Pre-requisites:

Part II Thermal Statistical

Part II Soft Condensed Matter [or self study]

Reading list:

Phillips, Kondev, Theriot, Garcia

Physical Biology of the Cell - 2nd ed., Garland 2013

Phil Nelson

Physical Models of Living Systems - Freeman 2015

Biological Physics -: Energy, Information, Life - Freeman 2007

Uri Alon

An Introduction to Systems Biology - Chapman and Hall 2007 + 2nd ed 20/20

Bruce Alberts et al.

Molecular Biology of the Cell- Garland (many editions, updated almost yearly)

Kim Sneppen and Giovanni Zocchi

Physics in Molecular Biology - CUP 2005

Kim Sneppen

Models of Life - CUP 2014 & free e-book

Warning, this is a biology book

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lecture 1

Useful Information:

Website, linked from TiS:

http://people.bss.phy.cam.ac.uk/courses/biolectures/

Send comments & errors to df390, or even better on slack channel

Supervisions available for students in Part III and MASt (3 supervisions during teerm, 3 medium-size groups).

Times (TBD, but approximately every 2-3 modules)

We will provide written answers to the question sheet 24 hours before supervision.

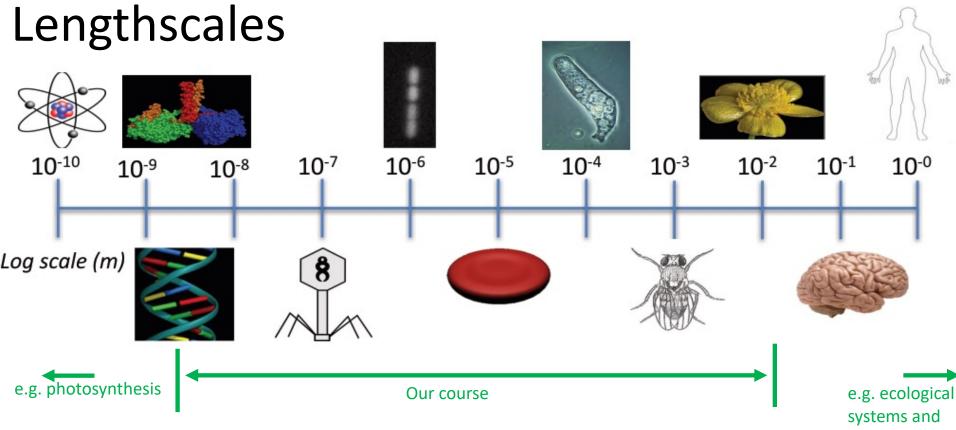
Subscribe to the Cambridge Centre for Physical Biology mailing list for seminars, PhD positions, and more...

Structure of the course:

24 lectures, in 7 modules:

- A context/overview/intro/basics, networks
- B evolution and growth of populations
- C dynamics in the cell
- D elements of neuro-physics
- E pattern formation in biology
- F protein production and regulation of gene expression
- G dynamical systems, switches and oscillations

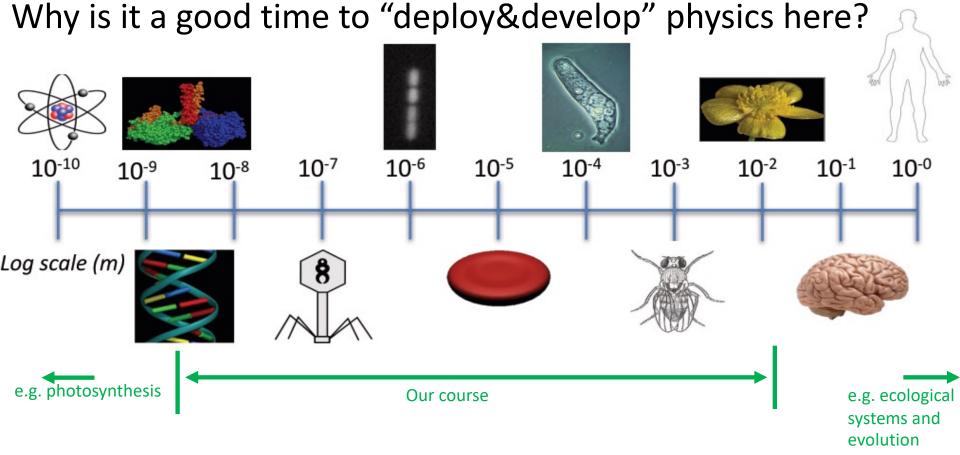
lecture 1



Biological systems have a hierarchical organization across many lengthscales $\stackrel{\text{evolution}}{\leftarrow}$ Lengthscale $\stackrel{\leftarrow}{\leftarrow}$ Timescale... hence "emergence".

Non-equilibrium (but considering separation of timescales, equilibrium often valid) Self assembly and self replication

We focus in this course on scales where thermal noise and small number noise are at play - classical statistical mechanics. $_{lecture\ 1}$



Fantastic detailed knowledge of the molecules that make up living systems from decades of "structural biology". Precise genetic code.

The broadly correct understanding of mechanisms of action of many of these constituents. Quantitative datasets resolved on relevant lengths & times.

Unique power of physics (stat mech, dynamical systems, soft matter) in linking up scales → models that have the "correct" mechanism, and that represent an understanding.

Physics is required.

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As in other fields (condensed matter, etc), what is our approach?

- Understand context here, cell biology context.
- Make order of magnitude estimates.
- Become familiar with tools for model building.
- Critical analysis to determine limitations, and suggest refinement to models.

lecture 1

Crick's legacy - Polymer Languages

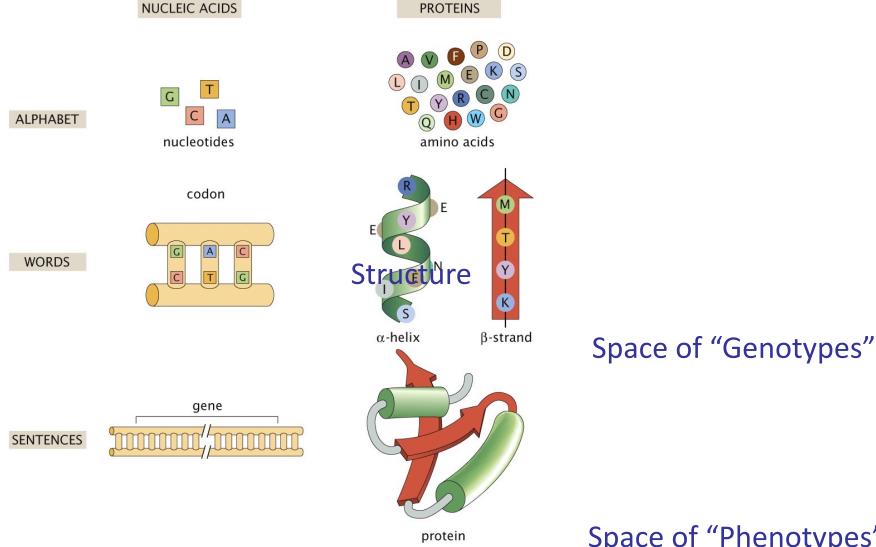
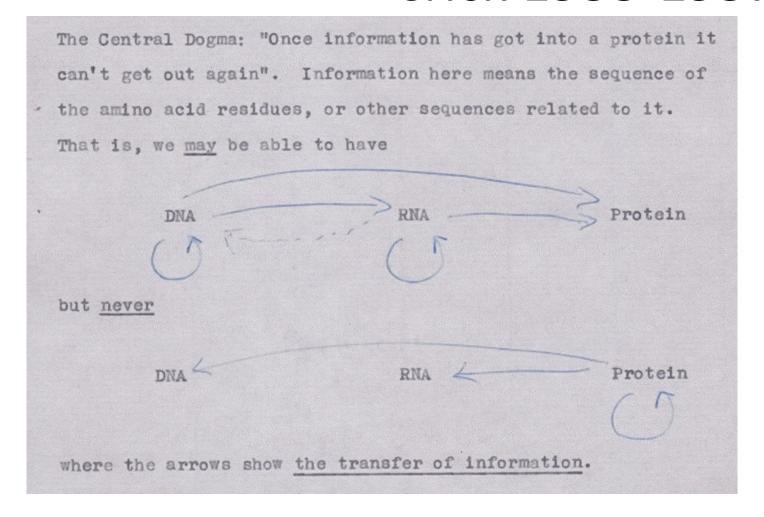


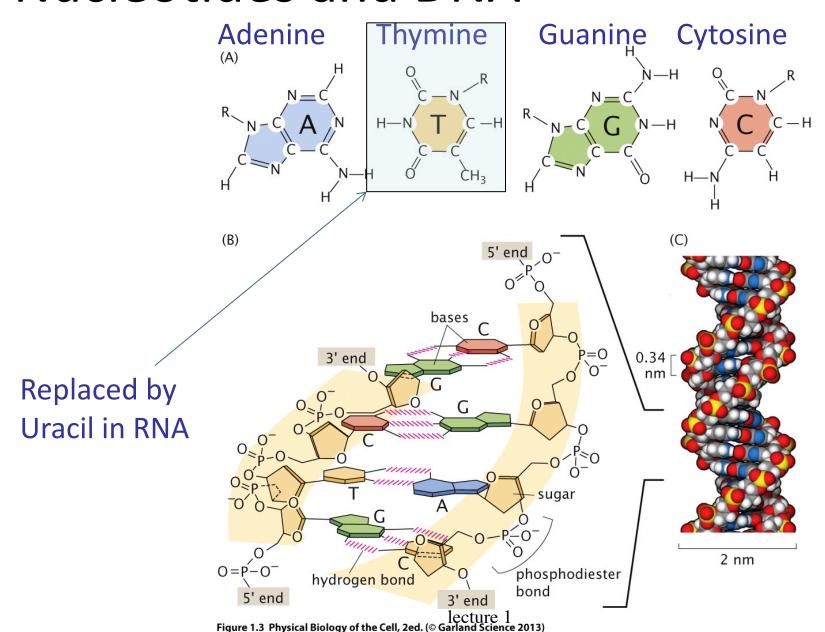
Figure 1.2 Physical Biology of the Cell, 2ed. (© Garland Science 2013)

Space of "Phenotypes"

Gene regulation: the "central dogma" Crick 1953-1957

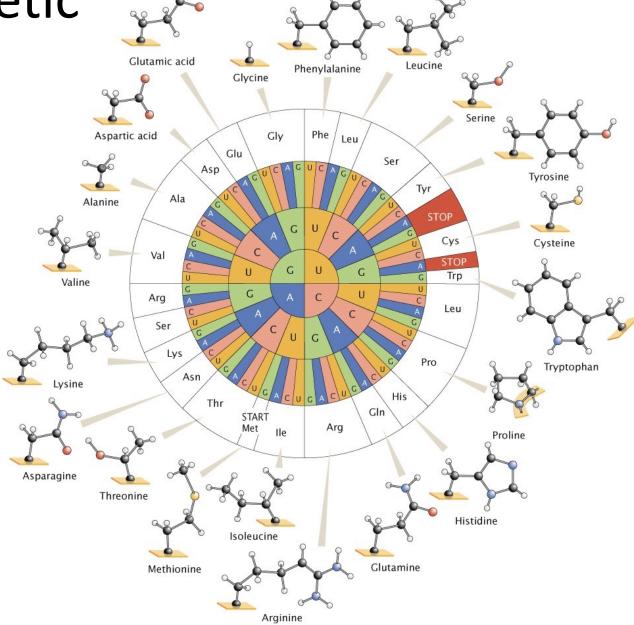


Nucleotides and DNA



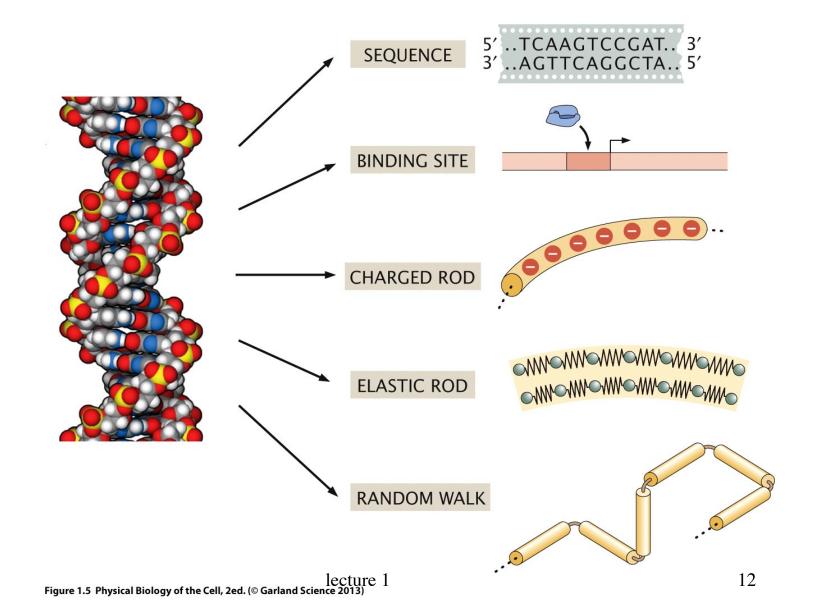
The genetic

code

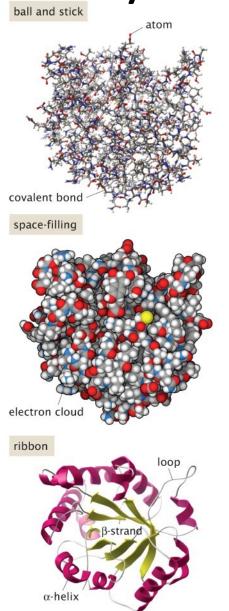


lecture 1 Figure 1.4 Physical Biology of the Cell, 2ed. (© Garland Science 2013)

Many ways to see a DNA double helix



Many ways to see a protein



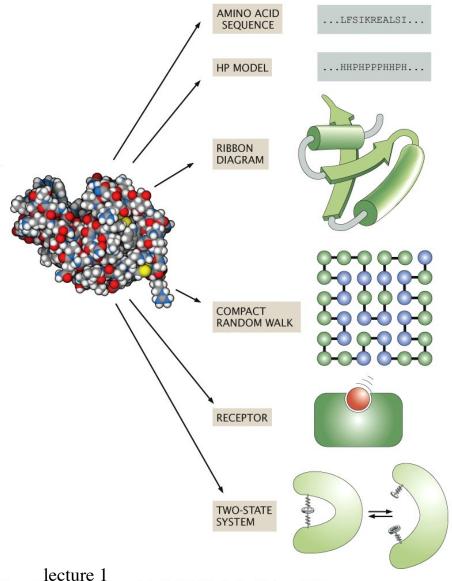
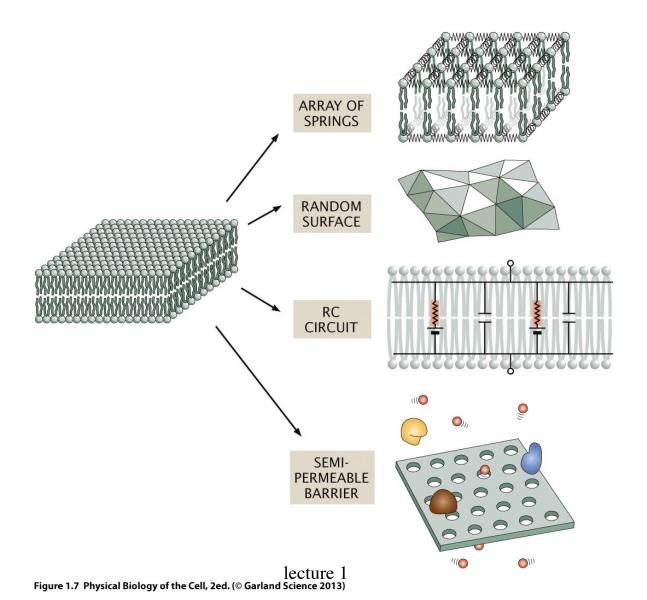


Figure 1.6 Physical Biology of the Cell, 2ed. (© Garland Science 2013)

Many ways to see a lipid membrane



To cells: Many ways to see a bacterium

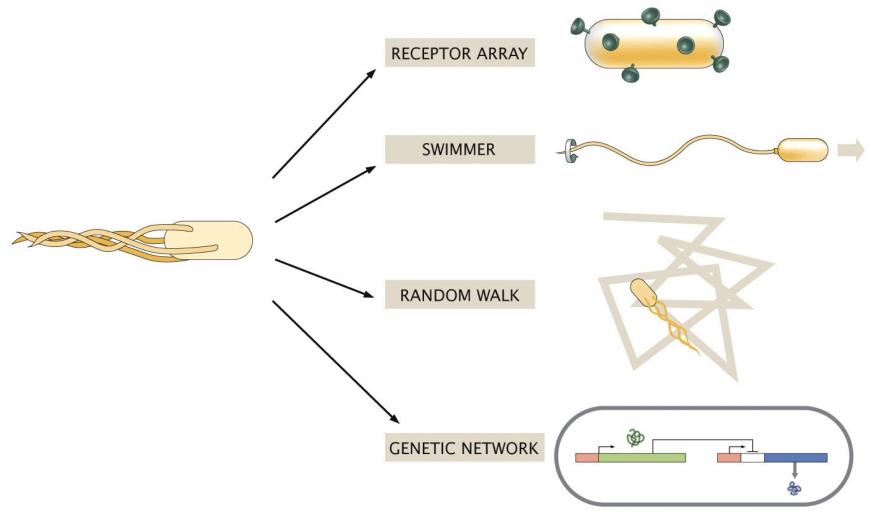
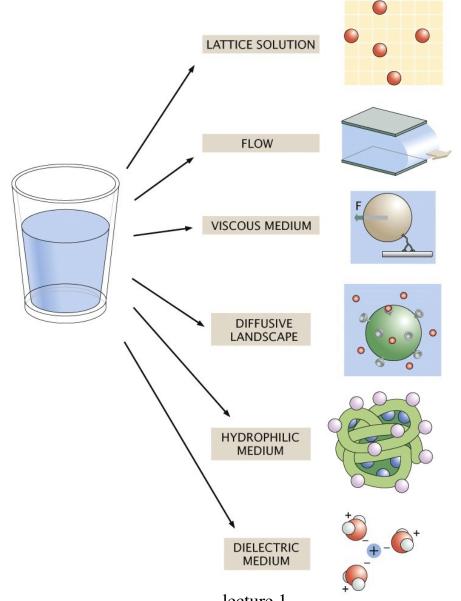


Figure 1.8 Physical Biology of the Cell, 2ed. (© Garland Science 2013)

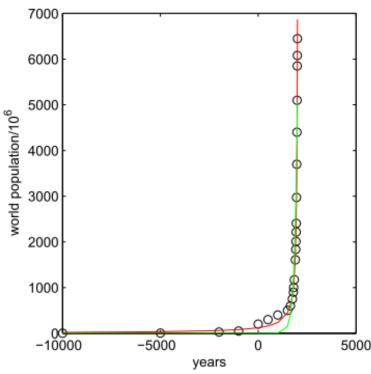
What is "right" level of description?



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What do we want to avoid?

Example of population growth data

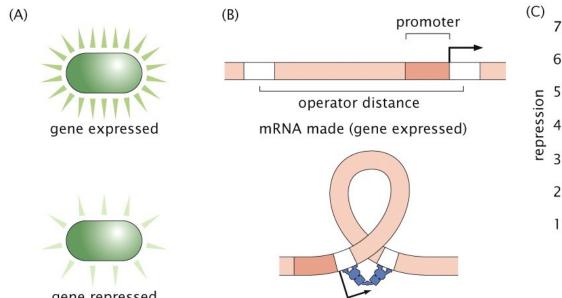


$$f(t) = a1/[a2 - (t/1yr)]$$
 a1=10000, a2=2050 works very well or $f(t) = a1 \exp (a2 t)$

Physical Models need to reflect a mechanism, and can point us to further key insights.

lecture 1

A success in quantitative biology (and still ongoing): The Lac repressor Where Stat mech and Polymer physics meet the biology of gene regulation



no mRNA made (gene repressed)

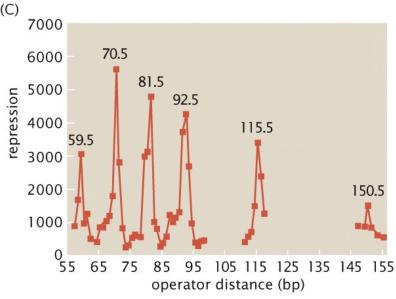


Figure 1.11 Physical Biology of the Cell, 2ed. (© Garland Science 2013)

Table 1.1: Rules of thumb for biological estimates.

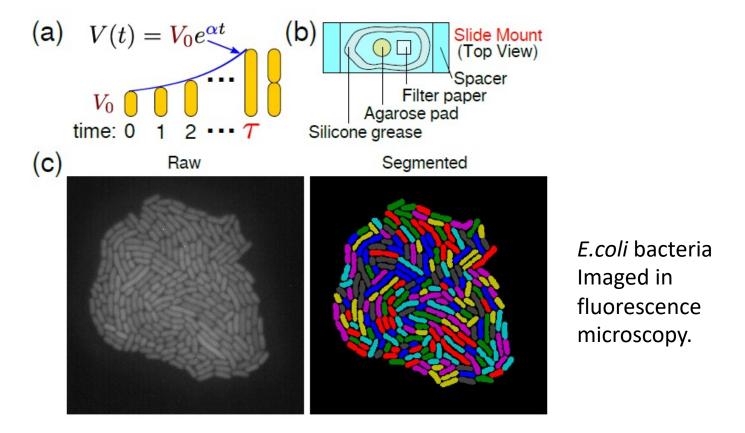
Table 1.1: Rules of thumb for biological estimates.					
	Quantity of interest	Symbol	Rule of thumb		
E. coli					
	Cell volume	$V_{E.coli}$	$\approx 1 \ \mu m^3$		
	Cell mass	m _{E. coli}	≈1 pg		
	Cell cycle time	t _{E. coli}	≈3000 s		
	Cell surface area	A _{E. coli}	$\approx 6 \mu m^2$		
	Macromolecule concentration in cytoplasm	cmacromol E. coli	≈300 mg/mL		
	Genome length	N ^{E. coli} bp	$pprox$ 5 $ imes$ 10 6 bp		
	Swimming speed	v _{E. coli}	${\approx}20\mu\text{m/s}$		
Yeast					
	Volume of cell	V _{yeast}	$pprox$ 60 μ m ³		
	Mass of cell	$m_{\rm yeast}$	≈60 pg		
	Diameter of cell	d_{yeast}	\approx 5 μ m		
	Cell cycle time	tyeast	≈200 min		
	Genome length	N ^{yeast} bp	$\approx 10^7 \mathrm{bp}$		
Organelles					
	Diameter of nucleus	d_{nucleus}	\approx 5 μ m		
	Length of mitochondrion	I _{mito}	\approx 2 μ m		
	Diameter of transport vesicles	d_{vesicle}	≈50 nm		
Water					
	Volume of molecule	V_{H_2O}	$\approx 10^{-2} \text{ nm}^3$		
	Density of water	ρ	1 g/cm ³		
	Viscosity of water	η	\approx 1 centipoise (10 ⁻² g/(cm s))		
	Hydrophobic embedding energy	$\approx E_{hydr}$	2500 cal/(mol nm ²)	19	
Table 1.1 (part 1 of 2) Physical Biology of the Cell, 2ed. (© Garland Science 2013)					

Table 1.1: Rules of thumb for biological estimates.

Table 1.1. Rules of thu	mb for biological estimates.		
	Quantity of interest	Symbol	Rule of thumb
DNA			
	Length per base pair Volume per base pair Charge density Persistence length	l _{bp} V _{bp} λ _{DNA} ξ _p	$\approx 1/3 \text{ nm}$ $\approx 1 \text{ nm}^3$ 2 e/0.34 nm 50 nm
Amino acids and			
proteins			
	Radius of "average" protein Volume of "average" protein Mass of "average" amino acid Mass of "average" protein Protein concentration in cytoplasm Characteristic force of protein motor Characteristic speed of protein motor Diffusion constant of "average" protein in cytoplasm	rprotein Vprotein Maa Mprotein Cprotein Fmotor Vprotein	≈2 nm ≈25 nm ³ ≈100 Da ≈30,000 Da ≈150 mg/mL ≈5 pN ≈200 nm/s ≈10 μ m ² /s
Lipid bilayers			
	Thickness of lipid bilayer Area per molecule Mass of lipid molecule	d $A_{ m lipid}$ $m_{ m lipid}$	$≈5 \text{ nm}$ $≈ \frac{1}{2} \text{ nm}^2$ $≈800 \text{ Da}$

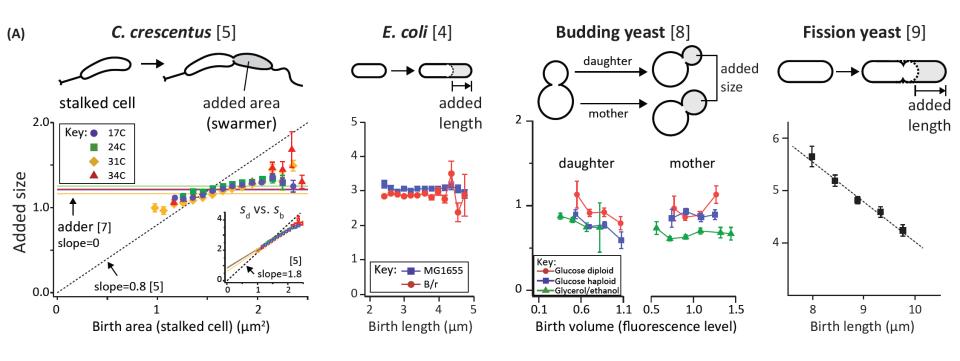
An example of "important question" that can be addressed in very different ways:

How do cells regulate division to have a mean size?



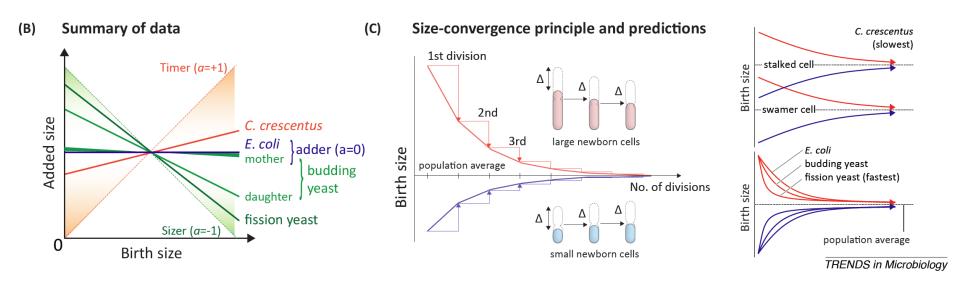
In principle, control could be through "sizer", "timer" or some combination.

Data on regulation of division



From S.Jun, S.Taheri-Araghi, Trends in Microbiology 4, 23 (2015)

How do cells regulate division to have a mean size?
One can also try to establish the general control theory, looking at the data.



One can search for the molecular mechanism, but certainly more complex than "the gene"!

Not so simple to come up with sizer mechanisms: plausible scenario put forward in yeast might involve sensing size through the balance between a species that has constant concentration in the cell volume as monomers, an adsorption equilibrium with the membrane (hence # prop to area), and a polymerisation "sink".

Concentration at the sink is then a membrane area sensor, triggering division.

Recap:

- Spirit and remit of this course.
- How physics contributes to this area of science.
- A first overview of cell machinery.
- Confidence in developing models and determining the right level of description.
- Next two lectures are "intro" to networks.

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