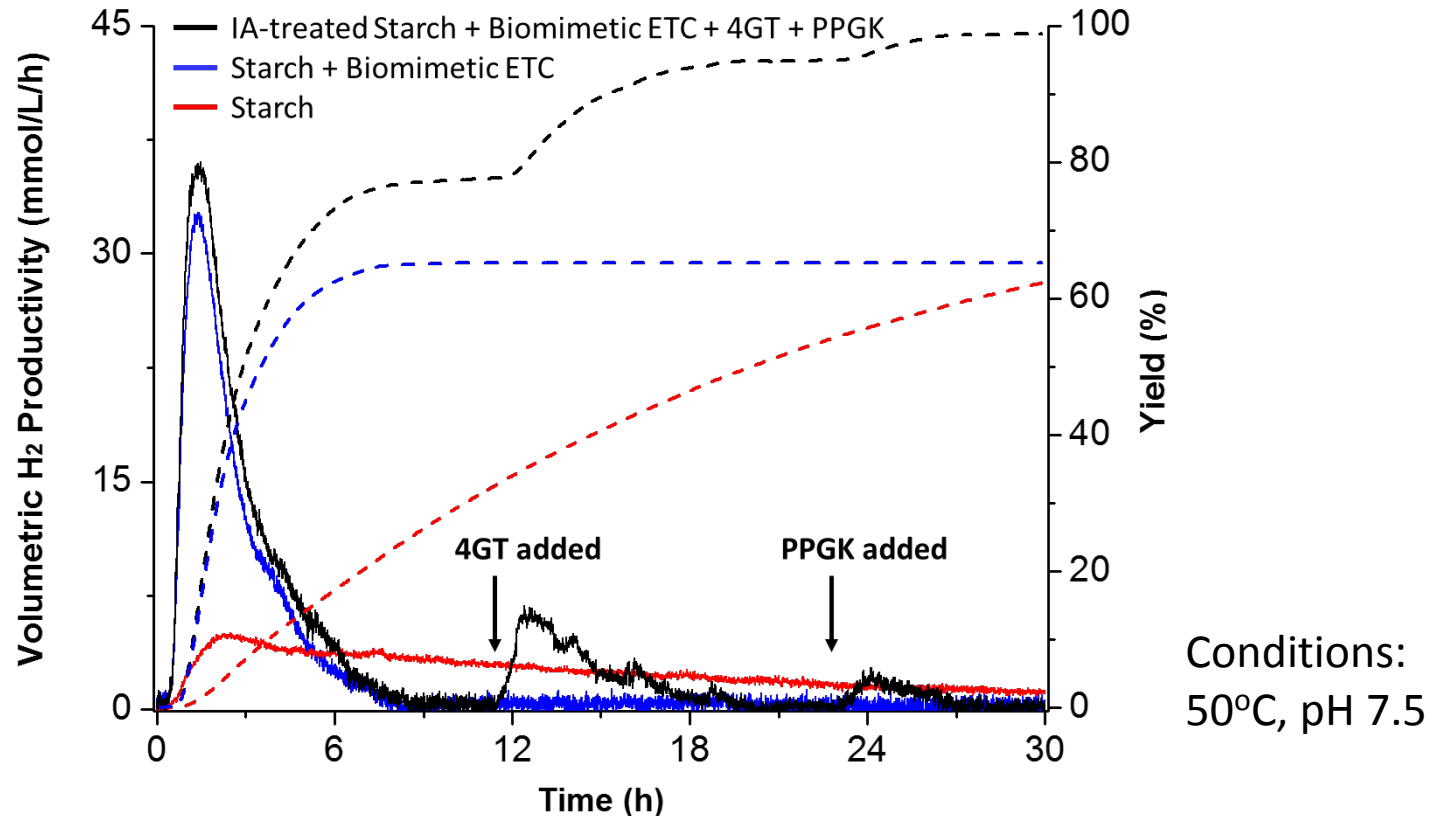
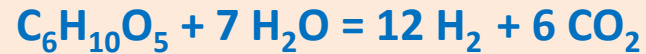
# Achievements: Complete starch utilization for H<sub>2</sub> generation

## Enzymatic starch phosphorylation without ATP



- Alternative phosphorolysis of starch, yielding G6P (better than simple hydrolysis).
- Use isoamylase to debranch branched starch (amylopectin) to increase H<sub>2</sub> yield by 30%
- Use 4-glucoanotransfer (4GT) to utilize maltose to increase H<sub>2</sub> yield about 4%.
- Polyphosphate glucokinase (PPGK) to utilize glucose to increase yield about 4%.

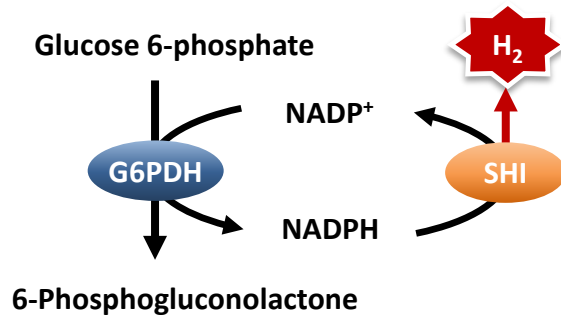
# Achievements: Complete starch utilization for H<sub>2</sub> generation (2)



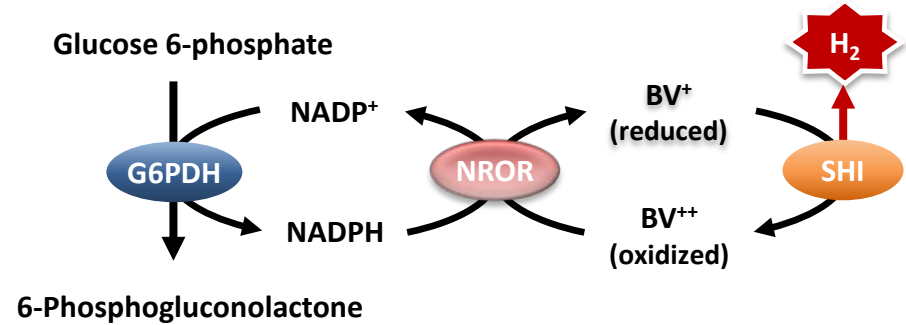
For the first time we can utilize all glucose units of starch without ATP for H<sub>2</sub> generation.  
**\$0.30/kg starch = \$2.02/kg of H<sub>2</sub>** (our goal is \$10/kg delivered H<sub>2</sub>).

# Accomplishments: Increasing Rxn by artificial electron transport chains

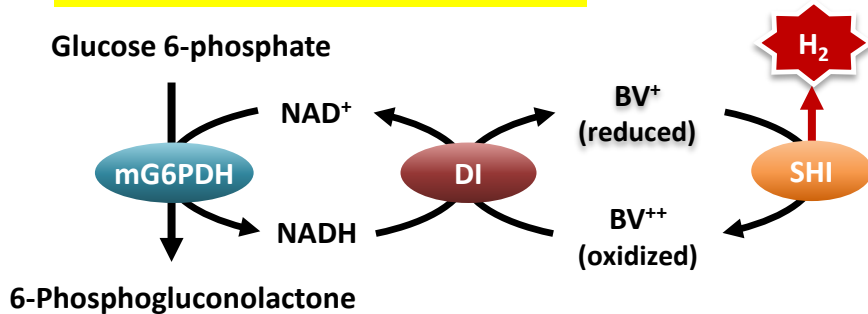
## a Original design (2007)



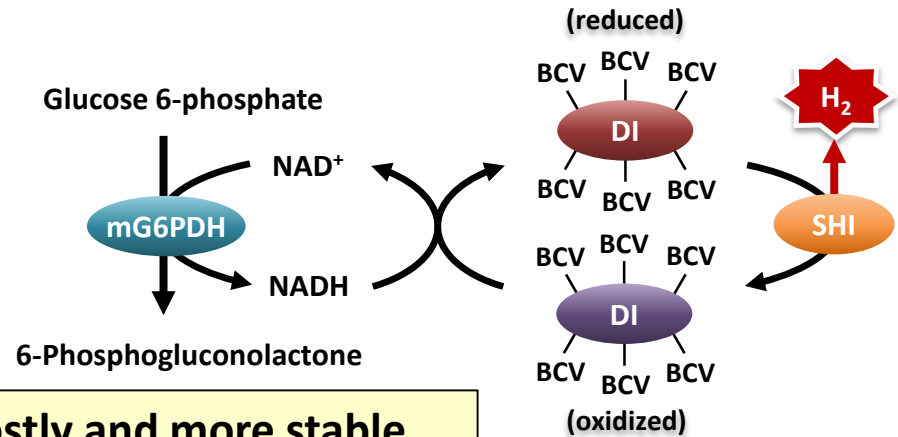
## b Revised 2016



## c NAD-based PPP 2017



## d NAD-based PPP + conjugate 2017



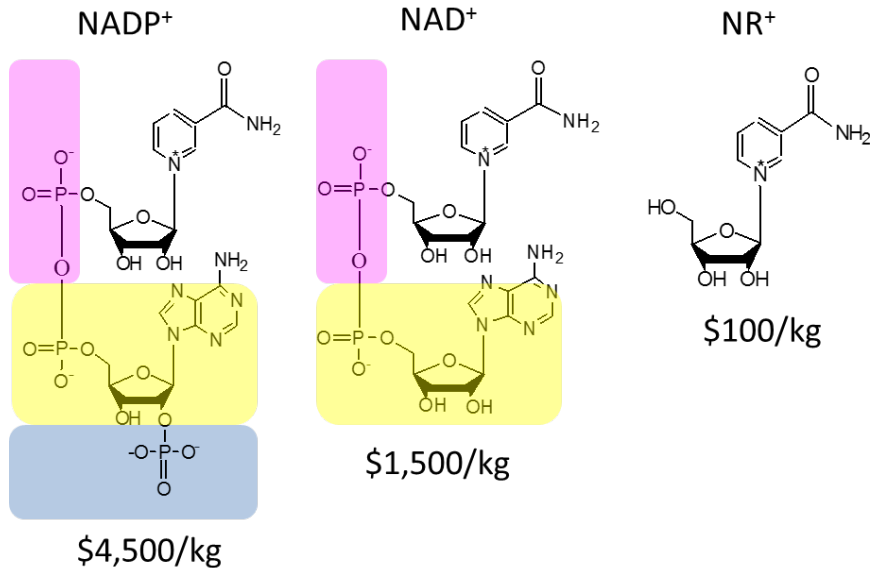
- Decreasing production costs by using less costly and more stable coenzymes instead of NADP<sup>+</sup>.

# Progress: Replace NAD(P) of dehydrogenases with NAD & NMN

Glucose 6-phosphate dehydrogenase (G6PDH)  
6-phosphogluconate dehydrogenase (6PGDH)

Specific activity order  
+++++ (NADP), ++ (NAD), - (NMN), - (NR)

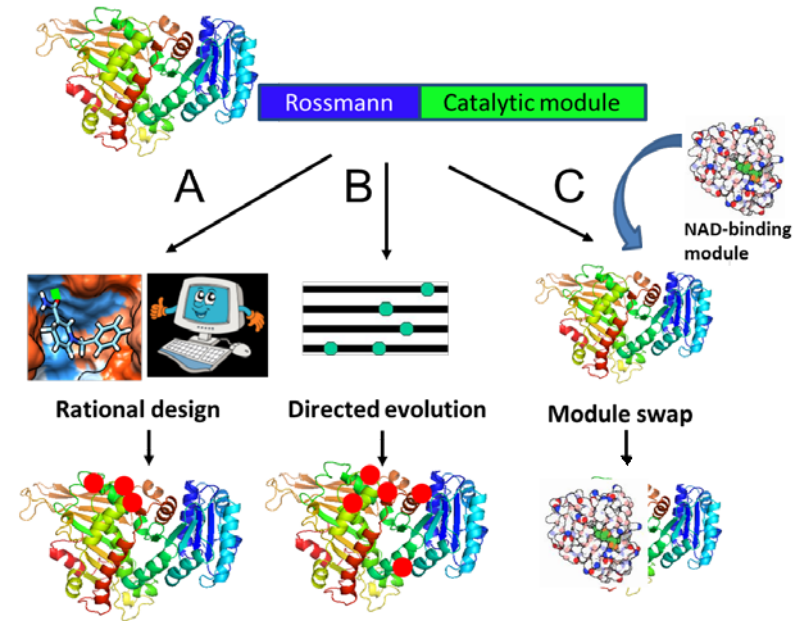
## Comparison of coenzymes



**NAD and nicotinamide riboside (NR) are better than NAD(P)**

- Less costly
- More stable
- Small size – better mass transfer

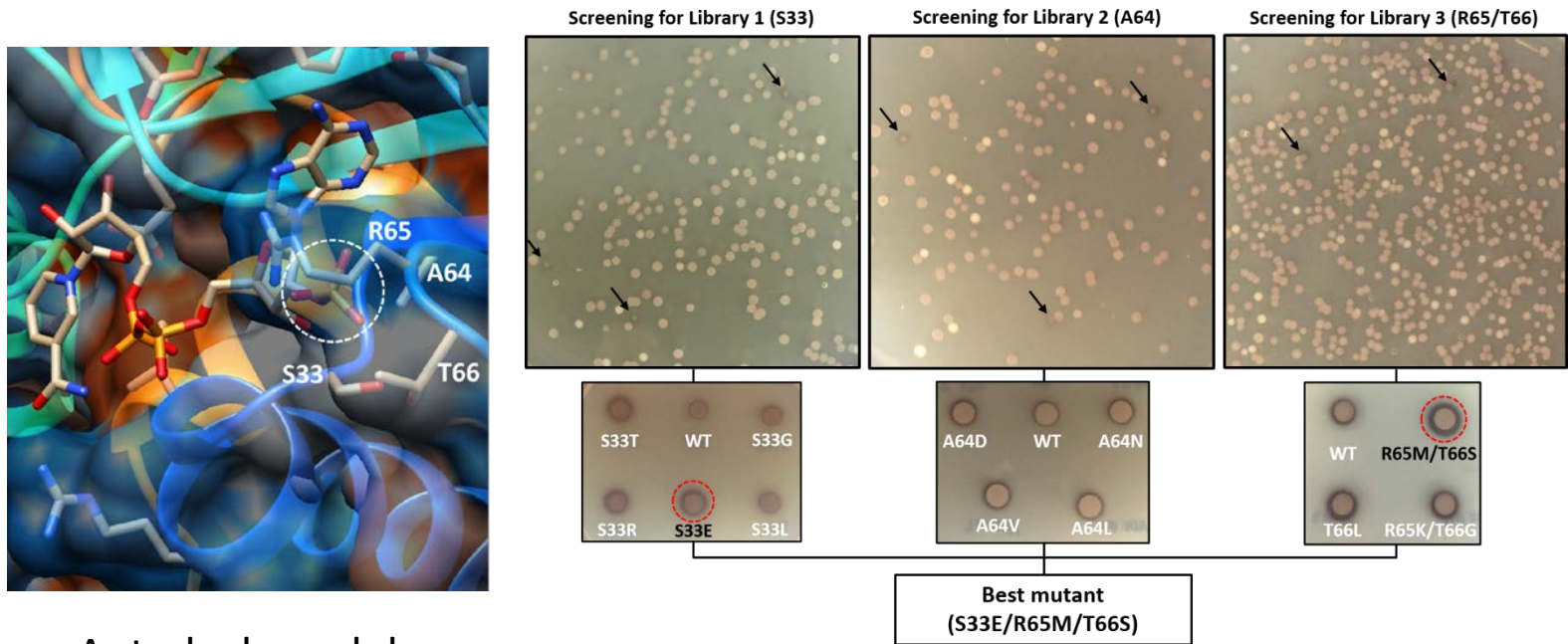
## Coenzyme engineering



- We have three strategies to change coenzyme preference of dehydrogenases
- New area – very high risk.

# Achievement: TmG6PDH coenzyme from NADP to NAD

## Identification of key amino acids followed by high-throughput screening



## Redox dye-based screening plate

The best mutant TmG6PDH (S33E/R65M/T66S) exhibited a preferred coenzyme from NADP<sup>+</sup> to NAD<sup>+</sup>.

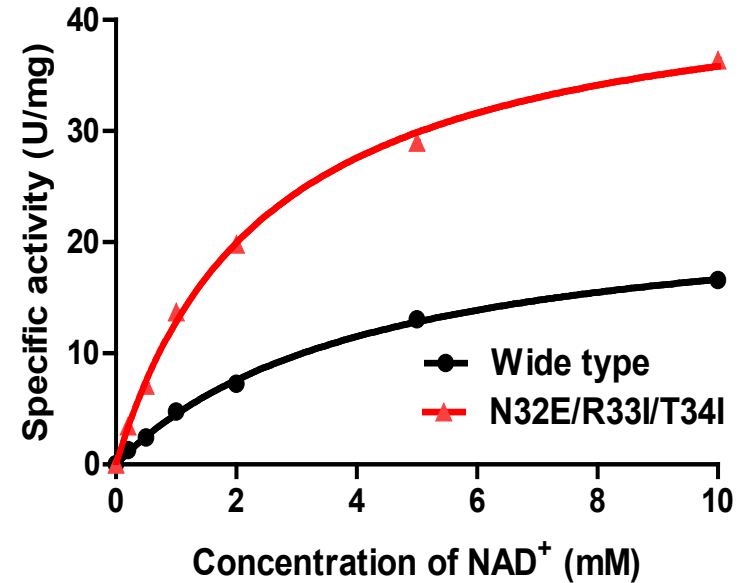
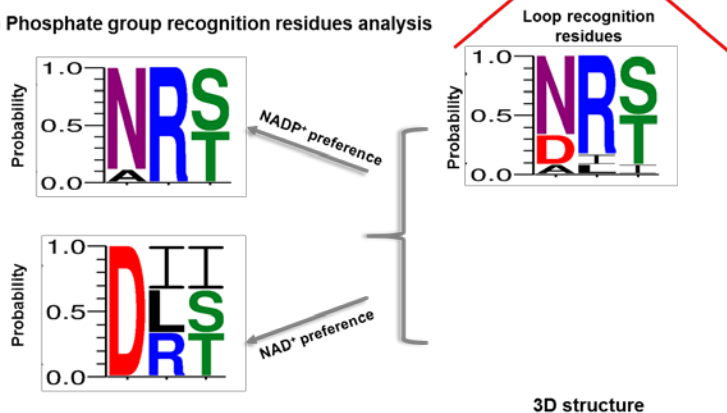
# Achievement: Tm6PGDH coenzyme from NADP to NAD

## Rational design

### A) Conventional sequence alignment

|                                                                                                                                                                                                                                                                                                        |                                                                                                                                                                                                                                                                                                                                                                                                                                    |
|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <i>Cyanophora paradoxa</i><br><i>Euglena gracilis</i><br><i>Saccharomyces cerevisiae</i><br><i>Corynebacterium diphtheriae</i><br><i>Lactococcus lactis</i> subsp<br><i>Bacillus subtilis</i><br><i>Escherichia coli</i><br>* <i>Thermotoga maritima</i><br>* <i>Morella thermoacetica</i> (wide type) | <b>NAD<sup>+</sup> preference 6PGDH</b><br>MAVMGQNLALNIAEEGLPTVSPNRS PDKVDDTVARA<br>LAVMGQNFALNMAEHGFTVAVCNRS PDKVDDTVERA<br>LAVMGQNLILNAADHGFTVCAVYNRIT QSKVDHFLANE<br>LAVMGSNLARNF AHHKGTVAVFNRIS FEKTOALMDQH<br>MAVMGKNLALNVE SRGYTVAIYNRTIT SKTEEVYKEH<br>LAVMGKNLALNIE SRGF SVSVYNNRS SSKTEEFLOEA<br>MAVMGRNLALNIE SRGYTVEIYNNRS SEKTEEVTAEN<br>LAVMGQNLALNI ARKGYKSVYNNRTAQRTTEFVKNR<br>LGRMGLNLALNMLDHGHEVRGYARTKATVDKAAAQG |
| * <i>Morella thermoacetica</i> (mutant)<br><i>Haloflexax volcanii</i><br><i>Gluconobacter oxydans</i>                                                                                                                                                                                                  | <b>NAD<sup>+</sup> preference 6PGDH</b><br>LGRMGLNLALNMLDHGHEVRGYDIIKATV----DKAAAQG<br>LGRMGRIVVDRVLDAGHEVVAFDLSAEAV----AAAADAG<br>LGRMGGINAVRLTRHGHDVVVDRITSEVTTSVVGRCEAGR                                                                                                                                                                                                                                                        |

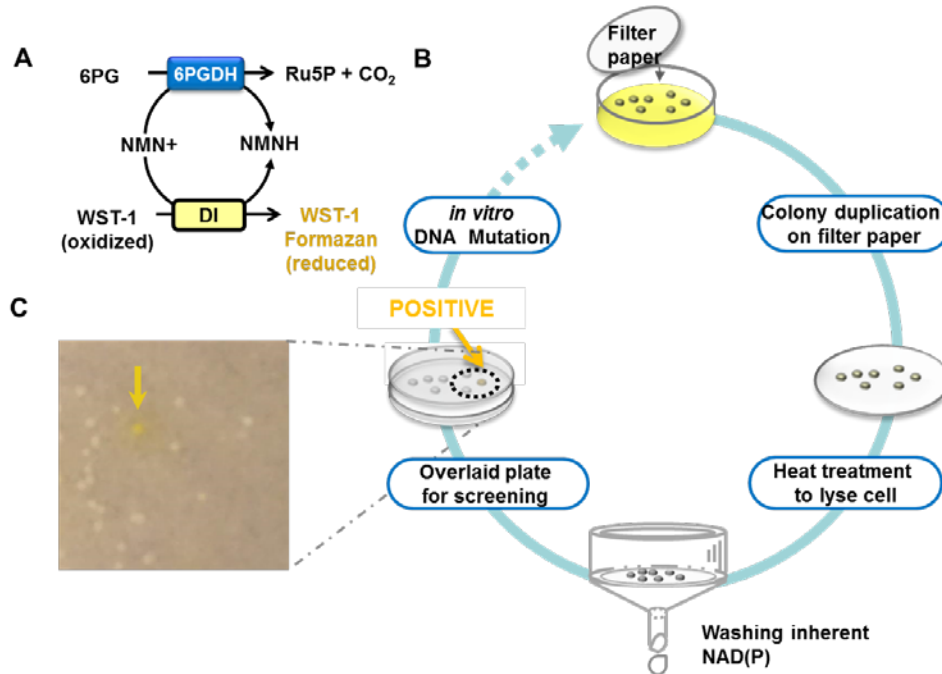
### B) Phosphate group recognition residues analysis



- Via rational design (amino acid alignment and homologous molecule structure), we identified the three amino acids binding to the phosphate group of NADP.
- The best mutant TmG6PDH (N32E/R33I/T34I) exhibited a  $6.4 \times 10^4$ -fold reversal of the coenzyme selectivity from NADP<sup>+</sup> to NAD<sup>+</sup>.

# Progress: Coenzyme engineering toward biomimics

## Novel screening method with minimal interference from intracellular NAD(P)

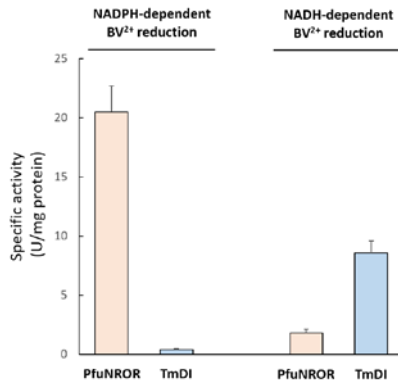
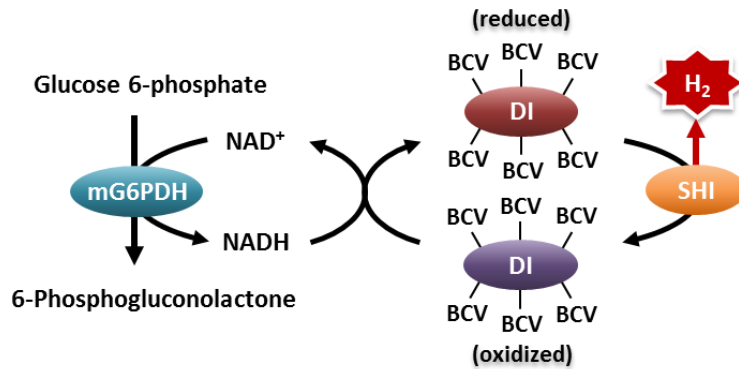


- We conducted three-round site-directed mutagenesis and one-round whole gene sequence mutagenesis, yield the best mutant Tm6PGDH having a specific activity of more than 1 U/mg on NMN.
- The best mutant has up to 10 amino acid changes (e.g., A11G/R33I/T34I/D81I/T82I/Q86I/D294V/Y383C/N387S/A447V).
- Met one of milestones of coenzyme engineering (> 1 U/mg on NMR).

- We are repeating our successful efforts in the other dehydrogenase – TmG6PDH.

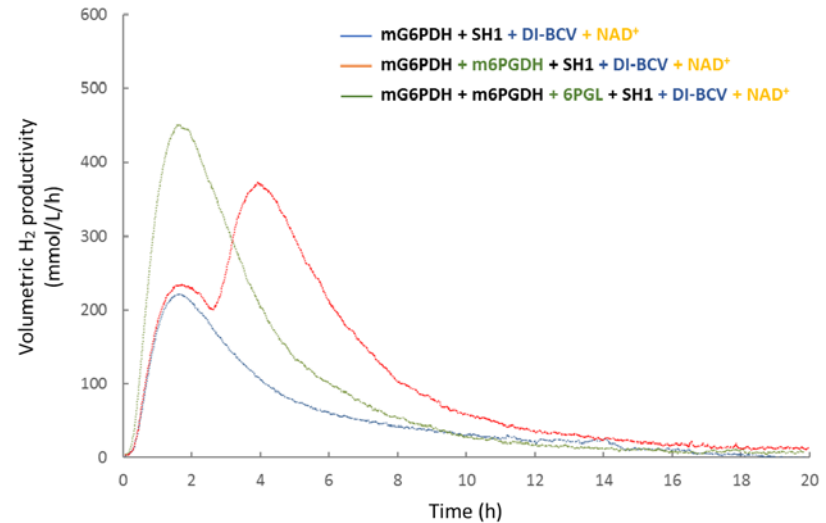
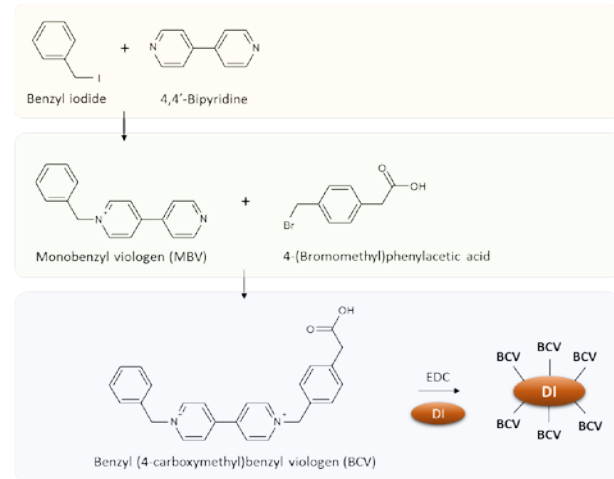
# Achievement: First NAD-based pentose phosphate pathway

## NAD-based PPP + conjugate 2017



Discover a new diaphorase (instead of NROR) for NAD-based ETC, increasing Rxn by 5-fold

## Making DI-BCV conjugate



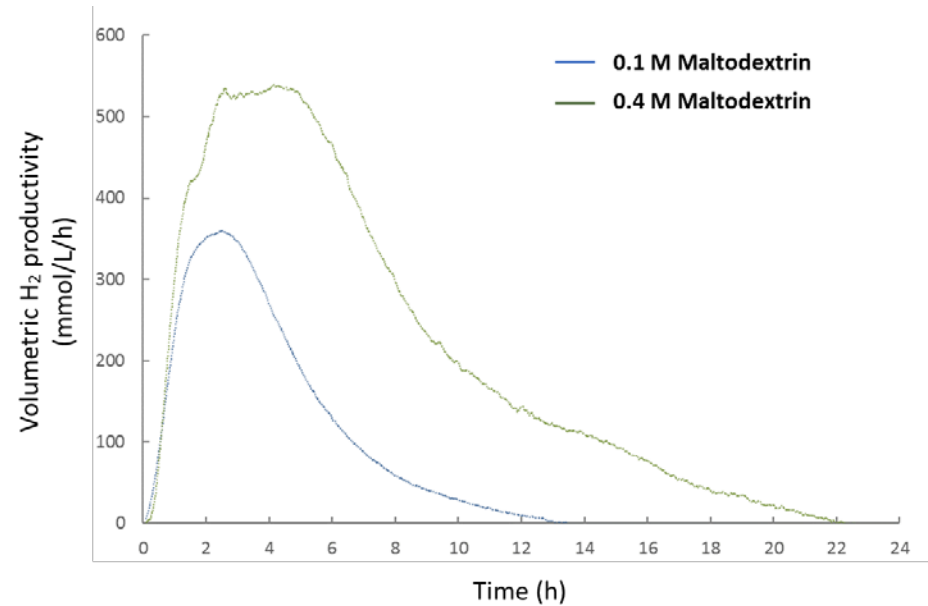
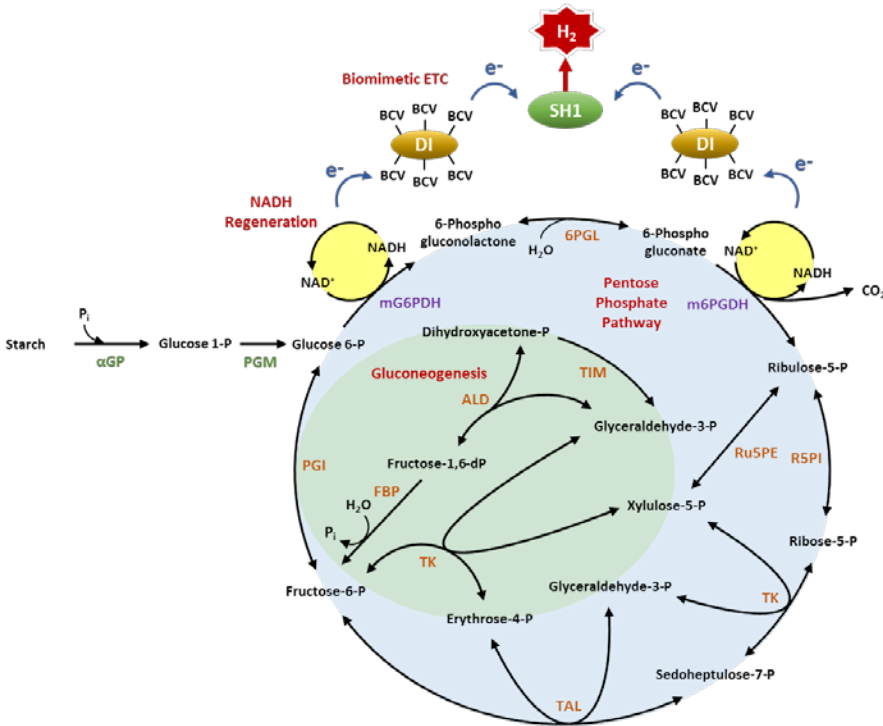
Doubling Rxn by 2-fold by using DI-BV conjugate.



# Achievements: Increasing Rxn to 1 g H<sub>2</sub>/L/h

Revised pathway based on NAD and BV-DI conjugate

Ultra-rapid in vitro H<sub>2</sub> generation

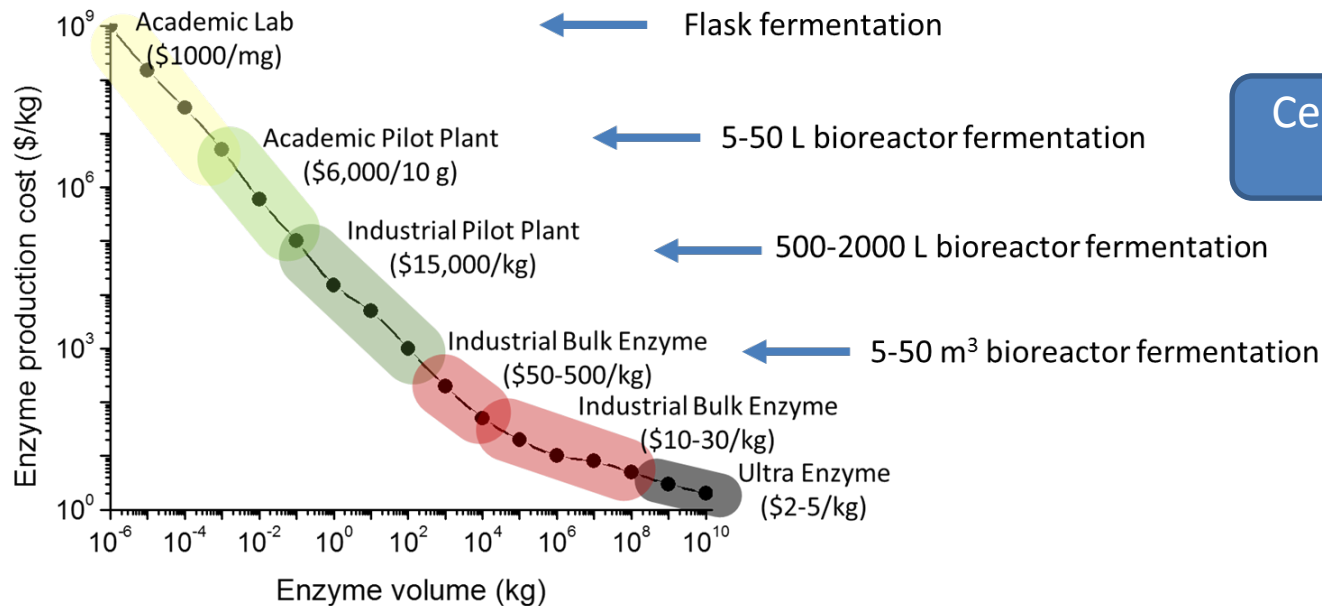


Conditions: 80°C, pH 7.5

- Maximum H<sub>2</sub> generation rate = 530 mmole H<sub>2</sub>/L/h = 1.06 g H<sub>2</sub>/L/h  
= 272 L of H<sub>2</sub>/L/d = ~8 g glucose consumption/L/h
- A plant producing 1,500 kg H<sub>2</sub>/day = ~70 m<sup>3</sup> (anaerobic digester, beer fermenters)

# Progress: Scale-up of high density *E. coli* fermentation

## Comparison of academic & industrial enzyme production cost



## Thermoenzyme prod. & purif.

High-density fermentation  
OD = 150-200, protein  
expression 20-30%

Cell lysis by heat treatment  
(~80 °C, 30 min)

Centrifugation

Ultra-filtration

Recombinant  
Enzymes  
(\$50/kg protein)

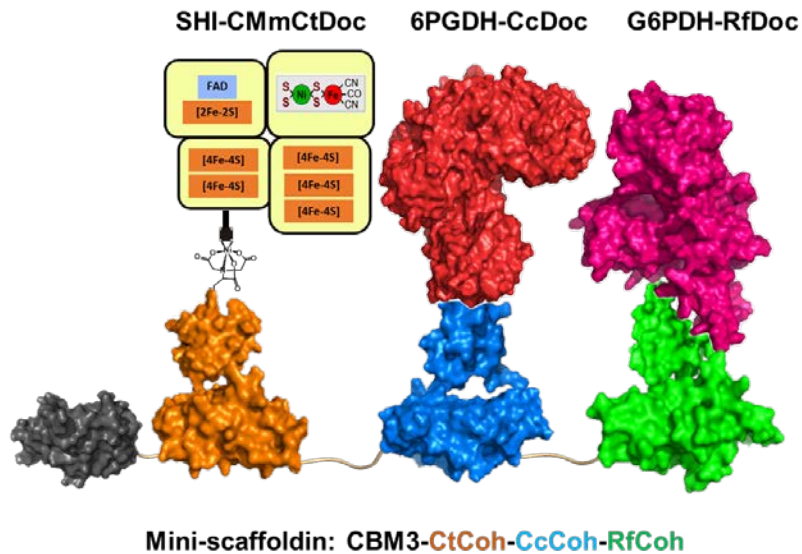
Enzyme costs are NOT a showstopper for *in vitro* synthetic biosystems.

Zhang et al. 2017. *Biomanufacturing: History and Perspective*. *J. Ind. Microbiol. Biotechnol.* 44: 773-786.

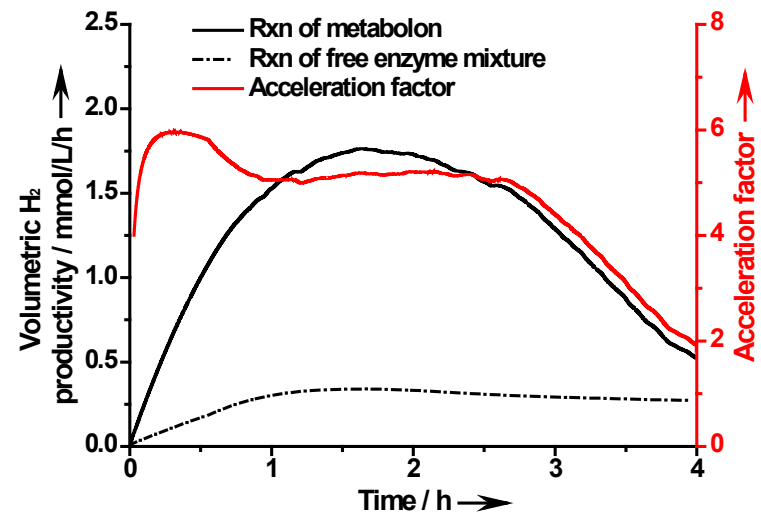
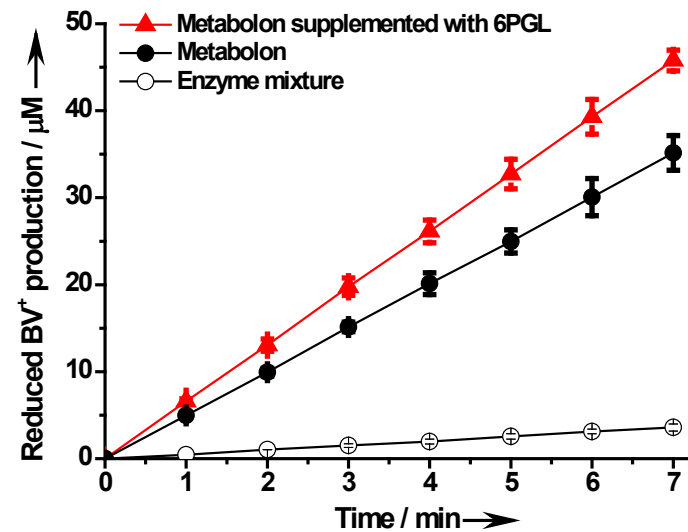
You et al. 2017. *An in vitro synthetic biology platform for the industrial biomanufacturing of myo-inositol from starch*. *Biotechnol. Bioeng.*:DOI: 10.1002/bit.26314.

# Achievements: Electron channeling in metabolon

## Self-assembling metabolon



- The metabolon exhibited an initial reaction rate 12.3 times that of the enzyme mixture based on reduction of oxidized benzyl viologen.
- The hydrogen generation rate catalyzed by the metabolon was approximately 5.6 times of that of the enzyme mixture.
- Such reaction rate enhancements suggested strong electron channelling among the adjacent redox enzymes of the metabolon.



# Progress: More metabolons constructed

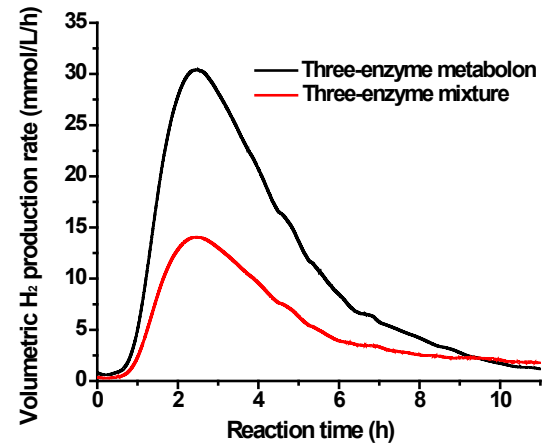
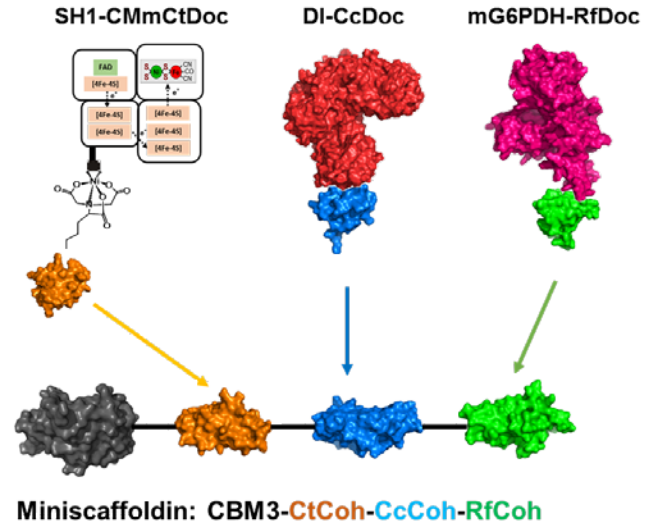
## Dehydrogenase-NAD conjugates



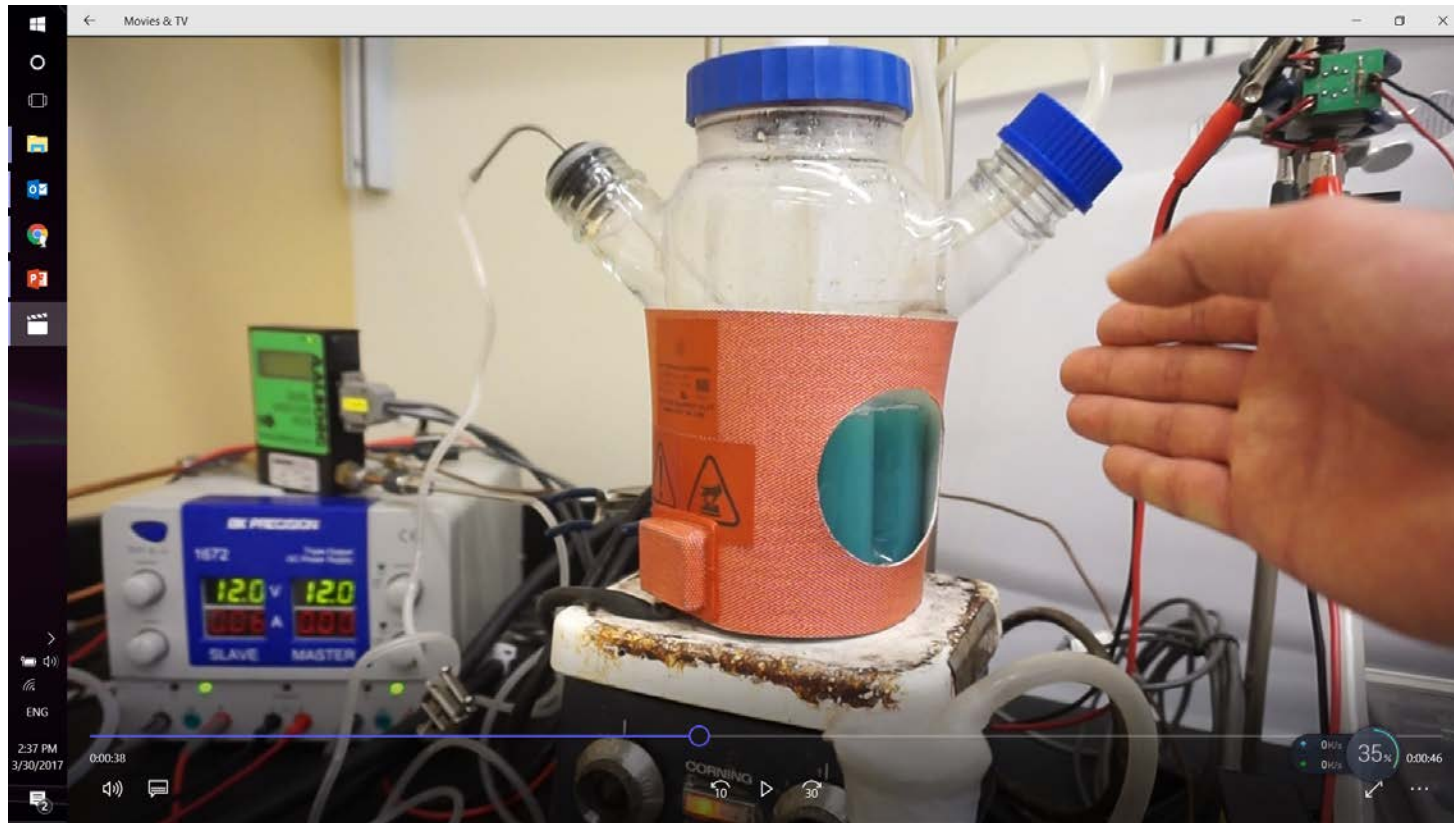
- Facilitate rapid electron transfer by 2-3 fold
- Decrease NAD use
- Stabilize NAD by 20-fold than free NAD.

- Increase H<sub>2</sub> generation rate by ~2-3 fold
- Stability of metabolon at high temperature needs to be addressed
- A possible solution is chemical cross-linking.

## Metabolon featuring ETC



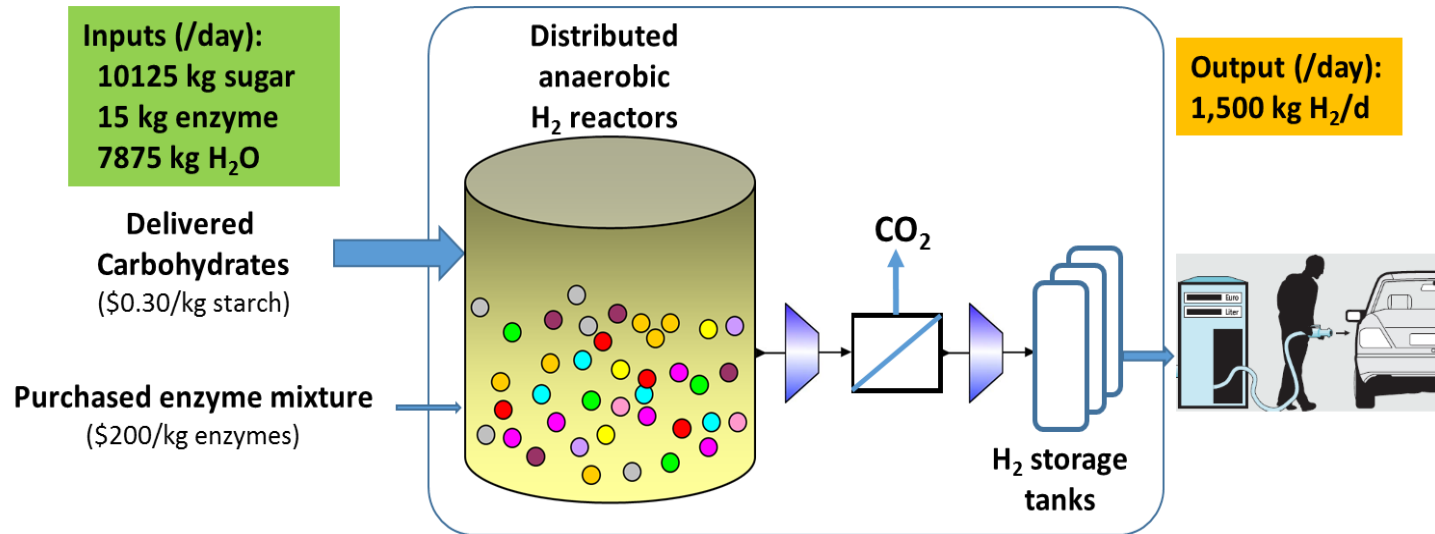
# Progress: 1-liter H<sub>2</sub> dry run production



- We have finished 10-mL experiment in June 2016, meeting the Go/No-Go milestones of Phase I.
- We have tested 1-L dry run H<sub>2</sub> production as shown above (at 80°C).
- We will conduct 1-L demonstration by integrating latest enzyme complexes and new SH1 by Dec. 2017.

# Progress: H<sub>2</sub> production cost analysis

## Distributed green hydrogen production stations (1500 kg H<sub>2</sub>/day)



### Key inputs

- (1) 0.30/kg starch (\$0.20/kg in the future)
- (2) Enzyme cocktail = ~\$200/kg
- (3) 1 kg of enzyme (TTN) = 100 kg of H<sub>2</sub>
- (4) Coenzyme (\$/kg) = 500/kg NADP; 1500/kg NAD; \$500/kg NMN; and \$250/NR
- (5) Coenzymes' total turn-over numbers: 20 k for NADP & NAD; 400 k for NAD-conjugate, NMN & NR

**Cost (/kg H<sub>2</sub>) = starch (\$2.025) + enzymes (\$2.00) + coenzyme (not including CapEx & OpEx)**

| Coenzyme      | Cost (\$/kg H <sub>2</sub> ) |
|---------------|------------------------------|
| NADP          | \$87.77                      |
| NAD           | \$28.90                      |
| NAD-conjugate | \$ 5.35                      |
| NMN           | \$ 4.23                      |
| NR            | \$ 4.09                      |

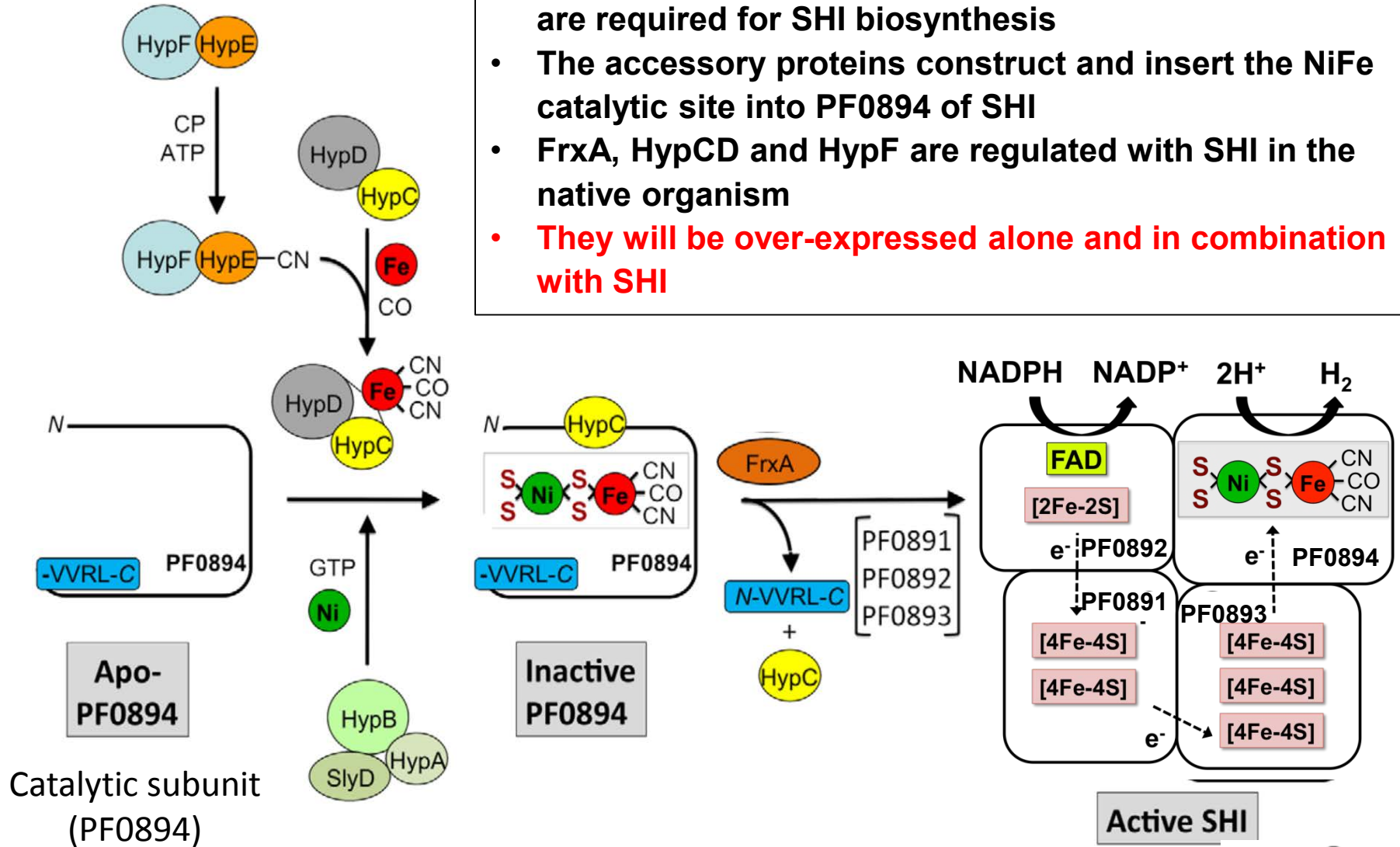


## Collaboration

- Subcontract – Co-PI (Prof. Mike Adams) at University of Georgia
- Focus on low-cost mass production of hyperthermophilic Fe-Ni soluble hydrogenase I (*P. furiosus* SH1) without a discount of specific activity of SH1.
- Related to **Objective 1** (decrease hydrogen production costs for enzymes) and **Objective 3** (scale-up of enzymatic hydrogen production)

# Biosynthesis of *P. furiosus* SHI is Complex and Requires Eight Accessory Genes

- Eight accessory genes (*hypABCDEFGF*, *frxA* and *slyD*) are required for SHI biosynthesis
- The accessory proteins construct and insert the NiFe catalytic site into PF0894 of SHI
- *FrxA*, *HypCD* and *HypF* are regulated with SHI in the native organism
- **They will be over-expressed alone and in combination with SHI**





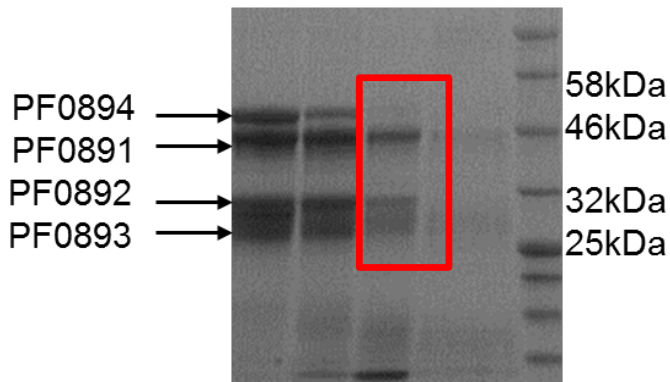
## Scale-up: SHI purification from 500 grams of cells

| Step    | Total Units*<br>( $\mu\text{mol min}^{-1}$ ) | Total Protein<br>(mg) | Specific Activity<br>(U/mg) | Yield<br>% | Fold<br>Purification |
|---------|----------------------------------------------|-----------------------|-----------------------------|------------|----------------------|
| S80     | 85,807                                       | 36,180                | 2.37                        | 100        | 1                    |
| Ni-NTA  | 38,147                                       | 680                   | 56.1                        | 44         | 11                   |
| Q HP E1 | 24,301                                       | 188                   | 128.9                       | 28.3       | 54.4                 |
| Q HP E2 | 10,913                                       | 136                   | 80.1                        | 12.7       | 33.8                 |
| Q HP E3 | 1,713                                        | 70                    | 24.5                        | 2          | 10.3                 |
| Q HP E4 | 557                                          | 20                    | 27.8                        | 0.6        | 11.7                 |

\*Based on MV-linked H<sub>2</sub> evolution assay

### QHP

E1 E2 E3 E4



**324 mg of SHI was purified**

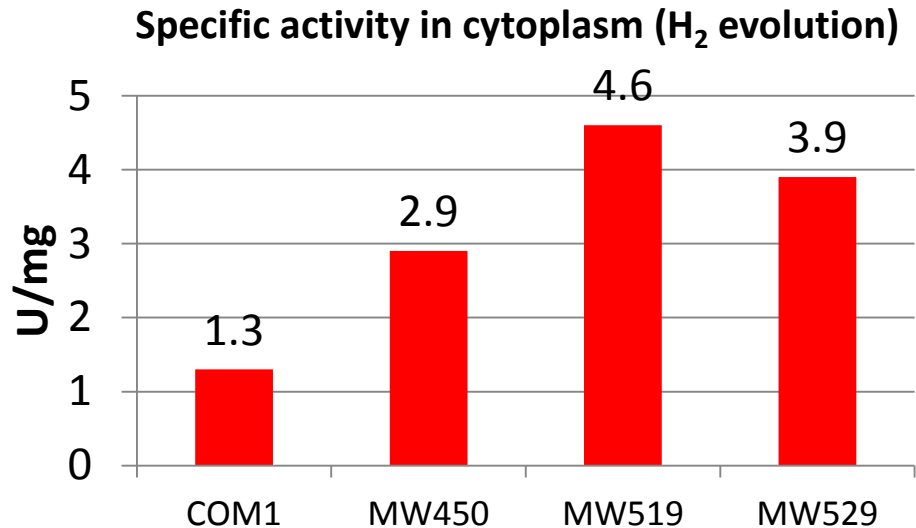
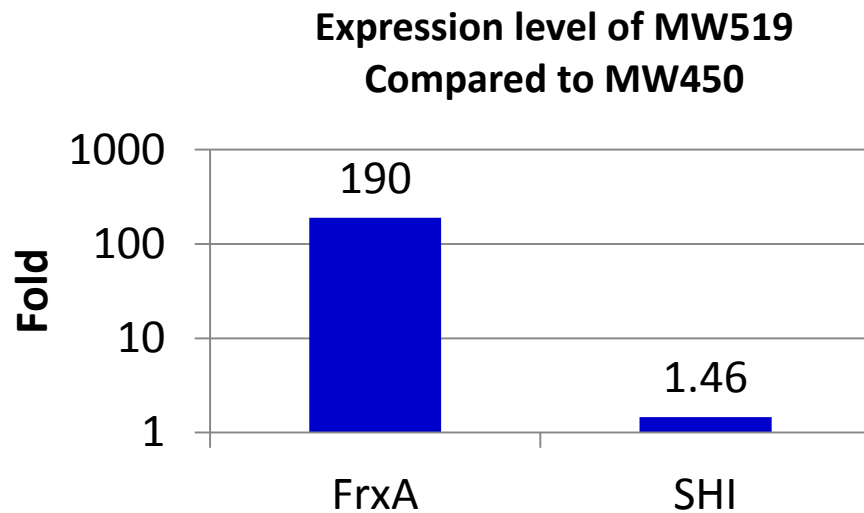
**Problem: >30% of the SHI protein lacks the PF0894 catalytic subunit (E3)**

**Assumed to be degraded as it lacks the NiFe catalytic site**

**Goal: to over-express the accessory proteins for SHI maturation to solve the problem**

## Over-expression of three SHI accessory proteins

Three accessory proteins (FrxA, HypC and HypD) necessary to synthesize the catalytic (NiFe) site of SHI were overexpressed



COM1: Original parent strain

MW450: OE 9xHis tagged SHI

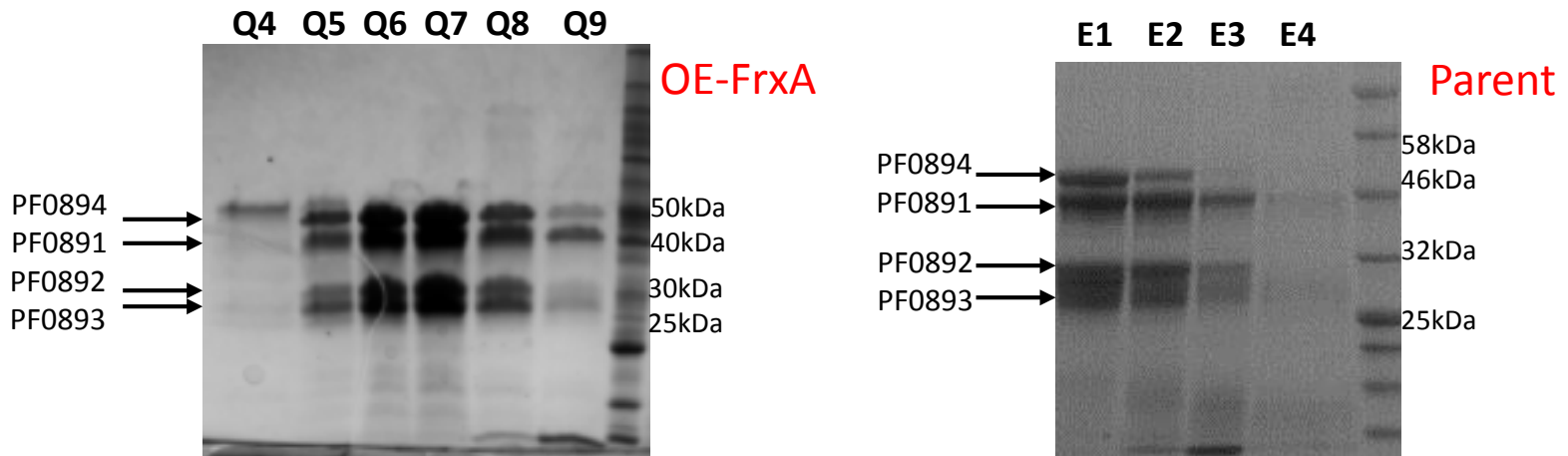
MW519: OE 9xHis tagged SHI plus OE-FrxA

MW529: OE 9xHis tagged SHI plus OE-FrxA/OE-HypCD

- Over-expression of *frxA* increased FrxA levels by almost 200-fold (strain MW519)
- **Over-production of FrxA resulted in three-fold higher SHI activity (H<sub>2</sub> evolution)**
- Over-production of HypC/D in the FrxA over-expression strain did not further increase SHI production

# SHI purification from cells over-expressing *FrxA*

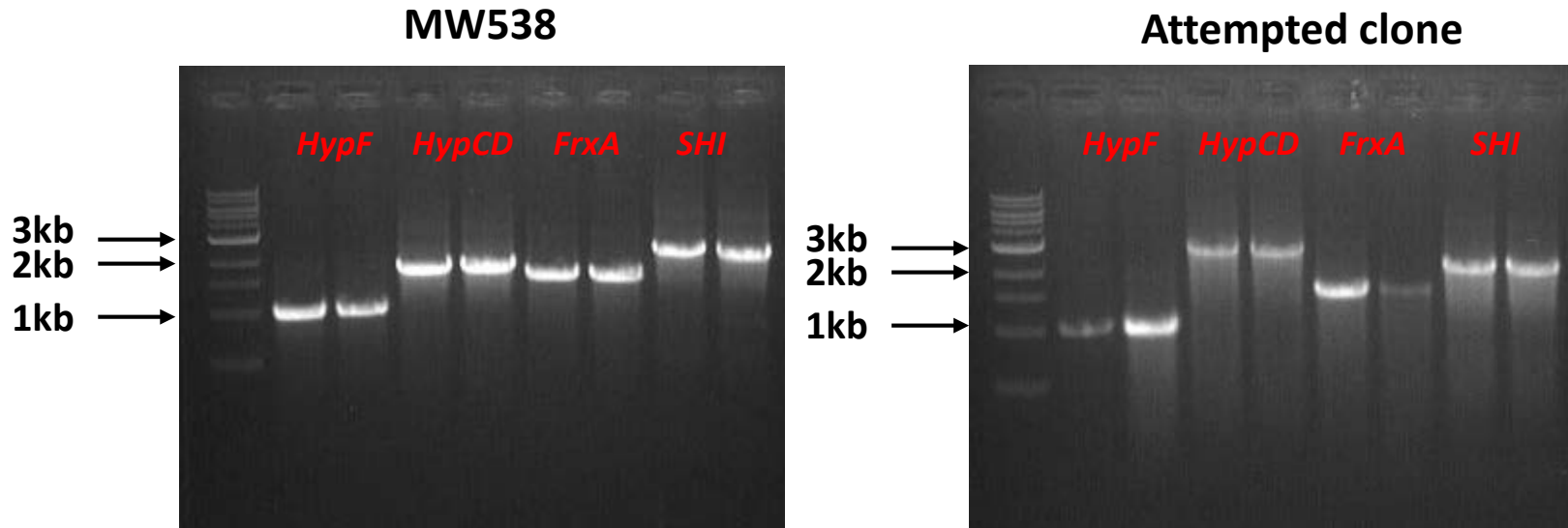
| Step        | Total Units*<br>( $\mu\text{mol min}^{-1}$ ) | Total Protein<br>(mg) | Specific Activity<br>(U/mg) | %<br>Yield | Fold<br>Purification |
|-------------|----------------------------------------------|-----------------------|-----------------------------|------------|----------------------|
| S100        | 4328.4                                       | 1334                  | 3.24                        | 100        | 1                    |
| Ni-NTA      | 3041.8                                       | 28                    | 108.8                       | 70.3       | 33.58                |
| Q HP (Q5)   | 115.8                                        | 0.75                  | 155.5                       | 2.7        | 48                   |
| Q HP (Q6-8) | 2265.3                                       | 14                    | 156                         | 52.3       | 48                   |
| QHP (Q9)    | 115.9                                        | 1.5                   | 77.7                        | 2.7        | 24                   |



- >30% trimer was observed in previous SHI prep (E3, 20U/mg)
- <10% of total SHI is the trimer in the OE-FrxA strain(MW519) SHI
- The yield from a 20L fermentation was low (20g) compared to MW430 (40g)
- **Over expressing *FrxA* dramatically reduces the amount of inactive trimeric SHI**

# Over-expression of processing protein *HypF*

## PCR confirmation of gene loci in strains

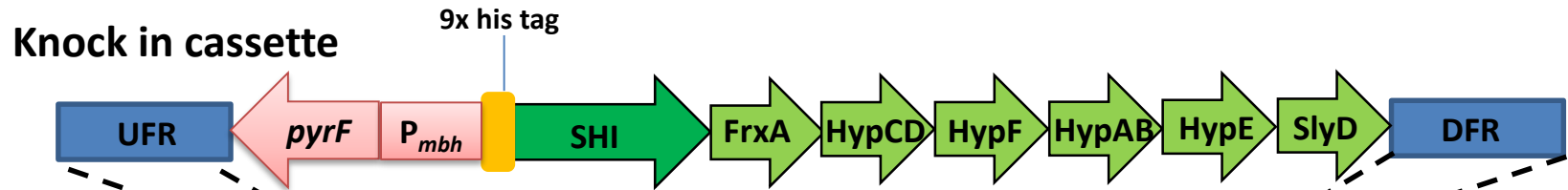


MW538: OE 9xHis tagged SHI plus OE-FrxA/OE-HypCD, along with *pyrF* recycled

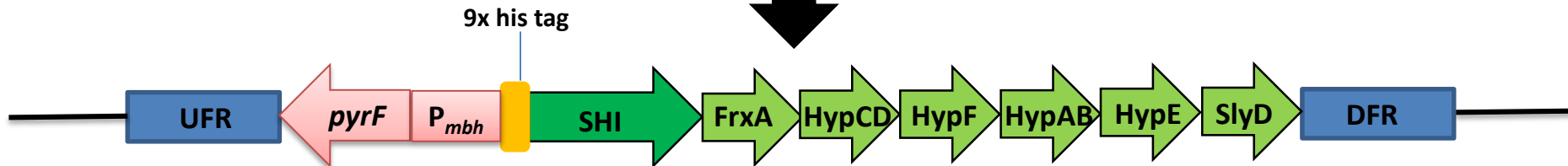
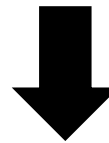
- Hundreds of colonies were screened for *HypF* over-expression
- Unable to purify a clean colony of MW538 for *HypF* over-expression, even after 5 rounds of plate purification
- The attempted clone still has *pyrF* selection marker in *HypCD* locus
- Over-expression cassette was not transformed into *HypF* locus
- ***HypF* cannot be over-expressed without deleterious effects**

# New strategy for over-expression of SHI and all 8 maturation genes

All eight maturation genes expressed in addition to a second copy of the SHI genes



Intergenic space in Pf genome

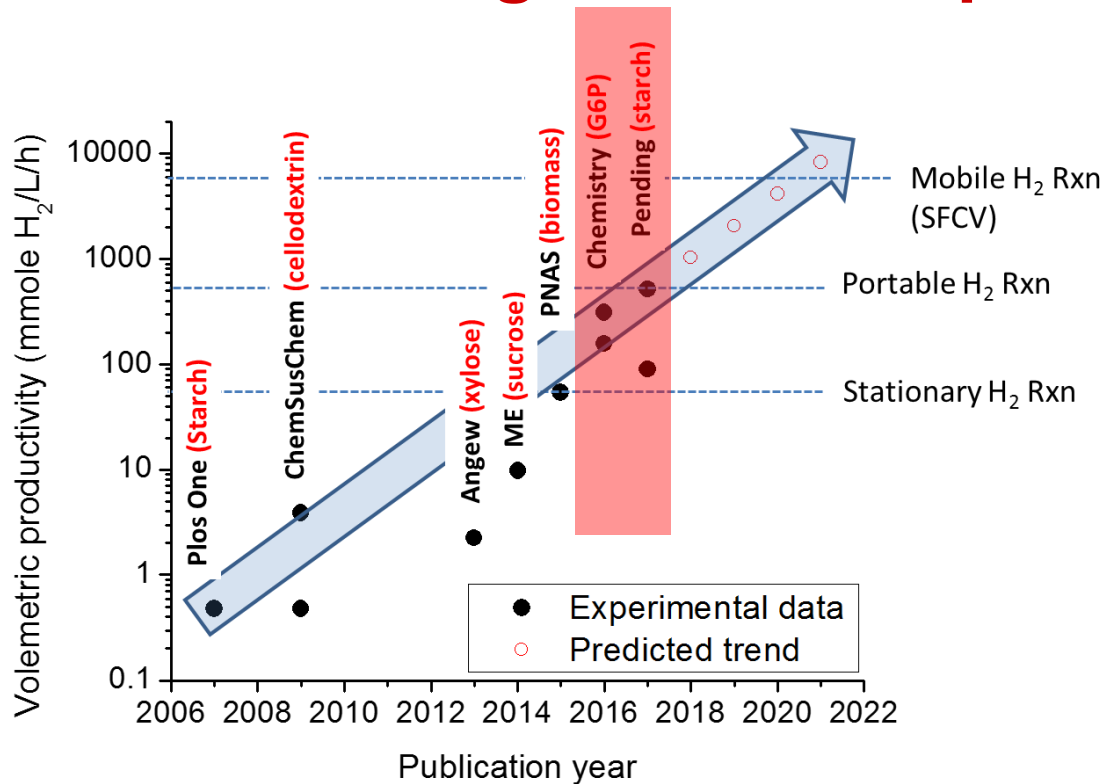


- A second copy of SHI and of its 8 maturation genes will be expressed at an intergenic space in the existing OESHI strain
- Genes expression controlled by the  $P_{mbh}$  promoter
- **Characterization of the new recombinant strain in progress**

# Remaining Challenges and Barriers

- Increase catalytic efficiencies ( $k_{cat}/k_M$ ) dehydrogenases on biomimics (NMN and NR) to comparative levels of to those on their natural coenzyme (NADP).
- Stabilize coenzymes for a long time running (partially addressed (via NAD-conjugate) here, future).
- Decrease SH1 production costs – better expression levels without a decrease in specific activity (i.e., coordinated co-expression of enzyme components, no-cost extension efforts).
- Construct very stable G6PDH-6PGDH-SH1 enzyme complexes featuring electron transport channeling.
- Scale up recombinant enzyme production to kg levels for pilot plant demonstration (need external support)

# Increasing volumetric productivity (Rxn)

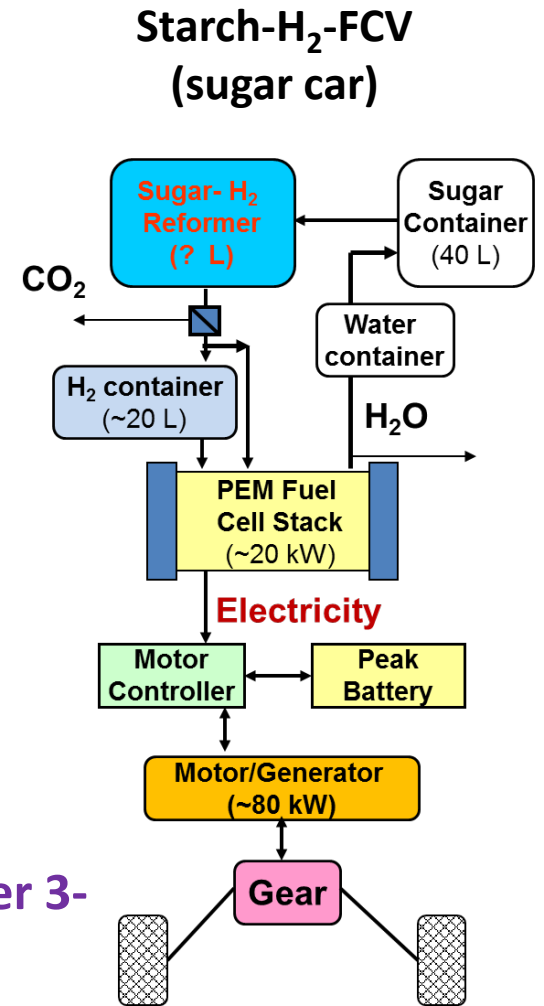


**Zhang's law – Doubling Rxn every year**  
**(increasing 1,000 fold, Rxn = 1 g/L/h)**

**Goal: On-board H<sub>2</sub> generation @ Rxn = 10 g/L/h (another 3-5 year efforts)**

## Remaining barriers

1. Concept (in vitro systems are too costly?)
2. Impacts (transportation revolution?)



Zhang. 2009. Is the sugar-powered car science fiction? *Energy Environ. Sci.* 2:272-282.

# Proposed future work (by Dec. 2017)

- **Decrease H<sub>2</sub> production costs by 1000 fold to \$10/kg, where production cost will be estimated using H2A model**
  - Task 1.2. Replace costly NADP by biomimic – NMN (cont'd)
  - Task 1.4. Detailed economic analysis of H<sub>2</sub> production (cont'd)
- **Increase H<sub>2</sub> production rate from 300 to 750 mmole H<sub>2</sub>/L/h**
  - Task 2.3. Construction of artificial electron transport chains
  - Task 2.4. Construction of synthetic metabolons (enzyme complexes)
- **Scale up H<sub>2</sub> reaction volume from 10 mL to 1000 mL**
  - Task 3.2. Mass production of hydrogenase (SH1) (UGa)
  - Task 3.3. Liter level demonstration (integrated work)



# Technology transfer activities

- Virginia Tech and Oak Ridge National laboratory received a US patent US 8,211,681. Biohydrogen production by an artificial enzymatic pathway. (2012).
- We are willing to provide technical help to any entities to commercialize the sugar-to-hydrogen technology and develop the hypothetical sugar-H<sub>2</sub>-fuel cell vehicles.
- No special business action taken due to a lack of hands and experienced businessmen.
- (Some Chinese research institutes and companies express strong interests in this project and the *in vitro* synthetic biology platform because they know enzymes better than US counterparts. China has established the first plant based on this *in vitro* platform for the production of value-added product – myo-inositol. )

# Summary

- We achieved the highest biological hydrogen generation rate of 550 mmole H<sub>2</sub>/L/h.
- We demonstrated the feasibility of changing coenzyme preference of engineered dehydrogenases from NADP to NAD and biomimics.
- Starch is an off-board H<sub>2</sub> storage compound (i.e., 14.8 H<sub>2</sub> wt. %) and could be a on-board H<sub>2</sub> storage compound.
- We scaled up recombinant *E. coli* enzyme production by 1000-fold and recombinant *P. furiosus* SH1 (hydrogenase) production by 50-fold.

| Targets         | Units                     | June 2016 Target   | Dec. 2017 Target (estimated) | Year 2020 Target (Plant gate)    |
|-----------------|---------------------------|--------------------|------------------------------|----------------------------------|
| Production cost | \$/kg H <sub>2</sub>      | 1000               | 10                           | 10 (year 2020)                   |
| Productivity    | mmole H <sub>2</sub> /L/h | 550 (achieved)     | 750                          | 2,000                            |
| Reactor volume  | L of reactor              | 1 (to be finished) | 1                            | 65,000*                          |
|                 |                           |                    |                              | *1,500 kg H <sub>2</sub> per day |

# Technical Back-Up Slides

# Appraisal of enzymes as biocatalysts

## Basic facts

- Most enzymes are proteins
- Biocatalysis catalyzed by enzymes has highly chemical selectivity (no side product)
- Most enzymes work at mild reaction conditions (low temperature, 1 atm, neutral pH, and aqueous phase)
- Enzymes do not require costly precise metals

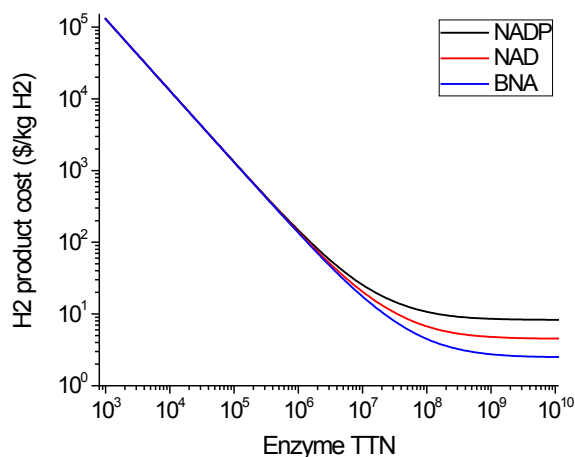
## Conflicting concepts (academic researchers versus industrial enzyme experts)

- × For academic researchers, enzymes are VERY costly (e.g., **billion dollars per kg**)
- √ For industrial enzyme experts, bulk enzymes are less costly (e.g., **10 dollars per kg**)
  
- × For academic researchers, enzymes are very labile, losing activities within hours or days
- √ For industrial enzyme experts, some enzymes (e.g., immobilized or engineered) are very stable, lasting months and years (e.g., glucose isomerase for HFCS production, protease in detergent, glucose dehydrogenase in blood sugar test strips).
  
- × For academic researchers, enzymes have narrow reaction conditions in terms of pH, temperature and solutions.
- √ For industrial enzyme experts, some enzymes (engineered or discovered from extremophiles) can work on a large temperature range from 0 – 100°C, pH from 1 to 14, from aqueous solution to 100% organic solvent

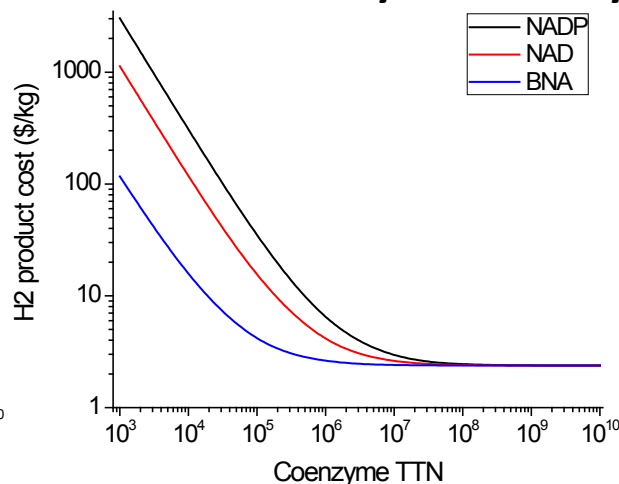
# Decreased H<sub>2</sub> production costs

Quantitative indicator of enzymes and coenzyme: **total turn-over number (TTN)**, mole product/mole enzyme

Effects of enzyme stability



Effects of coenzyme stability



## Preconditions for low-cost hydrogen production

- Low cost enzyme production (\$10-20/kg)
- High stability of enzymes (TTN = 10<sup>8-9</sup> mole product/mole enzyme)
- High stability of coenzymes (TTN = 10<sup>6-7</sup> mole product/mole enzyme)
- Low-cost of coenzymes (\$100/kg)

**Key directions: (1) discovery of better enzymes (Task 1.3), (2) engineering of dehydrogenases on biomimetic cofactors (Task 1.2), and (3) mass production of enzymes (Tasks 3.1 & 3.2).**

## Enzyme stability

| EC       | Enzyme name                    | Source                 | Form           | Cond.   | TTN                                            |
|----------|--------------------------------|------------------------|----------------|---------|------------------------------------------------|
| 1.1.1.44 | 6-phosphogluconate hydrogenase | <i>T. maritima</i>     | Free           | 80°C    | 2.4 x 10 <sup>8</sup>                          |
| 2.2.1.2  | Transaldolase                  | <i>T. maritima</i>     | Free           | 60°C    | 1.7 x 10 <sup>7</sup>                          |
| 3.1.3.11 | Fructose 1,6-Bisphosphatase    | <i>T. maritima</i>     | Free           | 60°C    | 2 x 10 <sup>7</sup>                            |
| 5.4.2.2  | Phosphoglucomutase             | <i>C. thermocellum</i> | Free           | 60°C    | 7.1 x 10 <sup>7</sup>                          |
| 5.3.1.5  | Xylose (glucose) isomerase     |                        | Immobilized    | 50-60°C | 5.0 x 10 <sup>8</sup>                          |
| 5.3.1.6  | Ribose-5-phosphate isomerase   | <i>T. maritima</i>     | Free           | 60-70°C | 2.2 x 10 <sup>8</sup>                          |
| 5.3.1.9  | Phosphoglucose isomerase       | <i>C. thermocellum</i> | Free<br>Immob. | 60°C    | 3.2 x 10 <sup>7</sup><br>1.1 x 10 <sup>9</sup> |

### Our goal:

1 kg of enzyme produces 300 kg of H<sub>2</sub> @ TTN = 4 E7 → \$0.10/kg H<sub>2</sub>