Source: [KBBiologyMasterIndex]

1 | Bio-Molecules Quiz Review

#disorganized

1.1 | Paul's Review Sheet

... is here

1.1.1 | Carbohydrates

- Set 1, carbs. See Luke De's video + [KBhBIO101Carbs]
 - · Glucose vs. fructose both monosacharrides, one is a 6-carbon ring and one is a 5-carbon ring
 - Mono vs. di. vs. polysaccharide carbohydrates made out of a single, double, and multiple monomer (single-unit) carbohydrates
 - Starch vs. glycogen vs. cellulose lots of alpha glucose in less branches, lots of alpha glucose
 in more branches, lots of beta glucose in organized lattice respectively.
 - Starch plant food reserve
 - Glycogen animal energy reserve
 - Cellulose cell wall in plants
- Set 2, lipids. See Luke De's video + [KBhBI0101Lipids]
 - Triglyceride vs. fatty acid vs. phosophilid see [KBhBIO101StructuresofCarbs]
 - · Glycerol => a fatty acid
 - Triglyceride => three of 'em above
 - Phospholipid => two fatty acid + phosphate head
 - Saturated vs unsaturated fatty acids see also [KBhBIO101StructuresofCarbs]
 - Saturated Fats => no double bonds in the carbon chain of fatty acids think! butter
 - Unsaturated Fats => double bonds in the carbon chain of fatty acids think! olive oil
- Identify functional groups
 - Amino acid groups see [KBhBIO101AminoAcids]
 - carboxyl/carboxylic acid H-O-C=O (left side of backbone) Screen Shot 2020-10-12 at 2.29.28 PM
 - carbonyl C=O part of carboxyl
 - amide RC(=0)NR'R" (frequently shown in side chains of amino acids see Amine)
 - amino/amine H3N+ (right side of backbone)
 - hydroxyl OH group. Need I say more?
 - ester take a carboxylic acid and replace the hydrogen with anything else #ASK. => What
 join fatty acid chains with the glycerol to make trigrcyeride
 - ether R-O-R structure. => glycocidic bonds are formed by ether bonds
 - alcohol group (H-O-R) as part of the carboxyl
- Monomers vs Polymers [KBhBIO101StructuresofCarbs]
 - Monomer single molecule (such as a monosacchride) that could be chained together to make polymers
 - Polymers complex molecues built from monomers
 - Building polymers dehydration reaction taking out water molecules
 - Destructing polymers hydration reaction adding in water molecules

1.1.2 | Cell Structures

- · Prokaryotic vs. Eukaryotic
 - Prokaryotic cells often in single-cellular cells, has a cell wall, and contained in capsules
 - Eukaryotic cells in multicellular cell elements, contains a plasma membranes and nucleus
- Compare and contrast a typical animal cell with a typical plant cell. Be able to label diagrams of each. (See... problem set 1)
 - Animal Cell
 - · No cell wall
 - No chloroplast
 - · Has Cytoplasm
 - · Has Ribosomes
 - · Has Mitochondria
 - · No plastids organelle pigments
 - · Has Cilla Hair-like items on the outer surface
 - Plant Cell
 - · Has cell wall
 - · Has chloroplast photosynthesis
 - Has cytoplasm
 - · Has Ribosomes
 - · Has Mitochondria
 - · Has plastics organelle pigments
 - Mostly has no Cilla
- · Endosymbiotic theory
 - Endosymbiotic theory states that organelles within our current eukaryotic cells the mitochondria and chloroplasts — are originally prokaryotic cells in their own right. This is because they divide independently through binary fission, and also contains circular DNA that is independent of the main cell itself.
- Organizing organelles based on membranes #ASK
 - Membranous organelles possess own plasma => regulates own macromolecure consumption, hormones, etc. Perhaps original prokarotic cells
 - Endoplasmic reticulum => forms the network of transferring proteins and other elements
 - Golgi body/Gioli apparatus => packs, sorts, and modifies proteins and other elements throughout the cell
 - Non-membranous organelles does not posess own plasma => mostly part of the cytoskeleton
 of a cell
 - Ribosomes => protein synthesizer in the cell
 - Centrosome => forms flangella, cilla, and handles cells divisions
 - Lysomoes => digesting large nutrients and changing them to what cells could process and work on energy metabolism
 - Mitochrondria => store ATP and extract energy from ATP
 - Vacuoles => storing water, nutrients, waste
 - Plastids => creates colours displayed in the chromoplasts
- · Cell Components. Basicall all of these exist only in Eukareotic cells
 - · chloroplast and mitochondria
 - Chloroplast found in plants + does photosynthesis
 - Mitochondria found in animals + store ATP and extract energy from ATP
 - · cell wall and plasma membrane

- Cell Wall found in plants => surround the cell: hard
- Plasma membrane found in animals => surround the cell: soft |KBhBI0101Lipids|
- · rough endoplasmic reticulum (ER) and smooth ER
 - Rough ER covered by ribosomes and folds [KBhBI0101Proteins]
 - Smooth ER not covered by ribosomes and makes [KBhBI0101Lipids]
- cytosol, cytoplasm and cytoskeleton
 - Cytosol => liquid found inside cells; the "cytoplasm" floats within it
 - Cytoplasm => all the stuff within the cell
 - Cytoskeleton => complex network of proteins + fibres that organize the rest of the cell
- · nucleus and nucleolus
 - nucleus => centre of the cell. stores DNA
 - nucleolus => largest part of the nucleous that makes ribosomes
- · lysosomes and food vacuoles
 - Lysosomes => vesticles that contains enzymes that breaks down biomolecules
 - Food Vacoules => vesticels that stores food and other resources
- · cytoskeleton and microtubules
 - Cytoskeleton => complex network of proteins + fibres that organize the rest of the cell
 - Microtubulues => Polymers of tubulin protein that provides the main structure of eukarotic cells
- · flagella and cilia
 - Flagella => a bacteria's tail allow them to move and also act as an sensory organ. longer than a cilla, and moves in sinosoidial pattern.
 - Cilium => a cell's "hair" provides sensory and communications functions. Motil cilla could move about to "grab" things, and non-motile cilla can't move. more abundant that the flagella, and moves in circular pattern if they do move, and moves in circular pattern if they do move
- · Ribosomes and Golgi apparatus
 - Ribosomes => synthesizes proteins
 - Golgi apparatus => packs, modifying, and moving proteins

1.1.3 | Plasma Membrane Structure + transport

- Lipid structure and substructures: [KBhBIO101Lipids]
- · Functions of cell membrane
 - Phosophilid structures [KBhBIO101StructuresOfLipids]
 - Transmembrane proteins KbhBIO101CellTransport
 - Hydrophobic + hydrophillic parts of a phosophilid ||KBhBI0101StructuresOfLipids|| + ||KBhBI0101FluidMosaic||
- Passive + active transport KbhBIO101CellTransport
- Cell transport process
 - Simple diffusion (#ASK same thing as passive transport + osmosis)=> non-polar molecules needed "fall in"
 - Facilitated diffusion => specific polar molecules go along the gradient to get into the cell through transporter proteins
 - Phagocytosis => take a piece of the membrane with you to form a vesticle to introduce large solid elements, recycling the membrane after done — "cell eating"
 - Pinocytosis => take a piece of the membrane with you to form a vesticle to introduce large area
 of the "outside" in fluid and solid and all, recycling the membrane after done "cell drinking"
 - Endocytosis => Phagocytosis + Pinocytosis
 - Extocytosis => opposite of endocytosis

- Defining...
 - Isotonic => inside and outside have the same level of "osmolarity": probablility for osmosis to happen through a semipermiable membrane
 - Hypertonic => inside has less osmolarity than the outside: water/other elems will flow out of the cell
 - Hypotonic => outside has less osmolarity than the inside: water/other elems will flow into the cell

1.1.4 | Proteins Structures and Function

- Overall structure, monomers/building blocks, functions, and examples of proteins => |KBhBIO101Proteins|
- "peptide" => a chain of amino-acids
- · Polymerization via dehydration
 - Take two amino acids, take the H-O out of the alcahol, take the H out of the Amine. Fill the hole
 with the other one
- · Protein structure
 - Primary structure, secondary structure => [KBhBIO101Proteins]
 - Amino acids, N & C terminus => |KBhBIO101AminoAcids||. N terminus (Amine), C terminus (Carbolixic.)
 - Secondary structure H bonds between H-O, H-N
 - Tertiary structure => see the [KBhBIO101Proteins] articles
- The functions of proteins are varied because the primary sequence can be varied, effectively building any shape protein to do its specific function
- Form = function is the idea that the shape or form a protein takes through the combination of primary, secondary, tertiary, or quaternary structure determines how it will then function. Any changes to the structure will have some impact on its function and the more the structure is affected the more the function is likely to impacted
- Functions => defense, movement, structure, transport, cell to cell signaling, etc.

1.1.5 | Cell Structure

• Enzymes? [KBhBIO101Enzymes]

OK, so. Apparently Paul just answered the rest of his questions.

And I quote

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Enzymes are catalysts. They speed reaction rates but do not affect the change in free energy of the reaction (the difference in potential energy between reactants and products).

- · Activation energy is the amount of kinetic energy required to reach the transition state of a reaction.
- Enzymes speed up a reaction by lowering the activation energy, often with the help of cofactors or coenzymes.
- Enzymes lower the activation energy by some combination of...
 - · Orienting the reactions substrate(s) to promote more effective collisions (and therefore reactions
 - Stressing or straining bonds to temporarily and/or slightly lower the strength of attraction to allow the bond to break more easily

 Involving amino acid R-groups or sidechains in creating the transition state between reactants and products

Enzymes have active sites that bring substrates together and may change shape to stabilize the transition state; known as Induced Fit upon binding active site and slight change in enzyme shape.

Most enzymes are proteins, and thus their activity can be directly influenced by modifications or environmental factors, such as temperature and pH, that alter their three-dimensional structure.

Enzyme activity may be regulated/inhibited by molecules that compete with substrates to occupy the active site (competitive inhibitor) or alter enzyme shape so that substrates become unable to enter the active site (non-competitive inhibitor).

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1.2 | Helpful review items



Figure 1: Screen Shot 2020-10-09 at 11.58.55 AM.png

Figure 2: Screen Shot 2020-10-12 at 2.34.16 PM.png