

L15 Metabolic Engineering

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Recall...

Parts assembled via variety of techniques

Can be delivered via plasmid or genome

Genomes have properties that are critical for function

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This lecture

- What is metabolism?
- Why metabolic engineering?
- Review Michaelis-Menten Kinetics
- Metabolic Engineering Strategies

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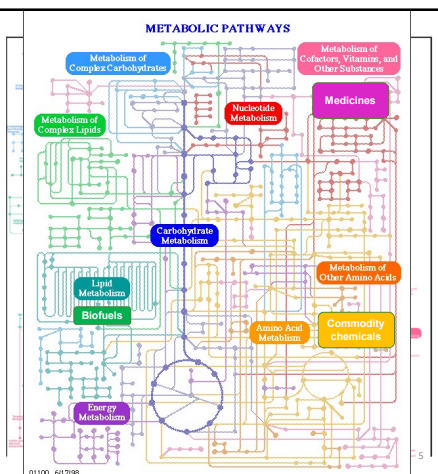
What is metabolism?

Metabolism is the network of reactions cells do to:

- Generate energy
- Create building blocks of biomass
- Protect itself from the environment and compete

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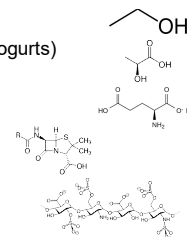
What is metabolism?



Metabolism

Examples of molecules possible from biology:

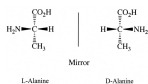
- Ethanol (yeast in beer/wine)
- Lactic acid (*Lactobacillus* in yogurts)
- MSG/flavors (*C. glutamicum*)
- Penicillin
- Heparin (anticoagulant)
- Proteins
- methane



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Microbial metabolism

- Microbes make diverse products
- Products are stereospecific
 - ex: L-amino acids only, no D-amino acids
- Use renewable feedstocks

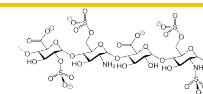


Stereospecificity and wealth of chemistry makes microbes attractive platforms for engineering

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Metabolic engineering vs traditional synthesis

Heparin Synthesis



Chemical Synthesis

- Up to 55 steps
- Non-stereospecific steps result in unwanted products

Enzymatic Synthesis

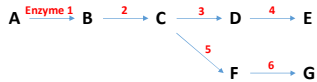
- 4 Enzymatic reactions
- Enzymes are stereospecific

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Metabolic engineering

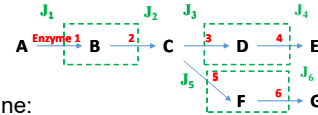
Can represent metabolism with:

- Products formed from metabolic reactions
- Reactions catalyzed by enzymes encoded by genes



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Metabolic engineering



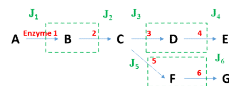
We can define:

- Products/Target Compounds (e.g. E)
- Substrates (reactants) (e.g. A)
- Side Products (e.g. G)
- Intermediates – metabolic nodes between products and substrates (e.g. B, C, D, F)
- Metabolic fluxes (J) – defined as flow of metabolite in or out of arbitrary control volume
 - Typically just reaction rate given by Michaelis-Menten kinetics

*What is the relationship between J_2 and J_1 ?
 J_2 and J_5 ?*

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Metabolic engineering



Goal: Maximize flux towards product

- System is most productive when concentration of intermediates are at steady state

$$\frac{dB}{dt} = \frac{dC}{dt} = \frac{dD}{dt} = \frac{dF}{dt} = 0$$

Why aren't $\frac{dA}{dt}, \frac{dE}{dt}, \frac{dG}{dt} = 0$?

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Metabolic engineering

Mass balances around branch points at steady state are useful

- Balance of C:

$$\begin{aligned} \frac{dC}{dt} &= \text{Influx} - \text{Outflux} \\ &= J_2 - J_3 - J_5 = 0 \\ J_2 &= J_3 + J_5 \end{aligned}$$

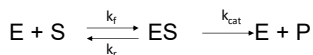
$$J_2 = J_3 + J_5$$

$$\frac{k_{cat2}[E_2]_0[B]}{K_{m2} + [B]} = \frac{k_{cat3}[E_3]_0[C]}{K_{m3} + [C]} + \frac{k_{cat5}[E_5]_0[C]}{K_{m5} + [C]}$$

Fluxes given by Michaelis-Menten kinetics

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Review: Michaelis-Menten Kinetics



E = Enzyme

S = Substrate

ES = Enzyme-substrate complex

k_f = Forward rate constant

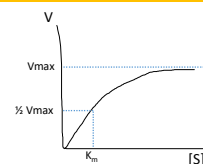
k_r = Reverse rate constant

k_{cat} = Catalytic rate constant (turnover number)
Molecules of S converted to P per second

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Review: Michaelis-Menten Kinetics

$$V = \frac{dP}{dt} = V_{max} \frac{[S]}{K_m + [S]} = k_{cat} [E_0] \frac{[S]}{K_m + [S]}$$



- V = reaction rate
- [S] = concentration of substrate S
- V_{max} = maximum rate achieved by system
 - At saturating substrate concentration
- K_m = Substrate concentration when rate is $\frac{1}{2} V_{max}$

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Review: Michaelis-Menten Kinetics

$$V = k_{cat} [E]_0 \frac{[S]}{K_m + [S]}$$

Reaction order:

[S] \ll K_m : **1st order**

Rate varies linearly with [S]

$$V \approx k_{cat} [E]_0 \frac{[S]}{K_m}$$

[S] \gg K_m : **0th order**

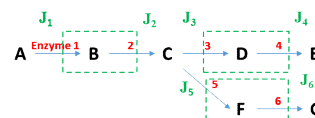
Rate approaches V_{max}

V_{max} = All E bound to S

$$V \approx k_{cat} [E]_0$$

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Example 1



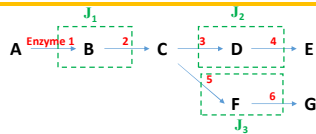
$$K_{m,enz1} = 0.25 \mu M$$

$$[A] = 0.10 \mu M$$

How does one improve production of B?

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Example 1



$$K_{m,enz1} = 0.25 \mu\text{M}$$

$$[A] = 0.10 \mu\text{M}$$

$$V = V_{max} \frac{[A]}{K_m}$$

1st Order

How does one improve production of B?

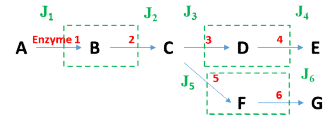
$$[A] < K_m$$

Rate is ~ linear in [A]

Increase [A] to increase rate

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Example 2



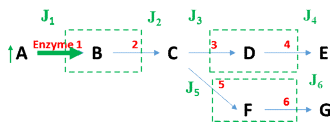
$$K_{m,enz1} = 0.25 \mu\text{M}$$

$$[A] = 5.0 \mu\text{M}$$

How does one improve production of B?

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Optimizing product formation



How do you optimize product formation (E)?

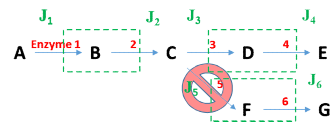
1. Push flux from A

- Increase [A]
- Increase $[E_1]_T$ (upregulate enzyme 1)

$$V = k_{cat}[E_1]_0 \frac{[A]}{K_m + [A]}$$

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Optimizing product formation



How do you optimize product formation (E)?

$$\frac{k_{cat2}[E_2]_T[B]}{K_{m2} + [B]} = \frac{k_{cat3}[E_3]_T[C]}{K_{m3} + [C]} + \frac{k_{cat5}[E_5]_T[C]}{K_{m5} + [C]}$$

2. Pull flux from branch point (C)

- Increase $[E_3]_T$ – overexpression
- Decrease $[E_5]$ – downregulate or knockout

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BRENDA (Braunschweig Enzyme Database)

Kinetic information

K _M VALUE [mM]	SUBSTRATE	ORGANISM	UNIPROT	COMMENTARY	LITERATURE	IMAGE
0.5 - 1.10	2-deoxy-D-glucose	[3] 4 entries				
0.07 - 3.05	ATP	[3] 36 entries				
5	CTP	Homo sapiens	-	[Mg]CTP2-, hexokinase I, at 37°C, pH 8.1	[640237]	
10 - 157.7	D-fructose	[3] 3 entries				
0.5 - 1.5	D-glucosamine	[3] 3 entries				
0.002 - 3.8	D-glucose	[3] 15 entries				
0.07 - 4.6	D-mannose	[3] 3 entries				
16.6	ITP	Homo sapiens	-	[Mg]ITP2-, hexokinase I, at 37°C, pH 8.1	[640237]	
5	UTP	Homo sapiens	-	[Mg]UTP2-, hexokinase I, at 37°C, pH 8.1	[640237]	
additional information		additional information			[640245]	

TURNOVER NUMBER [1/s]	SUBSTRATE	ORGANISM	UNIPROT	COMMENTARY	LITERATURE	IMAGE
29	2-deoxy-D-glucose	Homo sapiens	-	wild-type enzyme	[672188]	
4.92 - 62.3	ATP	[3] 10 entries				
0.00091 - 101	D-glucose	[3] 19 entries				
additional information		additional information				

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BRENDA

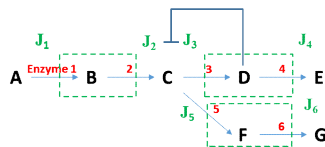
Kinetic information

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0.00091 - 1	D-glucose	[3] 19 entries				
additional information		additional information				

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Optimizing product formation



How do you optimize product formation (E)?

4. Introduce feedback to detect presence of key intermediates and control activity accordingly

- e.g. if D is essential for product but toxic to cell, want to keep [D] as high as is safe

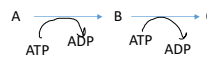
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Can also improve production with fermentation optimization

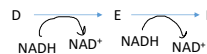
- Cofactors are used by many reactions
 - E.g. ATP, NADH, NADPH, FADH₂, etc

What are the functions of these cofactors?
- Altering production pathways will affect cofactor pools and thus cell health and growth

Ideal situation: Production = Consumption



Energy intensive pathway
- Can increase aeration



Pathway needs reduced co-factors
- Make process anaerobic

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We can also engineer *non-natural* products

Enzymes are *promiscuous*

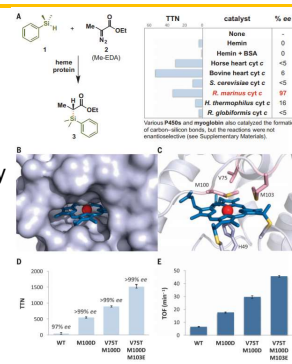
- will catalyze the same chemistry for similar molecules.
 - e.g. Alcohol dehydrogenase (ADH) reducing acetaldehyde to EtOH (C₂) or propanaldehyde to propanol (C₃)
- Given starting substrate and final product, we can identify synthesis pathway (enzymes) required. (*Retrobiosynthesis*)
- Can use databases to identify candidates to screen

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Enzyme promiscuity

- Directed evolution
- Cytochrome P450
 - Natural oxygenase activity
 - Mutated to incorporate Si

First biological C-Si bonds evolved in lab



Kan et al, 2016 Science 354: 1048-1051

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Metabolic engineering summary

Regulation of pathways critical to:

- Maximize product formation
- Minimize impact on host processes (e.g. cofactor balance)
- Prevent build-up of toxic intermediates

Achieved by:

- Push/Pull** - Expression of enzymes at appropriate levels (promoters, RBS, term)
- Feedback** - Programmable control of expression (biological feedback through inducible/repressible promoters or independent control)
- Homologs** - Identifying enzymes with sufficient activity
- Process optimization**

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Metabolic engineering

Engineering host may be required

- To reduce product/intermediate toxicity
- Remove native regulation of enzymes
- Reduce consumption of product

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Next time

- Practical examples: journal club

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