Bioengineering

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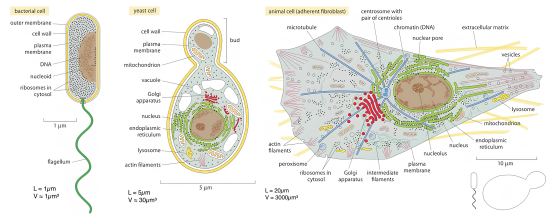
Tera	T	10^{12}	Kilo	k	10^{3}	Nano	n	10^{-9}
Giga	G	10^{9}	Milli	m	10^{-3}	Piko	p	10^{-12}
Mega	M	10^{6}	Mikro	μ	10^{-6}	Femto	f	10^{-15}

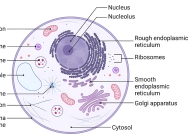
Orientation of the cell

Central Dogma of Molecular Biology

DNA $\xrightarrow{\text{transcription}}$ mRNA $\xrightarrow{\text{translation}}$ proteins

Cells:



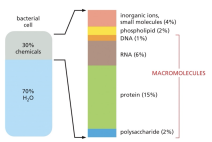


nucleus: houses DNA for EK
nucleolus: produces ribosomes/rRNA
mitochondria: cellular respiration (prod. ATP)
ribosome: produces proteins from mRNA transcripts

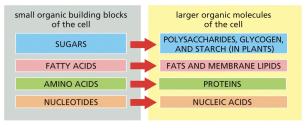
RER, SER and Golgi: involved in protein/lipid synthesis/processing
cytoskeleton: structure to cell, transport mol. in the cell or to enable the cell to move (cell migration)
centrosome: organizes microtubules during cell division allows the mother cell to split into 2 cells

Building Blocks

Cell composition:



Main building blocks:



- **Lipids** (fatty acids): long-term energy storage, cell membrane structure, signaling molecules.
- **Proteins** (amino acids): perform most of the cell's functions, including catalyzing reactions, signaling, and structural support. amino group $\mathbf{NH_2}$, carboxyl group **COOH**
- **Nucleic acids** (nucleotides): store and transmit genetic information (DNA, RNA), carry energy
- **Carbohydrates** (Sugars): short-term energy storages and for structural support.

Bounds

$$K_D = \frac{k_{off}}{k_{on}}$$

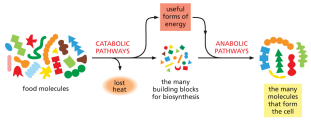
Covalent $\longleftrightarrow 100k_BT$
Ionic $\longleftrightarrow 1 - 10k_BT$
Hydrogen $\longleftrightarrow 1k_BT$
Van der Waals $\longleftrightarrow 0.1k_BT$
Electrostatic $\longleftrightarrow 0.1k_BT$

K_D: Equilibrium const.; indicates the ratio of free & bound units
k_{off}: Dissociation rate constant; inverse of time protein dissociates from the ligand
k_{on}: Association rate constant, speed of the reaction

Enzymes (aka catalysts)

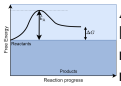
- Accellerate reaction by lowering the activation energy
- Are not consumed in the reaction
- Are specific to the reaction they catalyze
- Do not change the equilibrium point of the reaction.

Energy and Metabolism

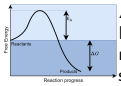


- **Catabolism:** Breakdown (glycolysis, citric acid cycle)
- **Anabolism:** Synthesis (protein synthesis, DNA replication)

Free Energy



$\Delta G > 0$
Endergonic
requires energy
not spontaneously



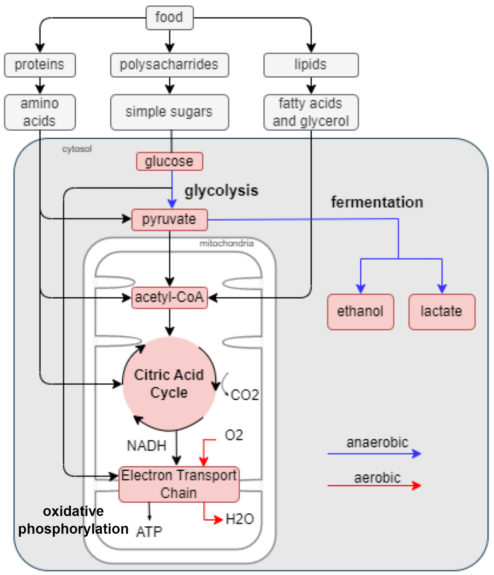
$\Delta G < 0$
Exergonic
releases energy
spontaneously

Redox

Oxidation loss of e^- $(H^+ + e^-)$ | Oxidation of glucose
Reduction gain of e^- $(H^+ + e^-)$| Reduction of pyruvate

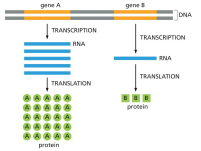
$\text{NADH} \rightleftharpoons \text{NAD}^+ + 2e^- + H^+$ | $\rightharpoonup$ oxidation, $\leftharpoondown$ reduction

Glucose Metabolism



Stage	Molecules	Glycolysis
Invest	2 ATP, 2 NAD^+, 1 glucose	
Payoff	4 ATP (2 net), 2 NADH, 2 H^+, 2 pyruvate	
Net gain	2 ATP, 2 NADH, 2 pyruvate	
Stage	**Molecules**	**Citric Acid Cycle**
Invest	2 Acetyl-CoA, 6 NAD^+, 2 FAD, 2 GDP (ADP)	
Payoff	6 NADH, 2 $FADH_2$, 2 GTP (ATP), 4 CO_2	
Net gain	2 ATP (GTP), 6 NADH, 2 $FADH_2$, 4 CO_2	
Stage	**Molecules**	**Oxidative Phosphorylation**
Invest	10 NADH, 2 $FADH_2$, 6 O_2	
Payoff	34 ATP, 6 H_2O	
Net gain	34 ATP, 6 H_2O	

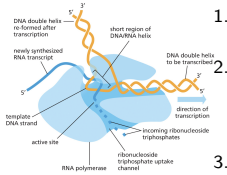
Biosynthesis



Genes are not always on and do not always produce the same number of transcripts or proteins.

Synthesis of proteins is a complicated and tightly regulated process.

Transcription

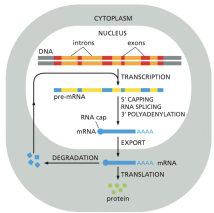


1. Small region of DNA opens and unwinds.
2. RNA polymerase (catalytic enzyme) interacts with one strand in the open region of DNA and adds ribonucleotides to grow an RNA polymer.
3. RNA polymerase further unwinds the DNA in the forward direction.

types of RNA

Type	Function
messenger RNA **mRNA**	code for proteins
ribosomal RNA **rRNA**	form core of ribosom structure and catalize protein synthesis
micro RNA **miRNA**	regulate gene expression
transfer RNA **tRNA**	serve as adaptors between mRNA and amino acids during protein synthesis
other non-coding RNA	RNA splicing, gene regulation, telomere maintenance, etc.

Eukaryotes vs Prokaryotes



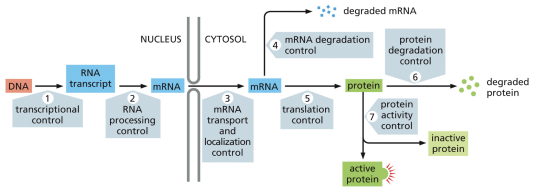
- Multiple types of RNA polymerases transcribe different classes of RNA.
- Transcription initiation is more complex.
- mRNA molecules undergo splicing, where introns (non-coding regions) are removed and exons (coding regions) are joined together to form the final RNA.

- 1 type of RNA polymerase transcribes all types of RNA.
- Transcription initiation is a simpler process.
- mRNA molecules are translated immediately after transcription

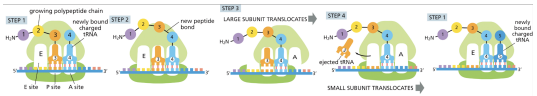
Both use gene regulatory proteins that bind to specific sequences of DNA:

- Repressors: Bind to sequences to turn genes off. Make it more difficult for RNA polymerase to bind to DNA.
- Activators: Bind to sequences to turn genes on. Make it more favorable for RNA polymerase to bind to DNA.

Gene Regulation



Translation by Ribosomes

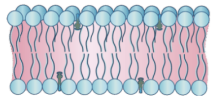


- On the mRNA, every three bases (= nucleotides) form one codon (needs ATP).
- A tRNA brings a matching amino acid. It has an anticodon that is complementary to the mRNA codon.
- The tRNA binds to the mRNA in the ribosome (at the A site).
- The amino acid is added to the growing polypeptide chain.
- The tRNA is ejected (from the E site), and the ribosome shifts forward by one codon.
- The process repeats until a stop codon is reached.

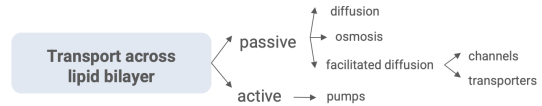
Multiple ribosomes can translate the same mRNA at the same time (making multiple proteins at the time), forming a polyribosome (or polysome).

Transport and Exchange

Cell Membrane



Phospholipid bilayer with hydrophilic head and hydrophobic tail act as barrier between compartements

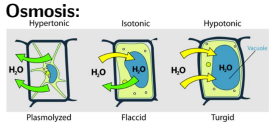


Passive Transport

Diffusion:
Along conzentration gradient, from high to low concentration.

$$\tau_D = \frac{l^2}{D} \quad D = \frac{k_B T}{6\pi\eta a}$$

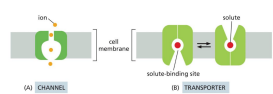
Osmosis:



Hypertonic: high solute concentration
Hypotonic: low solute concentration.

Membrane is **selectively permeable**, allowing some molecules to pass through while blocking others.

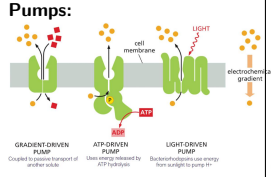
Facilitated Diffusion:



- selective to size and charge of solute
- bidirectional
- gated by external inputs
- Transporter good for large molecules

Active Transport

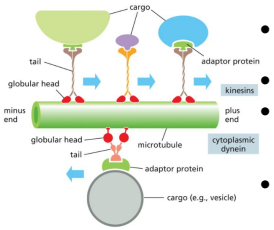
Pumps:



Vesicular Transport

Endocytosis: into cell
Exocytosis: out of cell
Pinocytosis: cell drinking
Phagocytosis: cell eating
Receptor-mediated endocytosis: specific uptake of molecules

Intracellular Transport



- chemical energy (ATP) to mechanical work
- directed motion
- Kinesin: carry out microtubule-based tiretrograde transport (towards cell edge)
- Dynein: carry out retrograde transport (towards cell nucleus)