

Introduction and Part I finished

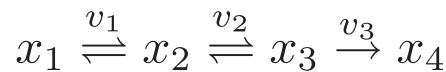
3 weeks down; 7 to go

Where you should be

- Comfortable with
 - P. Chem/Kinetics
 - Biochemistry as defined by the recommended reading materials
- Become familiar with
 - The simulation workflow, the MASSpy, computation
- Getting ready for
 - Visualization or results, pooling of variables, interpretation
 - Dealing with metabolic pathways and enzyme functions
 - Thinking about networks

On to Part II

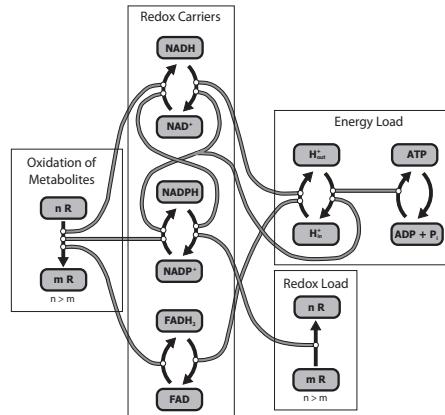
Part I



$$S = \begin{pmatrix} -1 & 0 & 0 \\ 1 & -1 & 0 \\ 0 & 1 & -1 \\ 0 & 0 & 1 \end{pmatrix} \quad \frac{dx}{dt} = Sv(x)$$

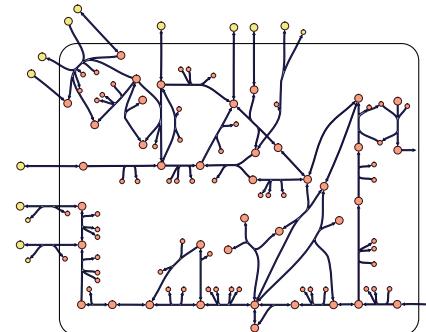
--Basics--

Part II



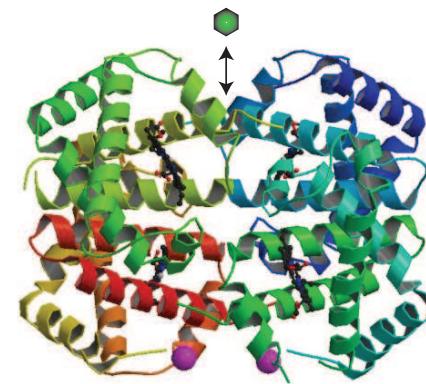
--Biological features--

Part III



--Metabolic networks--

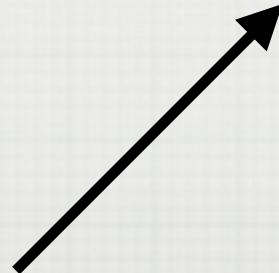
Part IV



--Macromolecular systems--

Parameterization

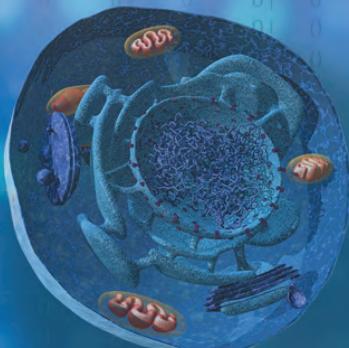
$$\frac{d\mathbf{x}}{dt} = \mathbf{Sv}(\mathbf{x}; \mathbf{k})$$



where do these numbers come from?

Systems Biology

Simulation of Dynamic
Network States



$$\frac{dx}{dt} = S \cdot v(x) \cdot k$$

Bernhard Ø. Palsson

Lecture #7

Estimation and
Orders of
Magnitude

Content

- Orders-of-magnitude
- Estimation
- Observations about Cells
- Examples from Metabolism
 - What are Typical Concentrations?
 - What are Typical Turnover Times?
- Kinetic Parameter Estimation
 - What to look for and where to search?
- Orders-of-magnitude example
- Summary



coffee bean

12 x 8 mm



grain of salt
sesame seed

grain of rice

8 x 2.5 mm

Times regular, 12 point



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Meter (m)	Centimeter (cm)	Millimeter (mm)	Micrometer (μ m) (aka micron)	Nanometer (nm)	Angstrom (\AA)	Picometer (pm)
10^0 m	10^{-2} m	10^{-3} m	10^{-6} m	10^{-9} m	10^{-10} m	10^{-12} m
1 m	0.01 m	0.001 m	0.000001 m	0.000000001 m	0.0000000001 m	0.000000000001 m
	1/100 m	1/1,000 m	1/1,000,000 m	1/1,000,000,000 m	1/10,000,000,000 m	1/1,000,000,000,000 m
	hundredth of a meter	thousandth of a meter	millionth of a meter	billionth of a meter	ten billionth of a meter	trillionth of a meter

<http://learn.genetics.utah.edu/content/begin/cells/scale/>

Estimation:

How many beans are in the jar?





Enrico Fermi (1901 - 1954) was an Italian physicist, particularly remembered for his work on the development of the first nuclear reactor, and for his contributions to the development of quantum theory, nuclear and particle physics, and statistical mechanics.

Famous for quick answers through back-of-the-envelope calculations

Introduction to Fermi problems

The classic Fermi problem is:
"How many piano tuners are there in Chicago?"



Estimation procedure...

- ***Assumptions***

- There are approximately 5,000,000 people living in Chicago.
- On average, there are two people in each household in Chicago.
- Roughly one household in 20 has a piano that is tuned regularly.
- Pianos that are tuned regularly are tuned on average about once per year.
- It takes a piano tuner about 2 hours to tune a piano, including travel time.
- Each piano tuner works 8 hours in a day, 5 days in a week, and 50 weeks in a year.
- ***From these assumptions we can compute that the number of piano tunings in a single year in Chicago is***

$$\frac{5 * 10^6 \text{ people}}{\left(\frac{2 \text{ people}}{1 \text{ household}}\right)} * \frac{1 \text{ piano}}{20 \text{ households}} * \frac{1 \text{ piano tuning}}{1 \text{ piano} * 1 \text{ year}} = \boxed{\frac{125000 \text{ piano tunings}}{1 \text{ year}}}$$

$$\frac{50 \text{ weeks}}{1 \text{ year}} * \frac{5 \text{ days}}{1 \text{ week}} * \frac{8 \text{ hrs}}{1 \text{ day}} * \frac{1 \text{ piano tuning}}{2 \text{ hrs} * 1 \text{ piano tuner}} = \boxed{\frac{1000 \text{ piano tunings}}{1 \text{ year} * 1 \text{ piano tuner}}}$$

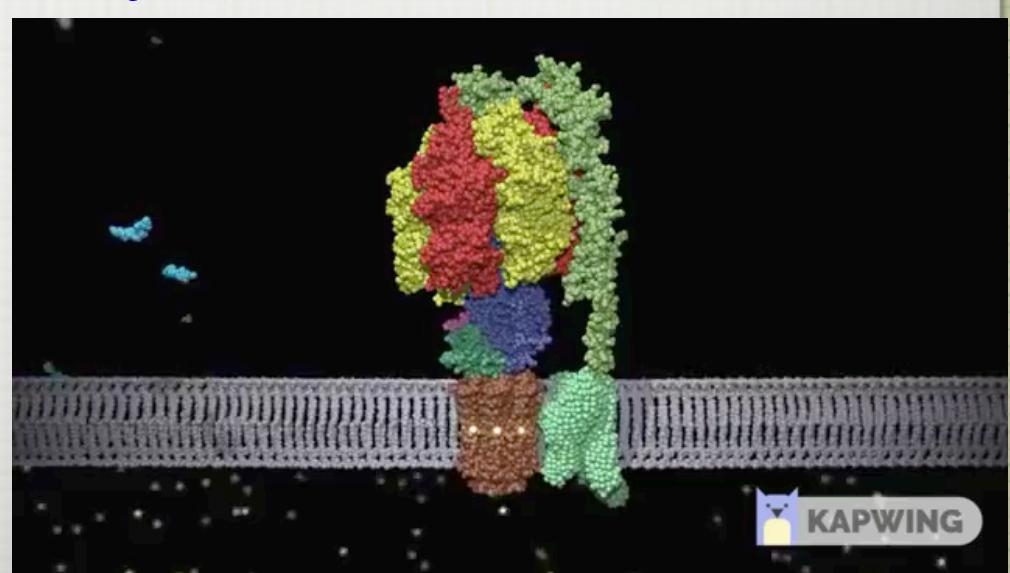
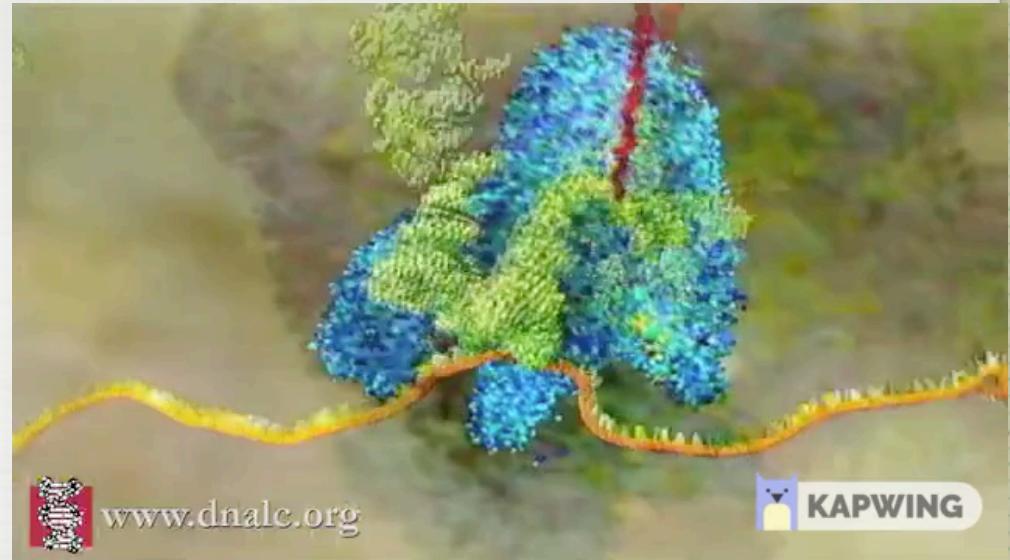
$$\frac{125000 \text{ piano tunings}}{1 \text{ year}} * \frac{1 \text{ year} * 1 \text{ piano tuner}}{1000 \text{ piano tunings}} = \boxed{125 \text{ piano tuners in Chicago}}$$

Real significance ...

- Possible to estimate key biological quantities on the basis of a few foundational facts and simple ideas from physics and chemistry.
- Numbers collected by the scientific community that initially appear unrelated are brought together as a tool of inference to shed light on biological mechanisms.

Biological examples

- How many proteins can be produced from a single mRNA in *E. coli*?
- How many ATP synthase complexes are required for optimal growth on glucose in *E. coli*?



Proteins/mRNA

- **Assumptions**

- The average length of mRNA is 1100 nt.
- A ribosome can bind to mRNA every 50 nt due to structural considerations.
- The rate of translation for one ribosome is 16 amino acids per second.
- The average length of a protein is 367 amino acids.
- The average half-life of mRNA is 6 minutes (360 seconds).

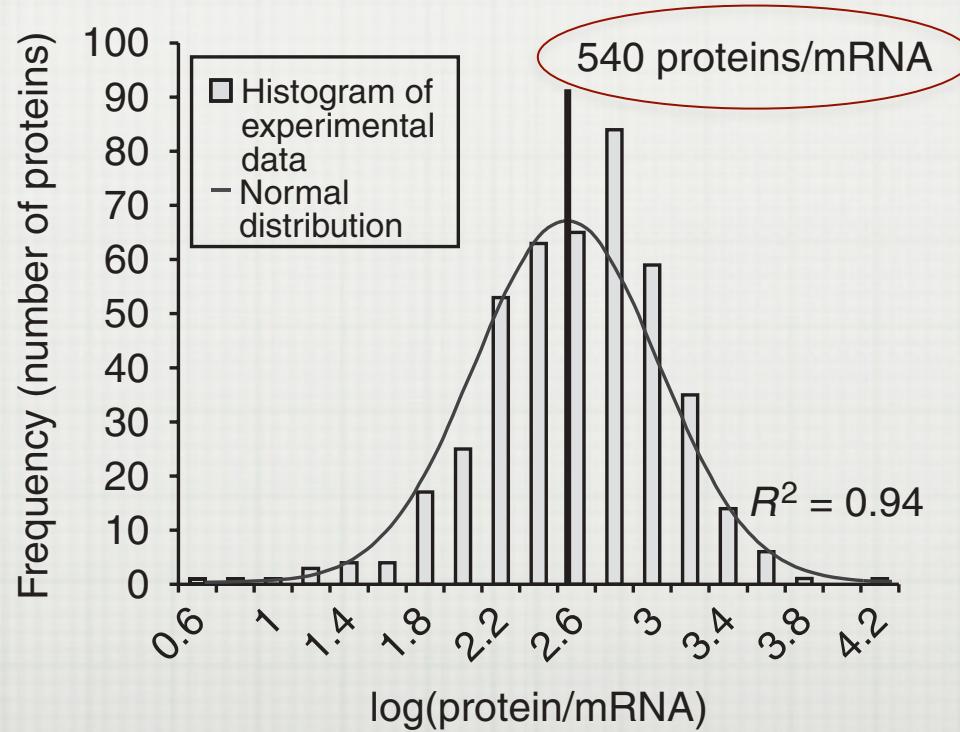
- **From these assumptions we can compute that the number of proteins translated over the lifetime of an mRNA transcript is**

$$\frac{1100 \text{ nt}}{1 \text{ mRNA transcript}} * \frac{1 \text{ ribosome}}{50 \text{ nt}} = \boxed{\frac{22 \text{ ribosomes}}{1 \text{ mRNA transcript}}} \text{ (Maximum ribosome loading)}$$

$$\frac{16 \text{ AA}}{1 \text{ sec} * 1 \text{ ribosome}} * \frac{22 \text{ ribosomes}}{1 \text{ mRNA transcript}} = \boxed{\frac{352 \text{ AA}}{1 \text{ sec}}} \text{ (Translation rate w/ maximum ribosome load)}$$

$$\frac{1 \text{ protein}}{367 \text{ AA}} * \frac{352 \text{ AA}}{1 \text{ sec}} \approx \frac{1 \text{ protein}}{1 \text{ sec}} \quad \rightarrow \quad \frac{1 \text{ protein}}{1 \text{ sec}} * \frac{(360 \text{ sec}/\ln 2)}{1 \text{ mRNA}} \approx \boxed{\frac{519 \text{ proteins}}{1 \text{ mRNA}}}$$

Let's see how we did...



Lu et al., *Nat Biotech* 2007

Another example: ATP synthase

- **Assumptions**

- Max velocity of ATP synthase: 230 revolutions/sec (828000 rev/ hr) [PMID 15668386]
- Three ATP produced each revolution of ATP synthase.
- For aerobic growth at an uptake rate of 10 mmol glucose gDW⁻¹ hr⁻¹, growth computations show that required flux through ATP synthase is 52 mmol gDW⁻¹ hr⁻¹
- The optimal growth rate is 0.7367 doublings/hr (approx. 1 doubling/hour).
- There is approximately 200,000 inner membrane proteins
- Each ATP synthase complex has 22 proteins.

- **From these assumptions, we can compute how many inner membrane proteins are**

$$\frac{0.052 \text{ mol}}{1 \text{ gDW hr}} * \frac{2.8 * 10^{-13} \text{ gDW}}{1 \text{ cell}} * \frac{6.022 * 10^{23} \text{ ATP}}{1 \text{ mol}} = \frac{8,768,032,000 \text{ ATP}}{1 \text{ hr}} \quad \begin{pmatrix} \text{ATP required per hour} \\ \text{for optimal growth} \end{pmatrix}$$

$$\frac{230 \text{ revs}}{1 \text{ sec}} * \frac{3600 \text{ secs}}{1 \text{ hr}} * \frac{3 \text{ ATP}}{1 \text{ rev}} \approx \frac{2.5 * 10^6 \text{ ATP}}{1 \text{ hr} * 1 \text{ ATP synthase}} \quad \begin{pmatrix} \text{ATP produced per hour} \\ \text{by 1 ATP synthase} \end{pmatrix}$$

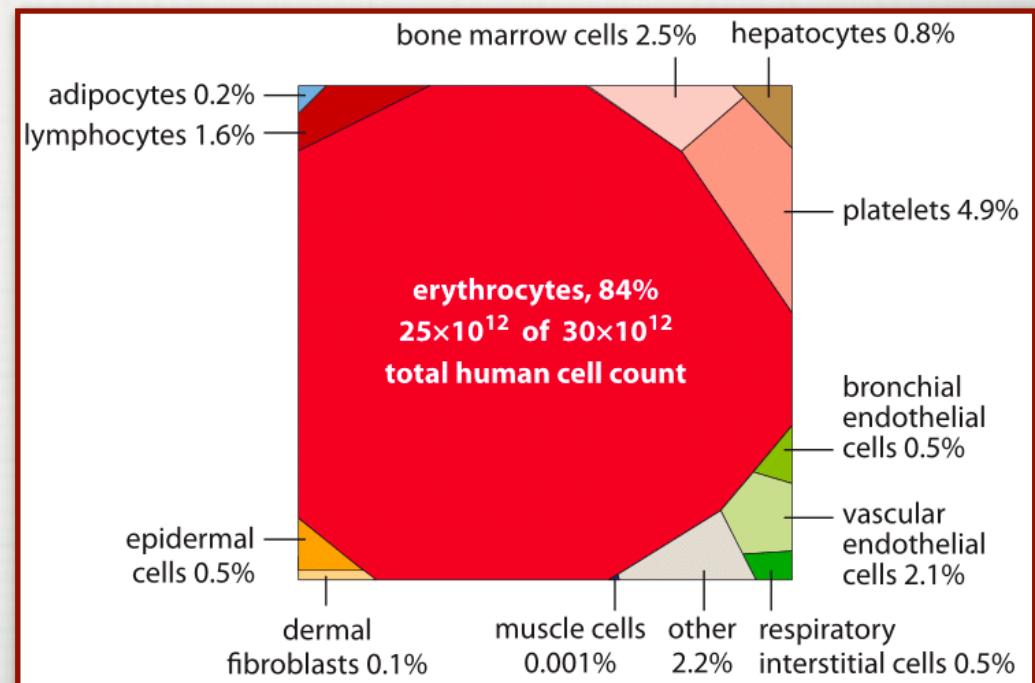
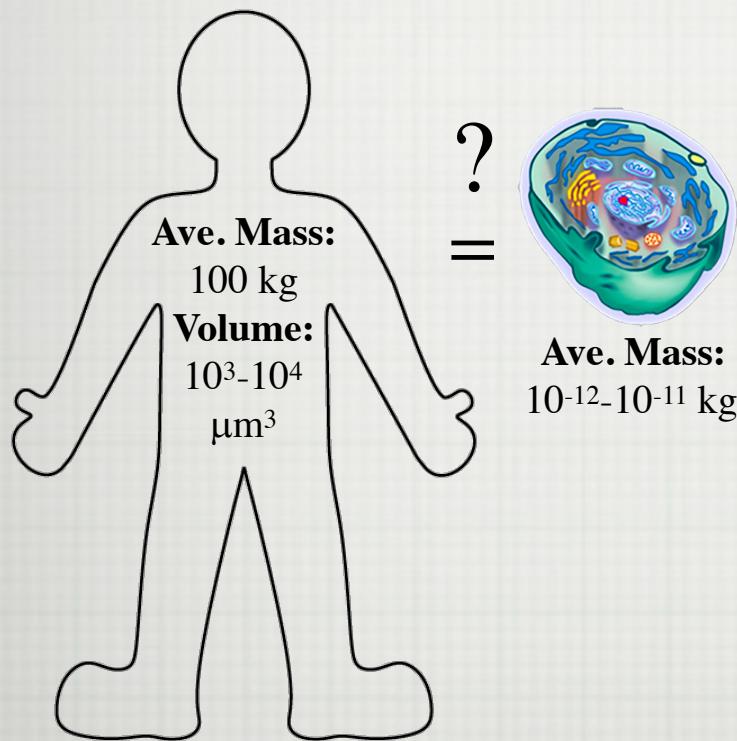
$$\frac{8,768,032,000 \text{ ATP}}{1 \text{ hr}} * \frac{1 \text{ hr} * 1 \text{ ATP synthase}}{2.5 * 10^6 \text{ ATP}} \approx 3510 \text{ ATP synthase complexes} \quad (\text{working at } V_{max})$$

$$\frac{3510 \text{ ATP synthase} * 22 \text{ proteins/1 ATP synthase}}{200,000 \text{ total proteins}} \approx 0.4$$

ATP synthase accounts for approximately 40% of inner membrane proteins

Cells in a Human

How many cells are there in the human body?



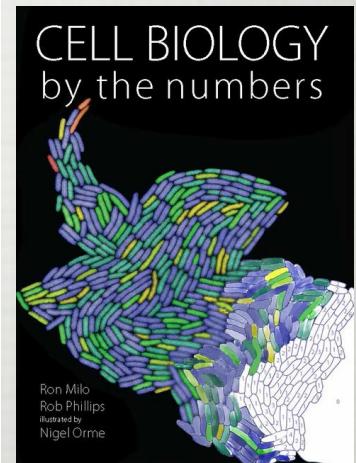
$$N \approx \frac{100 \text{ kg human}}{10^{-12} - 10^{-11} \text{ kg per cell}} = 10^{13} - 10^{14} \text{ cells in the human body}$$

Resource: BioNumbers database

Property	Organism	Value	Units	ID	Details
Average duration of a single eye blink	Human Homo sapiens	0.1-0.4	sec	100706	http://faculty.washington.edu/mminoli/BioNumbers/100706.html
Size of glucose molecule (open chain form)	Generic	1.5	nm	106979	Daniel Minoli, Nanotechnology...
One unit of OD600 corresponds to a cell wet weight and a cell dry weight of	Bacteria Escherichia coli	Cell wet weight 1.7g/L: cell dry weight 0.39g/L		109836	Glazyrina J, Materne EM...
Identification of total proteins in <i>A. baumannii</i> DU202 cultured in LB medium supplemented with imipenem	Bacteria Acinetobacter baumannii	Table - link	N/A	111523	Yun SH et al., Quantitative...
Estimations of periplasmic volume from distributions of different solutes in suspensions of <i>Salmonella typhimurium</i> LT2	Bacteria <i>Salmonella typhimurium</i>	Table - link	N/A	108473	Stock JB, Rauch B, Roseman...
Number of hairs on human head	Human Homo sapiens	90,000-150,000	unitless	101509	Amazing numbers in biology...
Number of cells in colony	Bacteria Escherichia coli	3.3e+9 (Table - link)	Cells/culture	104458	Mashimo K, Nagata Y, Kawata M...
Diameter of neutrophil	Human Homo sapiens	8.4 (7.3-9.7 Table - link)	µm	100510	Downey GP, Doherty DE...
Density and mass of each organ/tissue	Human Homo sapiens	Table - link	N/A	110245	International commission...
Diameter of HEK-293 cell	Human Homo sapiens	13	µm	108893	http://www.invitrogen.com

+3,000 monthly visitors

+50 countries



Source: <http://bionumbers.hms.harvard.edu/>

<http://book.bionumbers.org/>

BioNumbers is coordinated and developed by Ron Milo at the Weizmann Institute in Israel.

Key Concepts

- Characteristic order of magnitude for key quantities that characterize cellular functions can be estimated
- Data on cell size, mass, composition, metabolic complexity, and genetic makeup are available
- Numerous databases now available on the web
- Useful estimates of fluxes, concentrations, kinetics, and power densities in the intracellular environment can be made based on this data

Key Numbers for Cell Biologists

Orders of Magnitude

SOME OVERALL OBSERVATIONS ABOUT CELLS

Cell size

1. Bacteria (*E. coli*): $\approx 0.7\text{-}1.4 \mu\text{m}$ diameter, $\approx 2\text{-}4 \mu\text{m}$ length, $\approx 0.5\text{-}5 \mu\text{m}^3$ in volume; $10^8\text{-}10^9 \text{ cell/ml}$ for culture with $\text{OD}_{600}\approx 1$
2. Yeast (*S. cerevisiae*): $\approx 3\text{-}6 \mu\text{m}$ diameter, $\approx 20\text{-}160 \mu\text{m}^3$ in volume
3. Mammalian cell volume: $100\text{-}10000 \mu\text{m}^3$; Hela: $500\text{-}5000 \mu\text{m}^3$ (adherent on slide $\approx 15\text{-}30 \mu\text{m}$ diameter)

Length Scales Inside Cells

4. Nucleus volume $\approx 10\%$ of cell volume
5. Cell membrane thickness $\approx 4\text{-}10 \text{ nm}$
6. "Average" protein diameter $\approx 3\text{-}6 \text{ nm}$
7. Base pair: 2 nm (D) $\times 0.34 \text{ nm}$ (H)
8. Water molecule diameter $\approx 0.3 \text{ nm}$

Division, Replication, Transcription, Translation & Degradation Rates

- at 37°C with a temperature dependence Q10 of $\approx 2\text{-}3$
9. Cell cycle time (exponential growth in rich media): *E. coli* $\approx 20\text{-}40 \text{ min}$; yeast $70\text{-}140 \text{ min}$; human cell line (Hela): $15\text{-}30 \text{ hours}$
 10. Rate of replication by DNA polymerase *E. coli* $\approx 200\text{-}1000 \text{ bases/s}$; human $\approx 40 \text{ bases/s}$. Transcription by RNA polymerase $10\text{-}100 \text{ bases/s}$
 11. Translation rate by ribosome $10\text{-}20 \text{ aa/s}$
 12. Degradation rates (proliferating cells): mRNA half life $<$ cell cycle time; protein half life \approx cell cycle time

Concentration

13. Concentration of 1 nM in: *E. coli* $\approx 1 \text{ molecule/cell}$; Hela $\approx 1,000 \text{ molecules/cell}$
14. Characteristic concentration for a signaling protein $\approx 10 \text{ nM}\text{-}1 \mu\text{M}$
15. Water content: $\approx 70\%$ by mass; General elemental composition (dry weight) of *E. coli*: $\approx \text{C}_5\text{H}_{10}\text{O}_5\text{N}_1$; Yeast $\approx \text{C}_6\text{H}_{10}\text{O}_3\text{N}_1$
16. Composition of *E. coli* (dry weight): $\approx 55\%$ protein, 20% RNA, 10% lipids, 15% others
17. Protein conc. $\approx 100 \text{ mg/ml}=3 \text{ mM}$. $10^6\text{-}10^7$ per *E. coli* (depending on growth rate); Total metabolites (MW<1kD) $\approx 300\text{mM}$

Energetics

18. Membrane potential $\approx 70\text{-}200 \text{ mV} \rightarrow 2\text{-}6 k_B T$ per electron ($k_B T$ =thermal energy)
19. Free energy (ΔG) of ATP hydrolysis under physiological conditions $\approx 40\text{-}60 \text{ kJ/mole} \rightarrow \approx 20 k_B T/\text{molecule ATP}$; ATP molecules required to make an *E. coli* cell $\approx 10\text{-}50 \times 10^9$
20. ΔG^0 resulting in order of magnitude ratio between products and reactants concentrations: $\approx 6 \text{ kJ/mol} \approx 60 \text{ meV} \approx 2 k_B T$

Useful biological numbers extracted from the literature. Numbers and ranges should only serve as "rule of thumb" values. References are in the online annotated version at the BioNumbers website. Consult website and original references to learn about the details of the system under study including growth conditions, method of measurement, etc.

Diffusion and Catalysis Rate

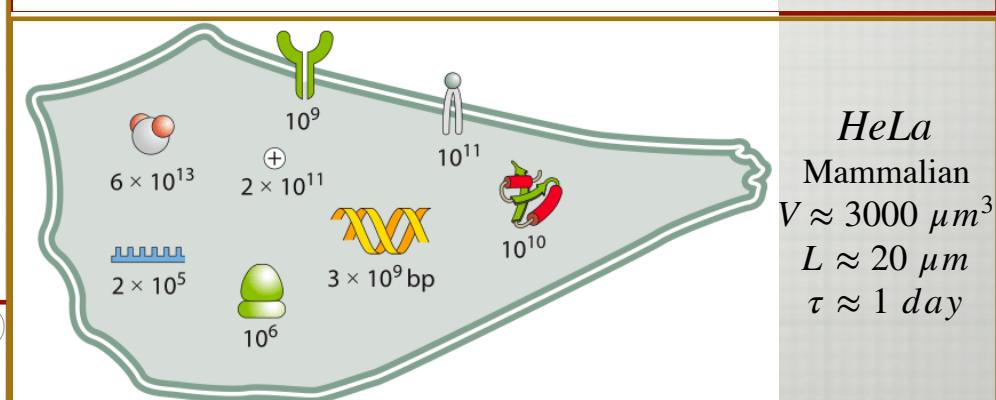
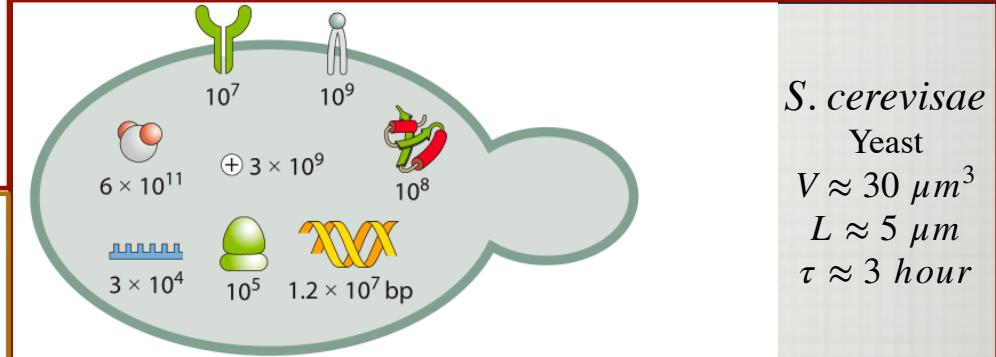
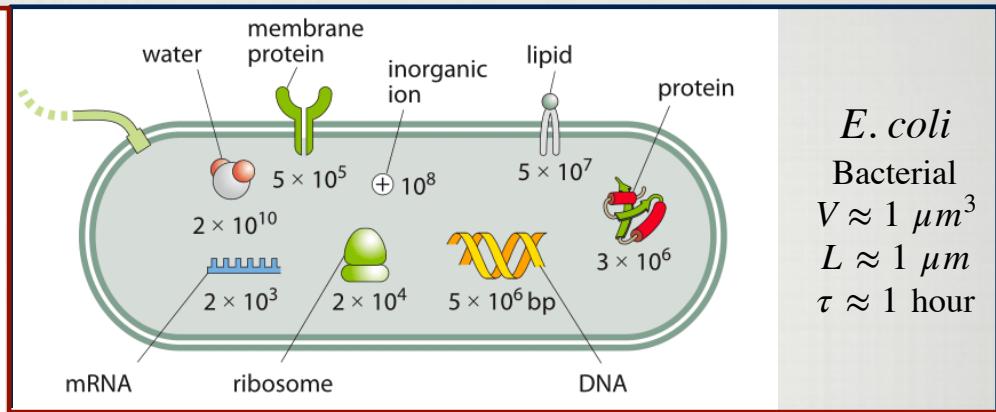
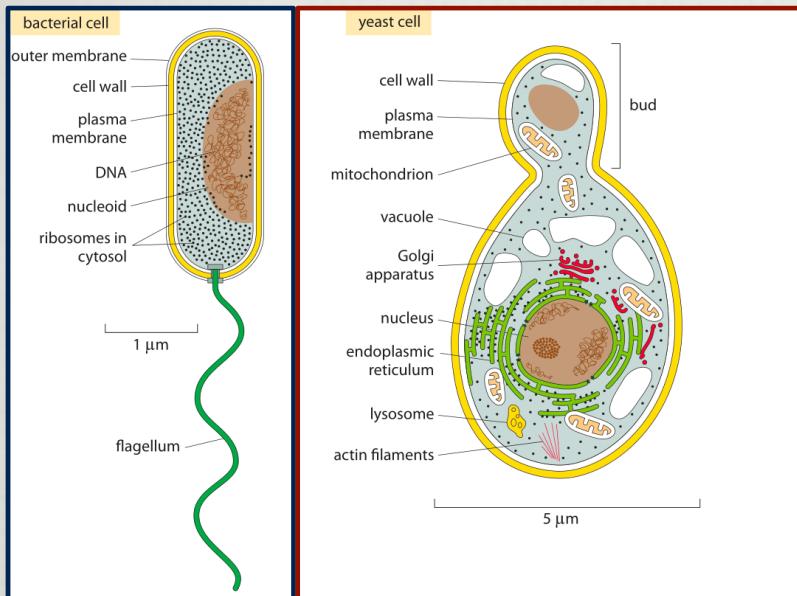
21. Diffusion coefficient for an "average" protein: in cytoplasm $D=5\text{-}15 \mu\text{m}^2/\text{s} \rightarrow 10 \text{ millisec}$ to traverse an *E. coli* $\rightarrow \approx 10 \text{ s}$ to traverse a mammalian (Hela) cell; small metabolite in water $D=500 \mu\text{m}^2/\text{s}$
22. Diffusion limited on-rate for characteristic protein $\approx 10^8\text{-}10^9 \text{ s}^{-1}\text{M}^{-1}$ \rightarrow for a protein substrate of concentration $\approx 1 \mu\text{M}$ the diffusion limited on-rate is $\approx 100\text{-}1000 \text{ s}^{-1}$ thus limiting the catalytic rate k_{cat}

Genome sizes & Error Rates

23. Genome size: *E. coli* $\approx 5 \text{ Mbp}$; *S. cerevisiae* (yeast) $\approx 12 \text{ Mbp}$; *C. elegans* (nematode) $\approx 100 \text{ Mbp}$; *D. melanogaster* (fruit fly) $\approx 120 \text{ Mbp}$; *A. thaliana* (arabidopsis) $\approx 120 \text{ Mbp}$; *M. musculus* (mouse) $\approx 2.5 \text{ Gbp}$; *H. sapiens* (human) $\approx 2.9 \text{ Gbp}$; *T. aestivum* (wheat) $\approx 16 \text{ Gbp}$
24. Number of protein-coding genes: *E. coli* $\approx 4,000$; *S. cerevisiae* $\approx 6,000$; *C. elegans*, *A. thaliana*, *M. musculus*, *H. sapiens* $\approx 20,000$
25. Mutation rate in DNA replication $\approx 10^{-8}\text{-}10^{-10} \text{ per bp}$
26. Misincorporation rate: transcription $\approx 10^{-4}$ per nucleotide; translation $\approx 10^{-3}\text{-}10^{-4}$ per amino-acid

Click on a number to see full description and reference www.BioNumbers.org

Cell Sizes and Interiors

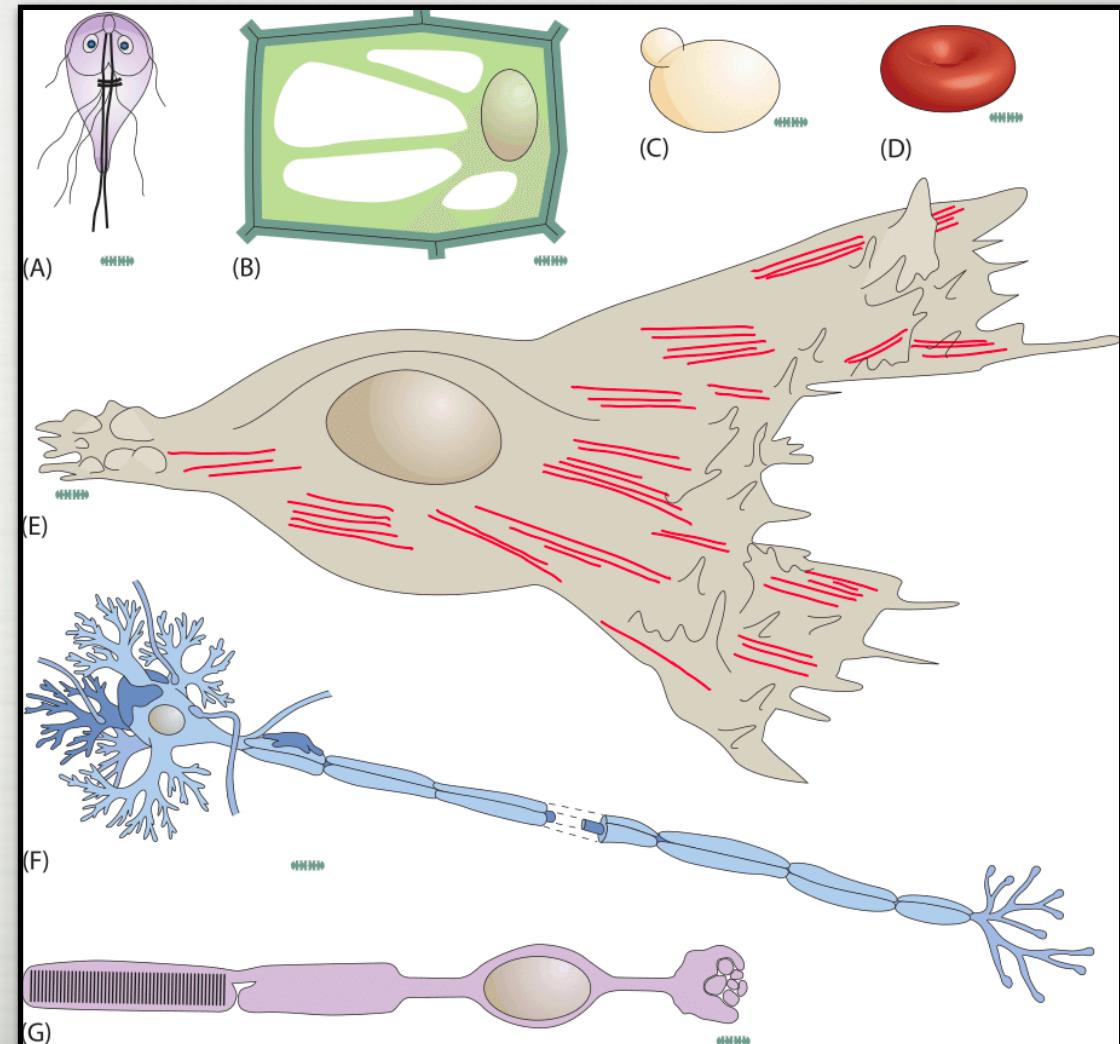


Range of Cell Sizes and Shapes

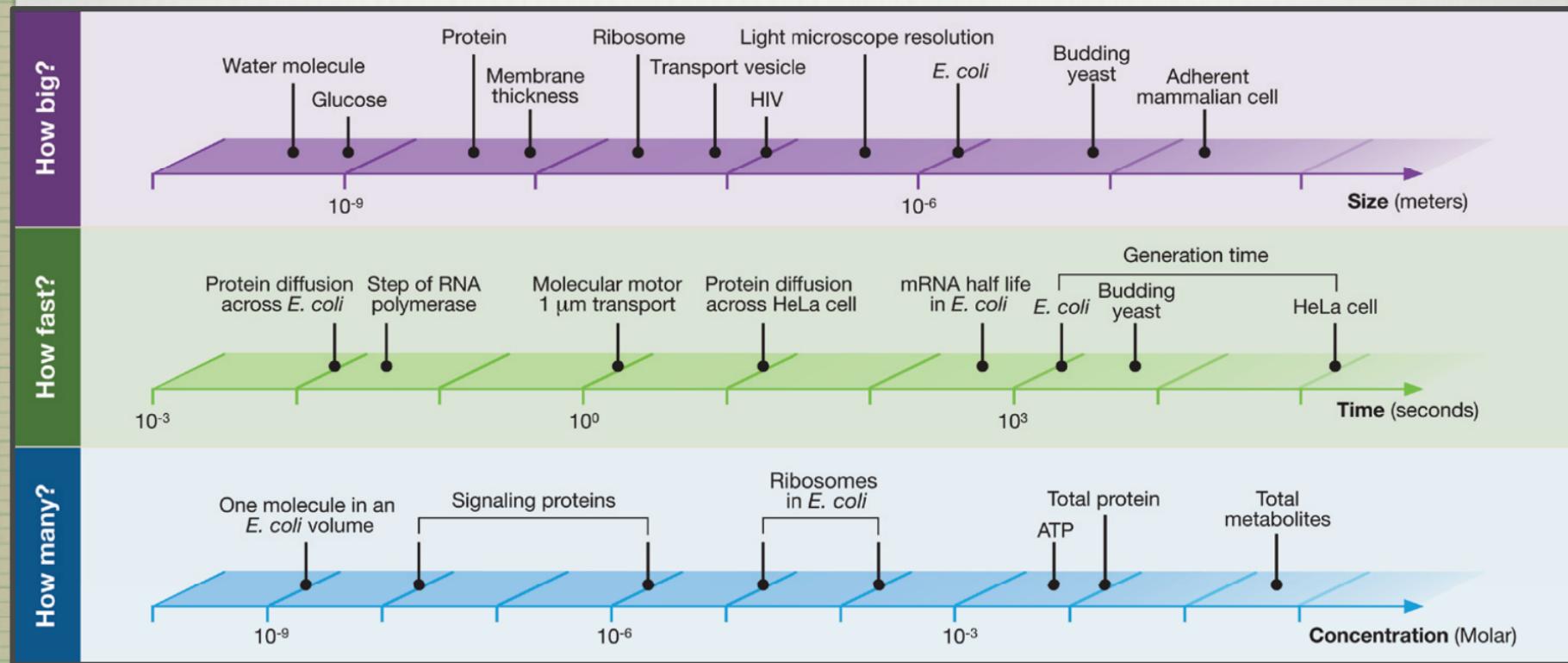
- Relative to *E. Coli*

1 micron

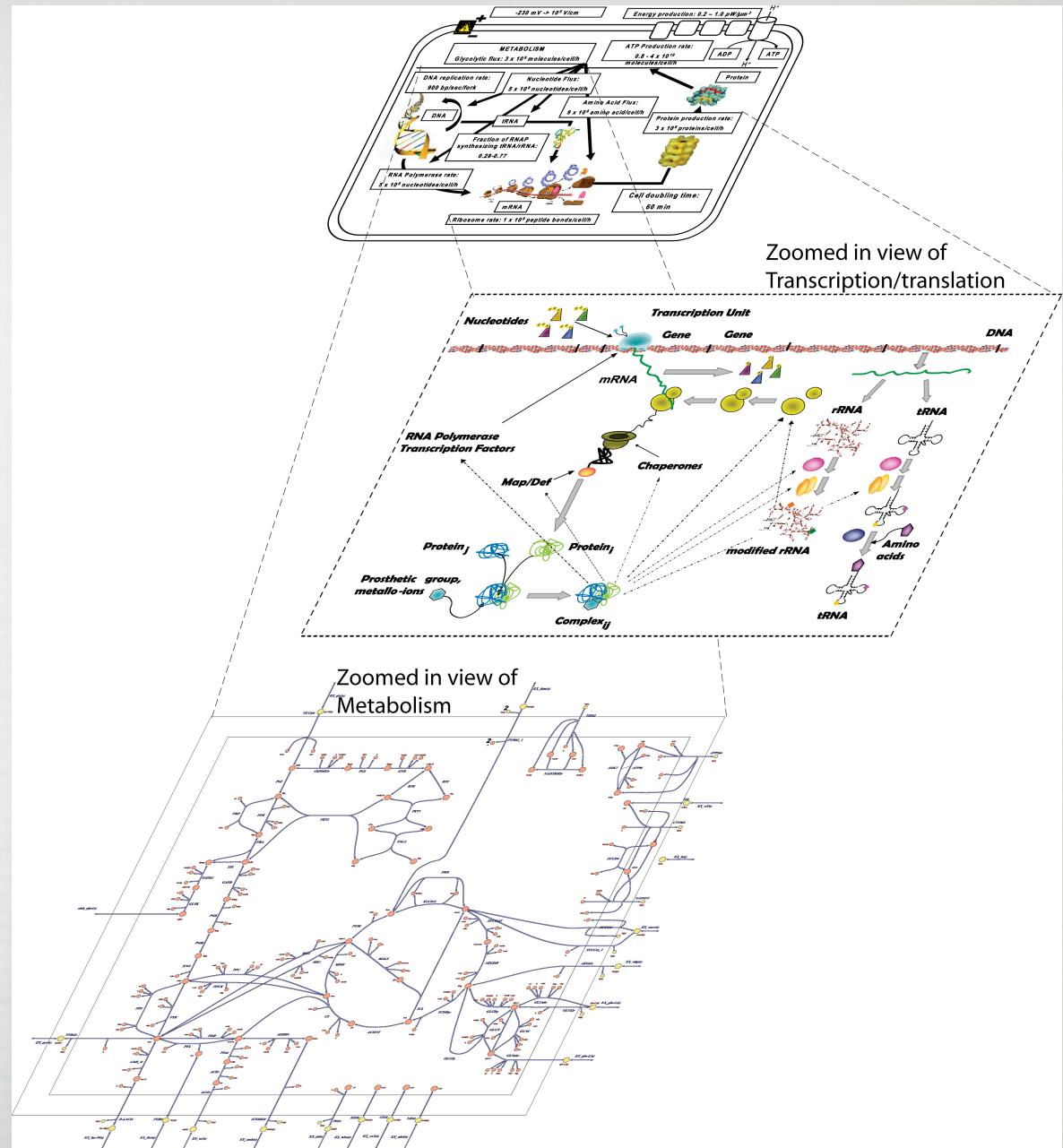
- (A) Giardia lamblia
- (B) Plant cell
- (C) Budding yeast cell
- (D) Red blood cell
- (E) Fibroblast cell
- (F) Eukaryotic nerve cell
- (G) Rod cell from retina



The Multi-Scale Nature of Biology

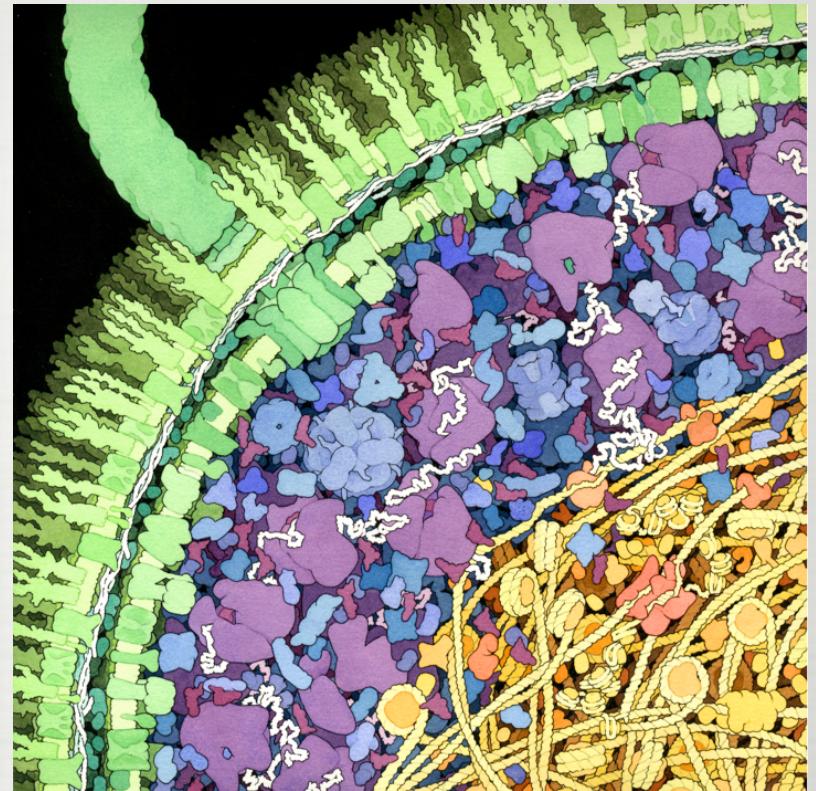


Multi-scale relationships (many processes to consider): metabolism, transcription, translation, phenotypes



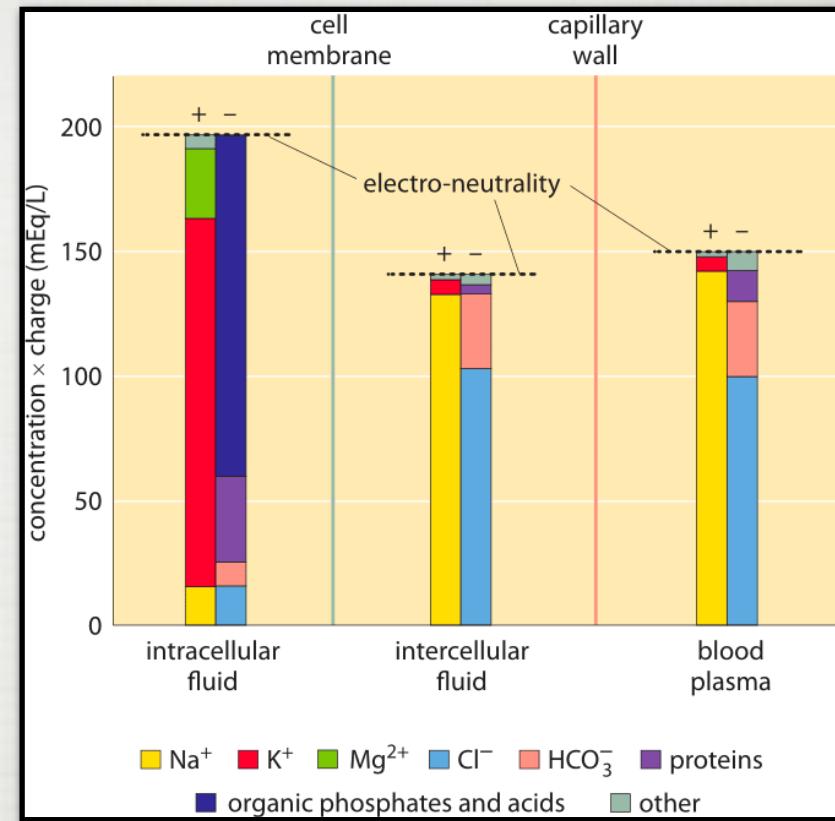
Small molecule scale

METABOLISM



Concentrations of Ions

- Census of ionic charges in a mammalian tissue cell
 - Law of electro-neutrality
 - Osmotic balance between compartments
- Intracellular Fluid
 - Many metabolites & **proteins** are negatively charged
 - **K⁺, Mg²⁺, Cl⁻** are most abundant ions
- Concentrations can change by more than an order of magnitude depending on...
 - Cell type
 - Physiological conditions
 - Environmental conditions



ion conc. (mM)	sea water	<i>E. coli</i>	<i>S. cerevisiae</i>	mammalian cell (heart or RBC)	blood plasma
K ⁺	≈10	30-300	300	100	4
Na ⁺	≈500	10	30	10	100-200
Mg ²⁺	≈50	30-100 (bound); 0.01-1 (free)	50	10 (bound) 0.5 (free)	1
Ca ²⁺	≈10	3 (bound); 100 nM (free)	2 (bound)	10-100 nM (free)	2
Cl ⁻	≈500	10-200 media dependent		5-100	100

Typical Metabolite Concentration

1. The number of different metabolites present in *E. coli* is on the order of 1000.
2. An average metabolite has a median molecular weight of about 312 gram/mol.
3. We estimate the typical metabolite concentration:

$$C_m \approx \frac{1 \text{ gm/ cm}^3 \cdot 0.01}{1000 \cdot 312 \text{ gm/ mole}} \approx 32 \mu\text{M}$$

and:

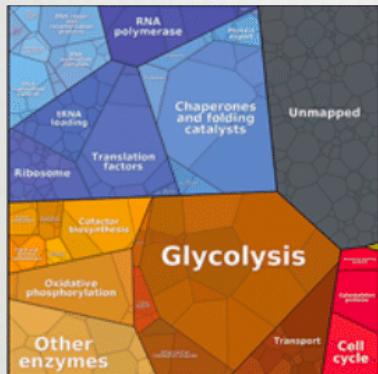
$$\begin{aligned} 1 \mu\text{M} &= \frac{10^{-6} \text{ moles}}{(10 \text{ cm})^3} \cdot \frac{\text{cm}^3}{(10^4 \mu\text{m})^3} \cdot \frac{6 \cdot 10^{23} \text{ molecules}}{\text{mole}} \\ &= 600 \text{ molecules/ } \mu\text{m}^3 \end{aligned}$$

A typical metabolite concentration translates
into about:

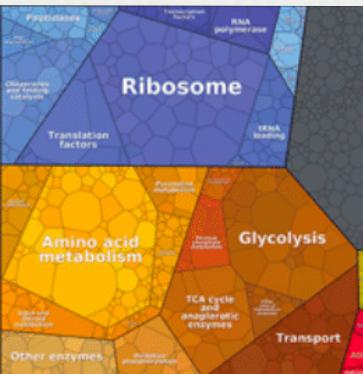
19,000 molecules per cubic micron!

Proteomaps

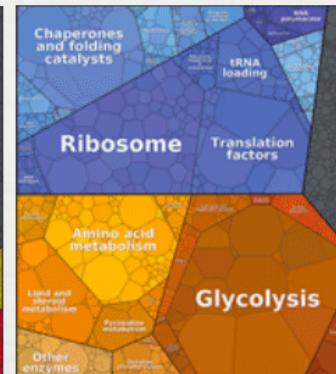
M. pneumoniae



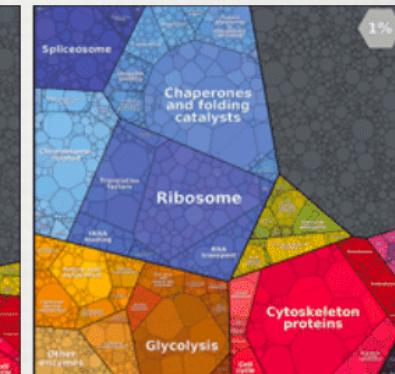
E. coli



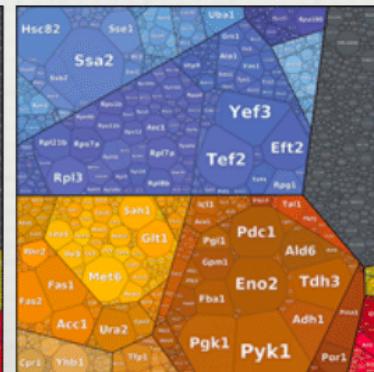
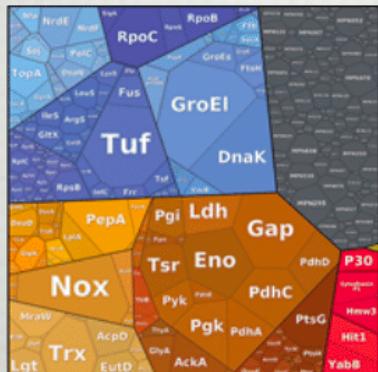
S. cerevisiae



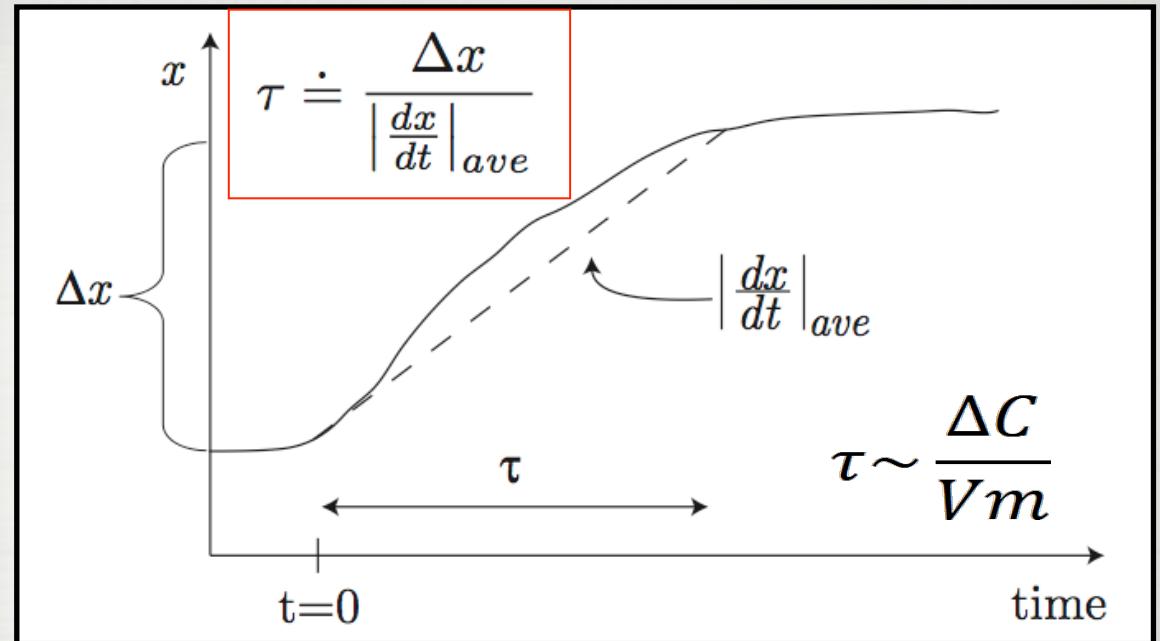
H. Sapiens (HeLa)



By functional category



By protein name

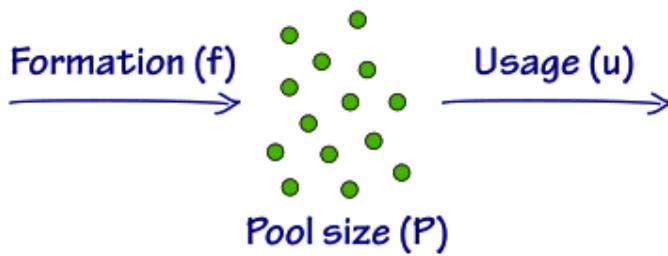


Timescales

TURNOVER TIMES

Defining the Turnover Time

defining the turnover time



metabolite	turnover time (s)		
	<i>Arabidopsis</i>	<i>S. cerevisiae</i>	<i>E. coli</i>
NADP	0.01	-	-
ADP	0.07	0.3	0.8
Calvin-Benson cycle intermediates (R5P, S6P, X5P, Ru5P, SBP, RuBP)	0.1-1	-	-
DHAP/G3P	0.2	-	-
ATP	0.3	1.4	2
3PGA	0.7	7	3
inorganic C	0.8	-	-
FBP	0.8	7	1.2
pyruvate	-	1.7	1.5
F6P	3	7	1.2
AMP	-	3	9
UDPG	40	-	-
G6P	40	17	4
glycerol-3-phosphate	-	60	13
TCA cycle (Suc, Fum, Mal)	-	4-30	0.7-9

Turnover Times of Glucose in *E. coli*

Estimating a glycolytic flux

1. The total stoichiometric amount of glucose that is needed to generate one *E. coli* cell is about 3 billion molecules per cell.
2. Doubling time for *E. coli* is 60 min.
3. Volume of the *E. coli* cell is $1-2\mu\text{m}^3$

$$\begin{aligned}\text{Glycolytic flux} &= \frac{3.0 \times 10^9 \text{ molecules}}{\text{cell} \times 60 \text{ min}} \\ &= \frac{3.0 \times 10^9 \text{ molecules}}{\text{cell} \times \text{hr}} \\ &= 4.2 \times 10^5 - 8.4 \times 10^5 \frac{\text{molecules}}{\mu\text{m}^3 \times \text{sec}}\end{aligned}$$

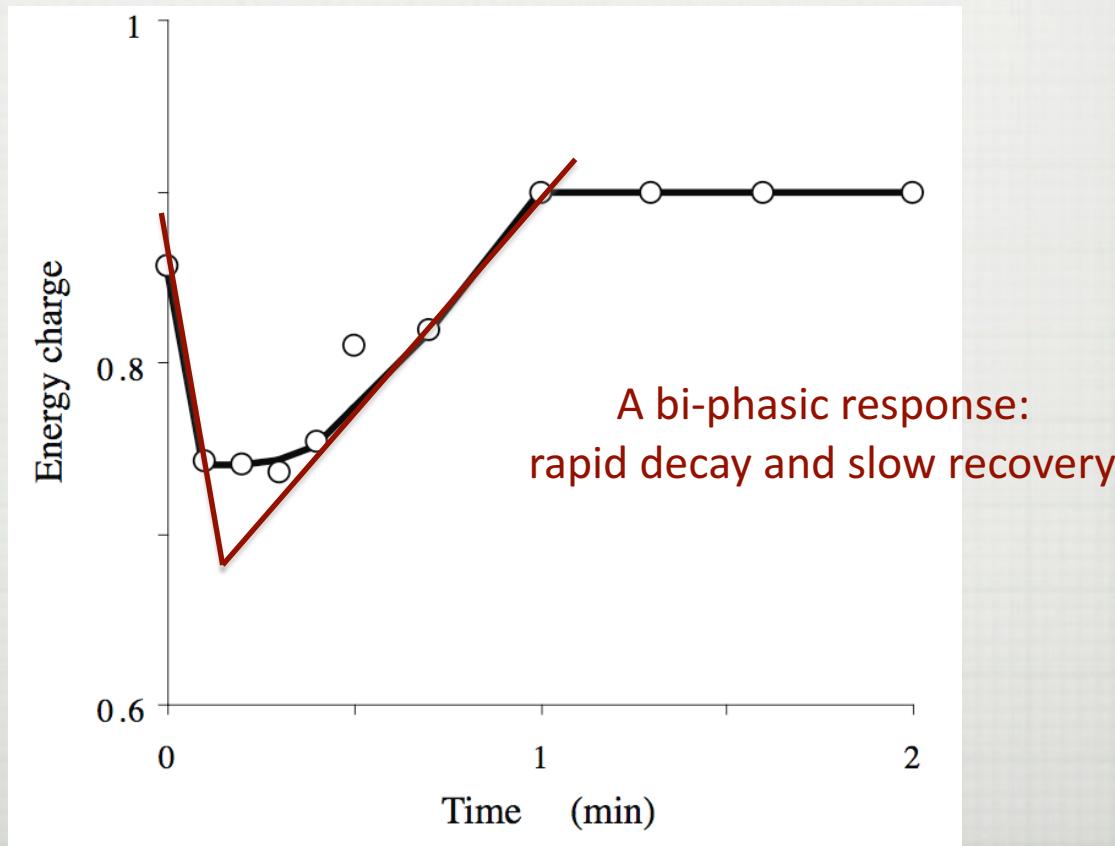
Glucose turnover in rapidly growing *E. coli*:

- Extracellular Glucose concentration: $1-5 \text{ mM}$ ($6-30 \times 10^5 \text{ molecules/cell}$)
- Turnover time is on the order of sec

$$\tau_{glu} = \frac{6 - 30 \times 10^5 \text{ molecules/cell}}{4.2 - 8.4 \times 10^5 \frac{\text{molecules}}{\mu\text{m}^3}/\text{cell}} = 1.5 - 4 \text{ sec}$$

The Measured Time Response of the Energy Charge

$$\frac{(2\text{ATP}+\text{ADP})}{2(\text{ATP}+\text{ADP}+\text{AMP})}$$



Turnover times in RBC glycolysis

$$\tau_R = \frac{C_m}{1.25mM/hr}$$

Compound	Concentration (mM)	Turnover time (sec)
G6P	0.038 ± 0.012	109
F6P	0.016 ± 0.03	46
FDP	0.0076 ± 0.004	21
DHAP	0.140 ± 0.08	7 (min)
GA3P	0.0067 ± 0.001	19
1,3DPG	0.0004	1.1
3PG	0.045	2 (min)
2PG	0.014 ± 0.005	40
PEP	0.017 ± 0.002	50
PYR	0.077 ± 0.05	4 (min)
LAC	1.10 ± 0.50	52 (min)

Fast
and slow:
Distributed
time
constants

NUMERICAL VALUES OF KINETIC CONSTANTS

Sources of Information

- BRENDA: Enzyme Database
- KEGG: Kyoto Encyclopedia of Genes and Genomes
- BioCyc: Pathway/Genome Databases (PGDBs)
- EXPASY: Bioinformatics resource portal
- UniProtKB: Universal protein knowledge
- KDBI: Kinetic Data of Bio-molecular Interactions
- NCBI
 - Entrez Gene, PubMed, PubChem, etc.

What to look for?

- Association rate ($k^+ / k_{\text{on}} / k_{\text{forward}}$)
- Dissociation rate ($k^- / k_{\text{off}} / k_{\text{reverse}}$)
- Catalytic (k_{cat})
- Equilibrium/Dissociation (K_{eq} / K_D)
- Inhibition (K_I)
- Michaelis Menten (K_M)
- Inhibitor Conc. at 50% response (IC50)
- Maximum reaction velocity (V_{max})
- Experimental Conditions
 - pH value/Temperature conditions (Standard?)



BRENDA



The Comprehensive Enzyme Information System

Text-based queries

- Full-text Search
- Advanced Search
- Enzyme & Disease



Structure-based queries

- Ligand Structure Search
- Metabolic Pathways
- Enzyme Structures



Explorer

- Enzyme Classification
- TaxTree
- Protein folding: CATH / SCOPe
- Ontologies



Visualization

- Word Maps
- Genomes
- Functional Parameter Statistics
- Metabolic Pathways



Prediction

- Membrane Helices
- Localization Prediction
- EnzymeDetector



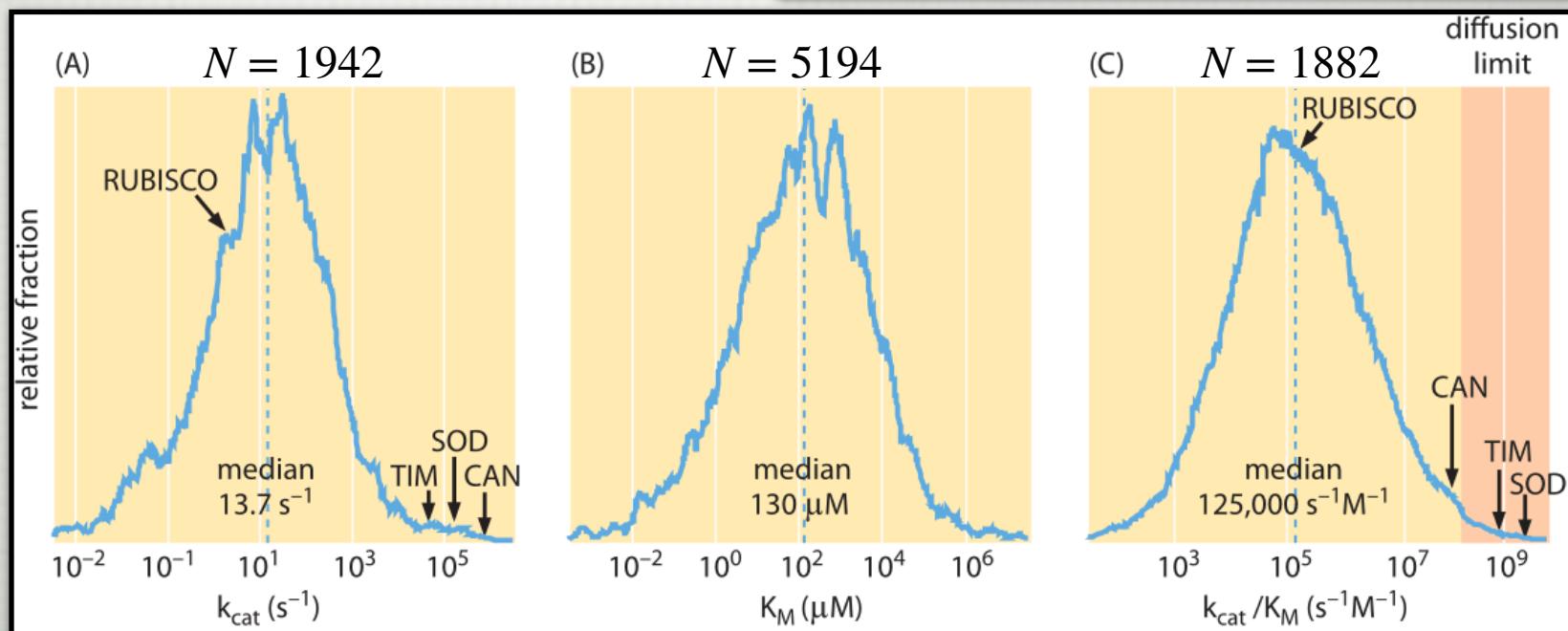
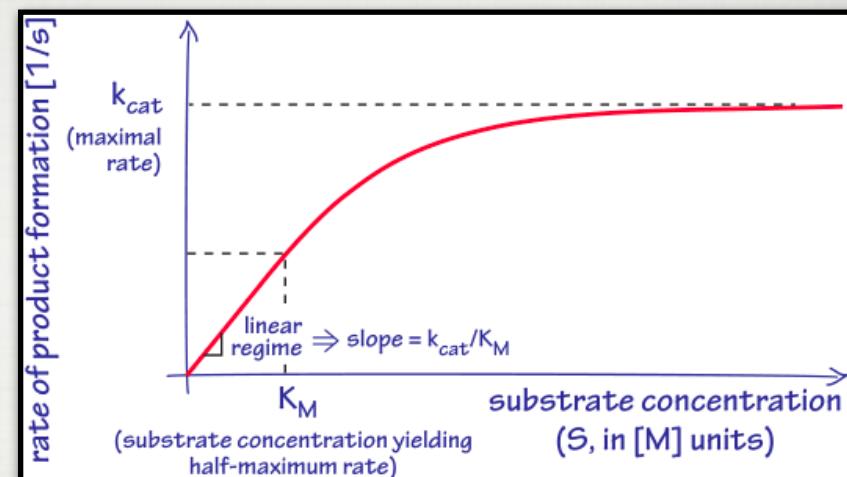
Supporting

- BRENDA
- Tissue Ontology
- Biochemical Reactions



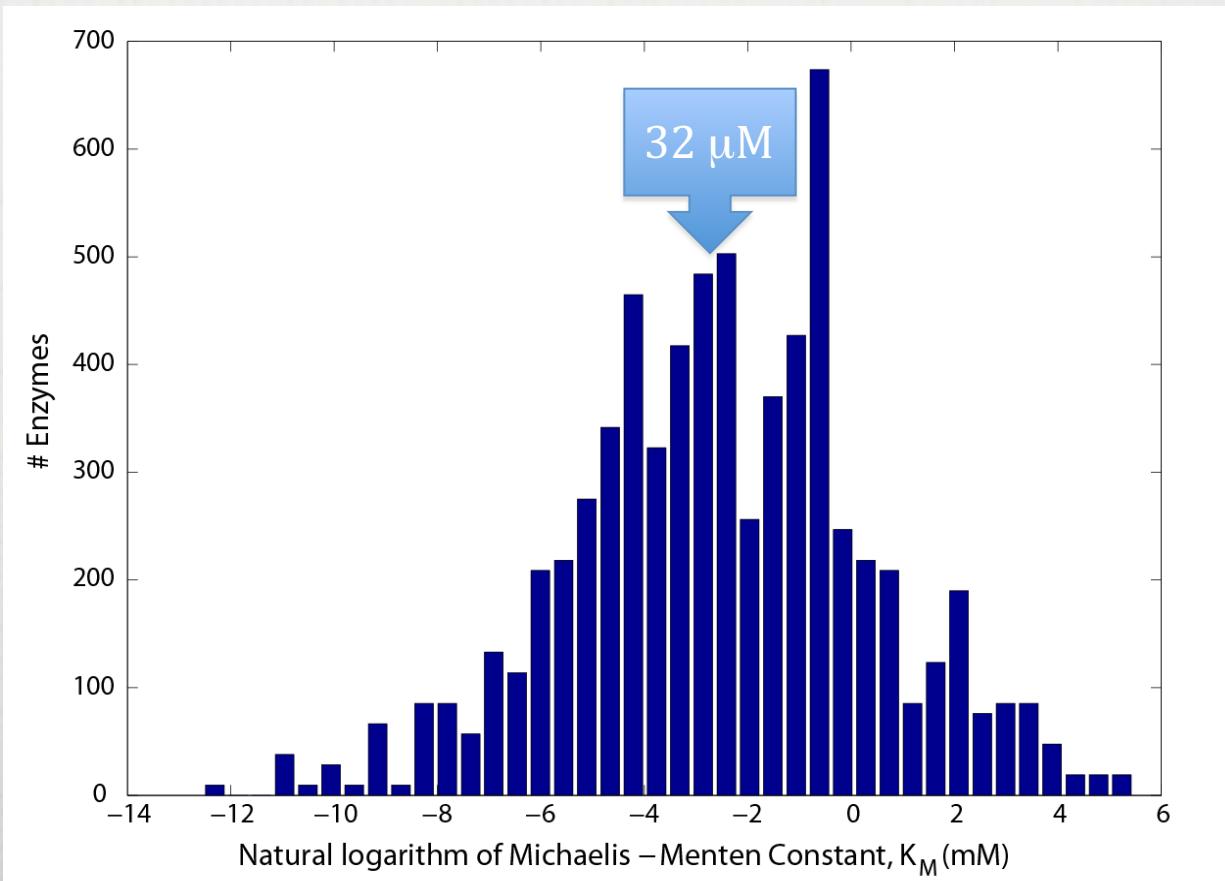
Distribution of Enzyme Kinetic Parameters

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

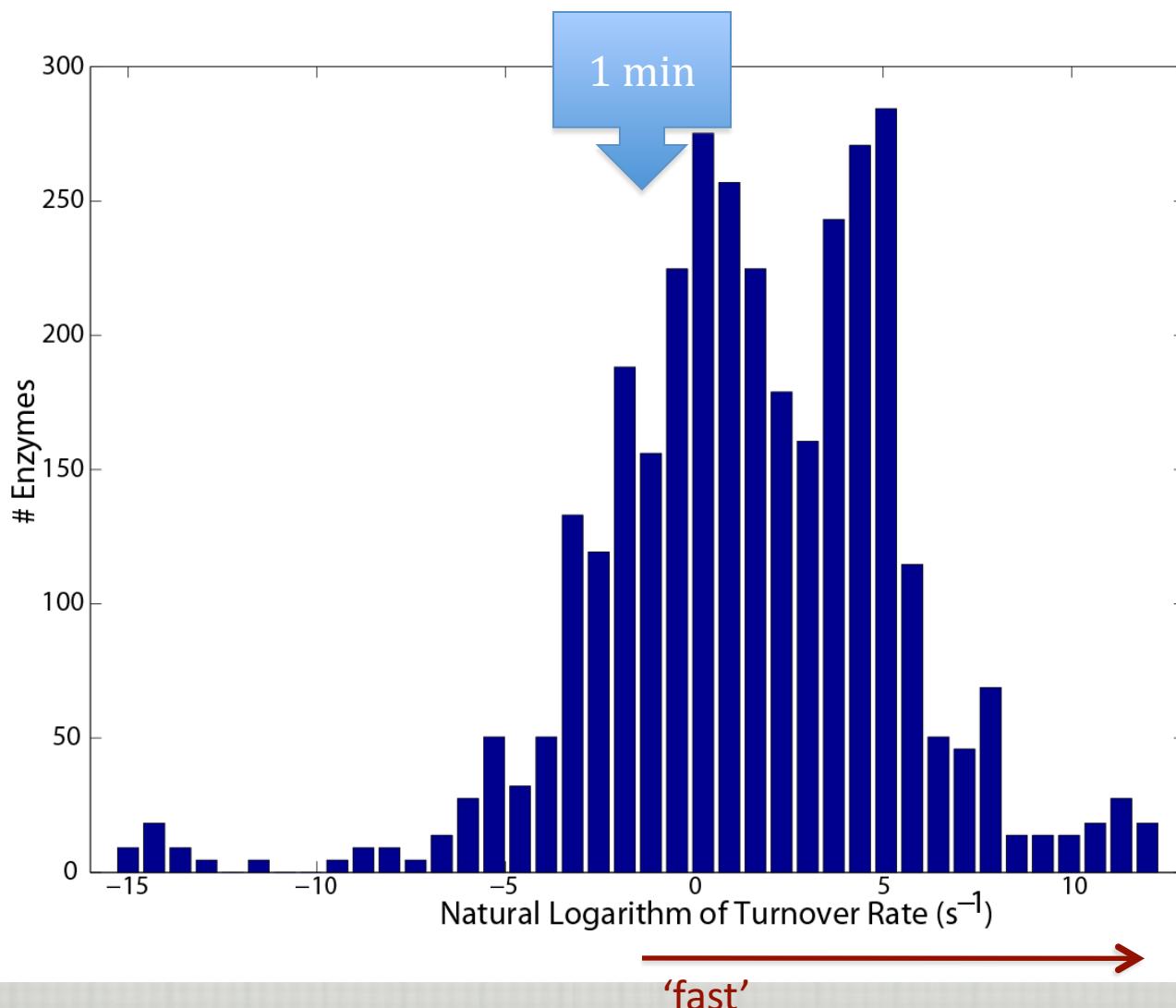


Kinetic Constants of *E. coli* Enzymes

- Average enzyme concentration is on the order of an average kinetic constant ($[E] \sim K_m$)

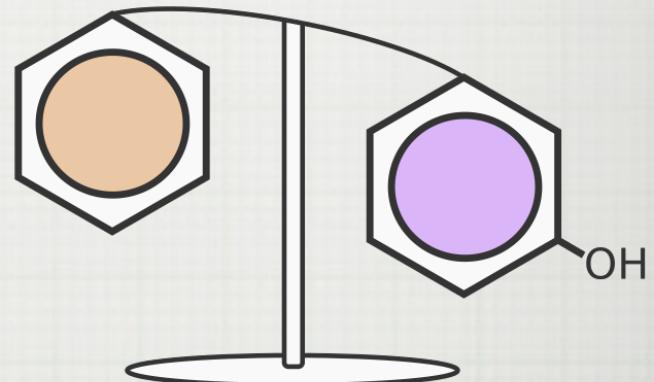


Typical Enzyme Turnover Times



eQuilibrator

- Thermodynamic analysis
 - Can account for different pHs and ionic strengths
- Estimation of
 - Gibbs Free Energy:
 - 1M at standard: ΔG°
 - 1mM at standard: ΔG^m
 - Nonstandard: $\Delta G'$
 - Reduction Potential
 - Equilibrium Constants



eQuilibrator

How much is a trillion dollars?

ORDER-OF-MAGNITUDE: AN EXAMPLE

\$1 Trillion & US Debt in Physical \$100 Bills



\$100 Dollars

DEMOCRACY.INFO
ECONOMIC INFOGRAPHICS

What does this mean for you and me?

The San Diego Mid-Coast Trolley Project

- Project Duration
 - Late 2016- Late 2021 (~5 years)
- Total Estimated Cost of Project:
 - \$2.1 billion
- Extension of Blue Line by
 - 11 miles

$$\frac{\$2.1 \text{ billion}}{11 \text{ miles}} \approx \$200 \text{ million per mile}$$



<https://www.keepsandiegomoving.com/Mid-coast/midcoast-FAQ.aspx>

<http://worldpopulationreview.com/us-cities/san-diego-population/>

https://www.sandag.org/uploads/publicationid/publicationid_2052_20893.pdf

The End