







# A genome scale model of *Cupriavidus necator* for 3-HP production

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# **Gas fermentation for platform chemical production**



3-HYDROXY-PROPRIONIC ACID





# **Gas fermentation for platform chemical production**







### **3-HP production – first intermediate chemical**



- > Important building block for **biorenewable** polymers
- Chemical synthesis is not commercially feasible due to low yield and high production cost





## C. necator an ideal chassis for biotechnology

- Facultatively chemolithoautotrophic bacteria - grow with organic substrates or H<sub>2</sub> and CO<sub>2</sub> under aerobic conditions
- Grow to high-cell densities under lithoautotrophic or heterotrophic conditions
- Produces large amounts of a biodegradable polymer polyhydroxybutyrate (PHB)



Flagellation of strain N-1. Bar, 1.0 µm



STEM picture of *Cupriavidus necator* harbouring PHB granules





### C. necator lithoautotrophic metabolism

- Carbon dioxide is fixed via the Calvin cycle
- Membrane bound hydrogenase directly connected to the electron transport chain (ETC) for generating ATP
- Soluble hydrogenase that is coupled to NADH synthesis that is required for the Calvin cycle or ETC
- Oxygen final electron acceptor (under anaerobic conditions nitrate is used)



Lithoautotrophic metabolism





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Lithoautotrophic metabolism





# Metabolic modelling of C. necator for 3-HP production

#### Assess 3-HP pathways

- Product yield
- Oxygen requirements
- ATP yield
- Simulate behaviour of bacteria for
  - insertion/knockouts of reactions
  - varying feeding ratios
- Predict metabolic interventions for increasing 3-HP production





#### **Assessing 3-HP pathways using EMA**



Computed **elementary modes** of a small model of *C. necator* to assess two **3-HP pathways** 

- > Calvin cycle (13)
- Electron transport chain (6)
- > 3-HP pathways (13)
- > PHB pathway (3)





## **Assessing 3-HP pathways using EMA**

Pathway	Theoretical max. Yield (mol/mol co <sub>2</sub> )	O <sub>2</sub> requirement (mol/mol product)	H <sub>2</sub> requirement (mol/mol product)	No. of gene additions
РНВ	1.0	[5.57, 6.5]	[20.14, 22]	0
3-HP I. BAPAT	1.0	[3.0, 3.66]	[12.0, 13.33]	0
3-HP II. MCR	1.0	[3.27, 3.83]	[12.28, 13.66]	1

Net stoichiometry of example elementary mode:

 $3 \text{ CO}_2 + 12 \text{ H}_2 + 3 \text{ O}_2 \longrightarrow 1 3 \text{-HP} + 1 \text{ H}_2\text{O} + \text{H}^+$ 





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Deletion of membrane hydrogenase removes all nonoptimal modes in terms of oxygen requirements





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#### Genome scale model of *C. necator*

- GSM construction based on BioCyc database for C. necator H16
- Model constructed as set of modules:
  - BioCyc reactions (1101)
  - Transport reactions (58)
  - Electron transport chain (10)
  - Additional reactions (152)
- Minimal aerobic media: fructose, oxygen, sulfate, phosphate and ammonium

Final model consists of 1301 reactions and 1200 metabolites





## Simulating reaction knockouts

















### **3-HP growth coupling strategy 1: results**







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Chorismate

## **3-HP growth coupling strategy 1: results**

GLN Simulated single knockouts Pyruvate on each reaction in model PRPP α-L-alanine Anthranilate PPI Kyrurenine Identified **7 candidate** N-5-phosphoribosyl-• Formate anthranilate reactions whose deletion resulted in 3-HP production N-formylkynurenine Carboxyphenylaminodeoxyribulose Highest 3-HP flux was **0.98**, with 0.05 carbon yield Trptophan  $CO_2$ Indole-3-glycerol-p GAP Biomass Serine





























## **Production envelopes to assess growth coupling**

maximise or minimise  $v_{3-\text{HP}}$ subject to :  $\begin{cases} N\mathbf{v} = 0 \\ v_j = t_j \\ v_{\text{ATPase}} = J_{\text{ATPase}} \\ v_{\text{ko}} = 0 \\ 0 \le v_{\text{fru}} \le 8.8 \end{cases} \longleftarrow \text{Constrain maximum}$ 

LP resolved for increasing growth rates for both minimising and maximising 3-HP flux





#### **Production envelopes to assess growth coupling**







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#### Flux varied analysis to assess growth coupling







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# **3-HP scan analysis**

minimise : 
$$|\mathbf{v}|$$
  
subject to : 
$$\begin{cases} Nv = 0\\ v_j = t_j;\\ v_{\text{ATPase}} = J_{\text{ATPase}}\\ v_{\text{ko}} = 0\\ v_{\text{3HP}} = k \end{cases}$$

- LP resolved for increasing values of 3-HP flux
- Identify reactions that respond to increase 3-HP, fluxes that decrease are candidate knockouts





## **3-HP scan analysis**



Simulate knockout combinations of the reactions that decreased as 3-HP increased





# **3-HP scan analysis: results**

#### We identified 1 reaction that increased 3-HP carbon yield from 0.24 to 0.30







### **Conclusions & Outlook**

- Example of biology and modelling working together:
  - Using model to assess knock out strategy, and then to identify further knock outs that coupled 3-HP to essential biomass precursors
  - Using experimental data to refine model test the predictions in the lab





### **Conclusions & Outlook**

- Example of biology and modelling working together:
  - Using model to assess knock out strategy, and then to identify further knock outs that coupled 3-HP to essential biomass precursors
  - Using experimental data to refine model test the predictions in the lab
- Future work:
  - Implement MOMA (minimisation of metabolic adjustments) for simulating knockouts
  - Implement techniques such as OptGene, for predicting knockouts





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