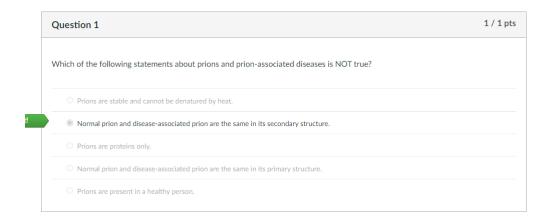
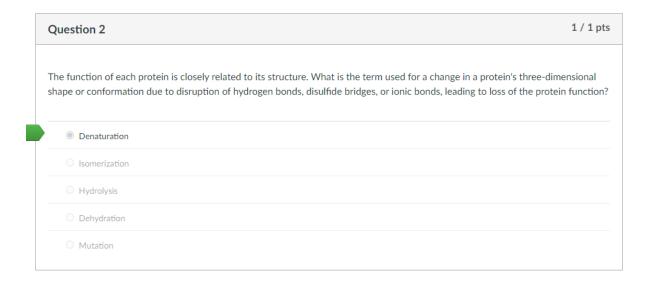
LSM 1301 - CA1 (Quiz Compilation)

Cell and molecular biology

Chemistry of Life



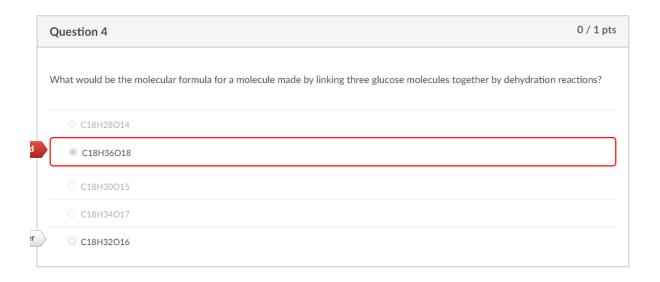
- normal prion and disease-associated prion are different in its secondary structure due to the way they fold
 - o normal alpha helical
 - diseased beta sheet
- disease-associated prion induces normal prion to transform



- denaturation disrupt bonds that hold the ternary structure together, changing the protein's shape
- hydrolysis destroys the whole polypeptide chain



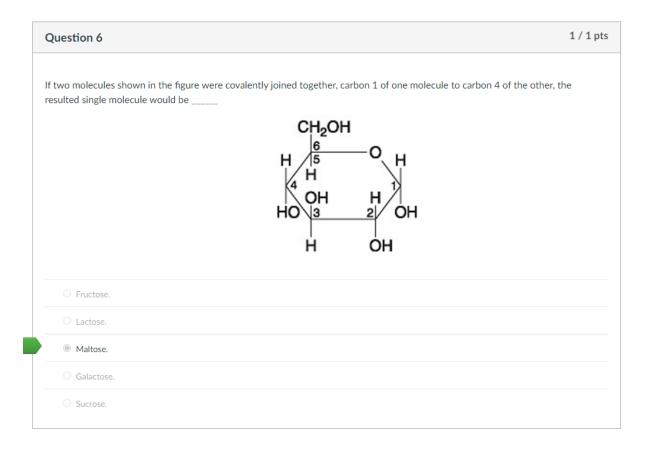
- glycogen question says dont want extra weight, so this option correct
- adipose tissue fat can store more energy than glycogen



- chem formula of glucose C6H12O6
- 3 glucose molecules joined together 3 x C6H12O6 2H2O (H and OH lost from forming 2 bonds) → G-G-G

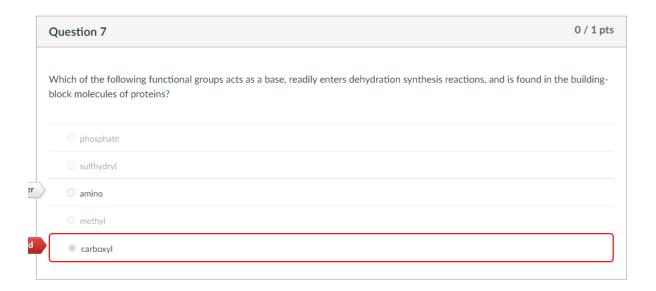
Question 5	1 / 1 pts
A quick scan of any health magazine would leave most people with the impression that cholesterol is always for you. Howe cholesterol is actually crucial to life:	ver,
As a source for producing vitamin D.	
As a keeper of fluidity in cell membrane at low temperature.	
As a material for synthesis of sex hormone.	
As a component of cell membrane.	
All of the above.	

- cholesterol
- keeper of fluidity in cell membrane at low temp sit in between phospholipids in the lipid bilayer to prevent them from clumping together at low temp
 - high temp decreases fluidity by pulling the phospholipids together

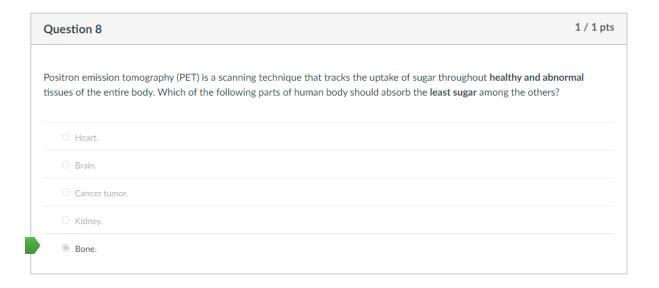


• maltose = glucose + glucose

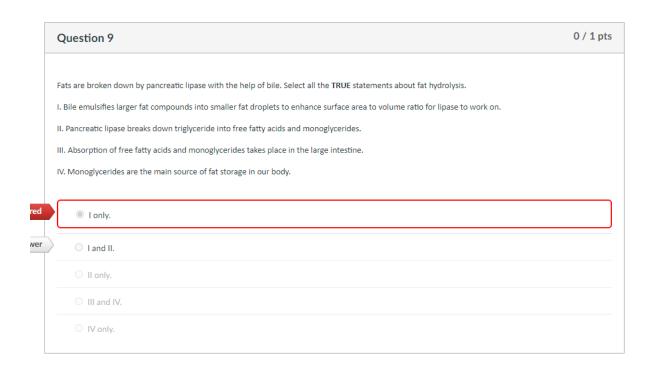
- fructose isomer of glucose (same chemical formula but lined up differently)
- lactose = galactose + glucose
- galactose = isomer of glucose
- sucrose = glucose + fructose



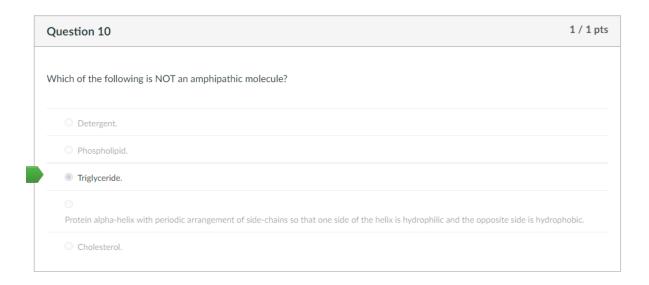
- base → positively charged
- · acid negatively charged
- amino positively charged (base)
- carboxyl and phosphate negatively charged (acidic)



 bone → structural material, dont have a lot of metabolic activities → use less sugar



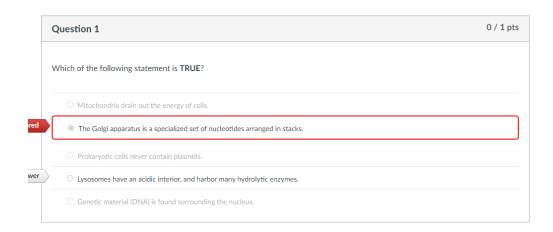
- bile salt emulsifies fat → increase surface area so lipase can work on them more efficiently
- pancreatic lipase breaks down the ester bond between glycerol and fatty acids
- · triglyceride are the main source of fat storage



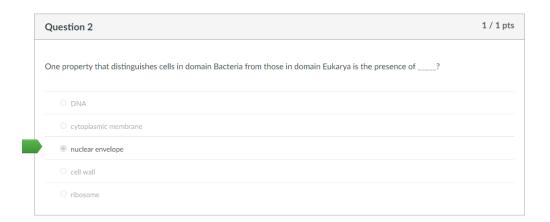
· amphipathic - both hydrophilic and hydrophobic

• triglyceride - hydrophobic

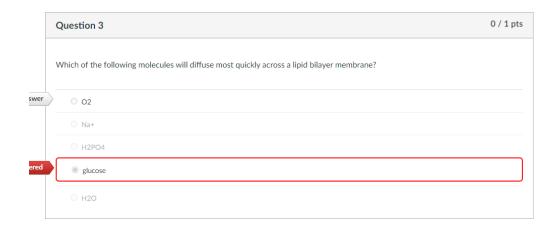
Cell Structure and Function



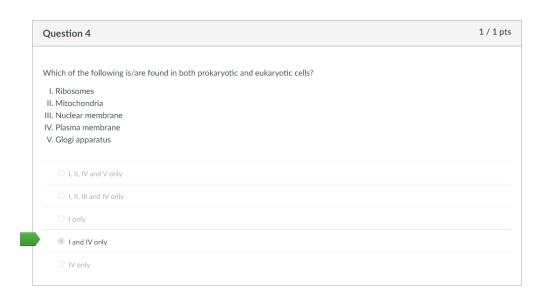
- · lysosomes stomach of cells, has a lot of enzymes to break down stuff
- no nucleotide in golgi apparutus
- bacteria (prokaryotic) has plasmid
- DNA in nucleus



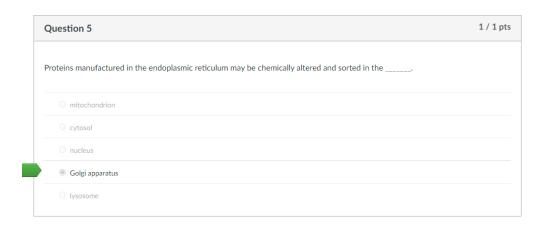
- presence of membrane-bound organelles
- cytoplasmic membrane cell membrane, exists in all cells including bacteria (prokaryotic) cells



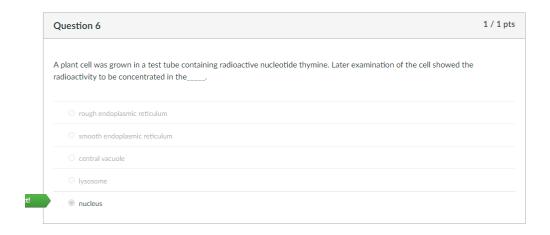
smaller and not charged → easier to diffuse across cell membrane



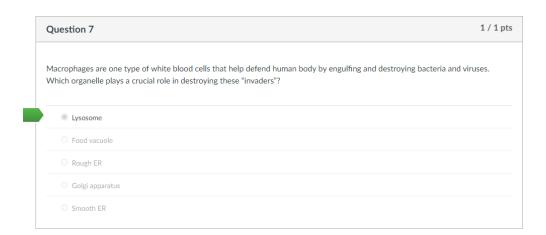
• prokaryotes dont have membrane-bound organells



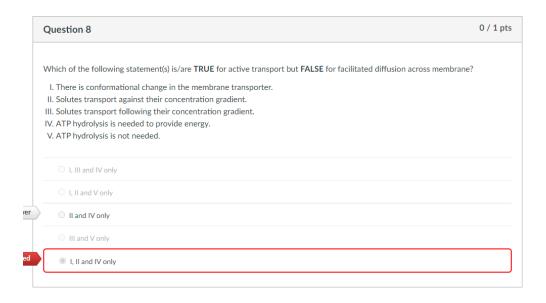
- after synthesis in rough ER, transported to golgi apparatus for modification (folding)
- cytosol fluid component of cytoplasm
 - cytoplasm includes everything in the cell except for the nucleus



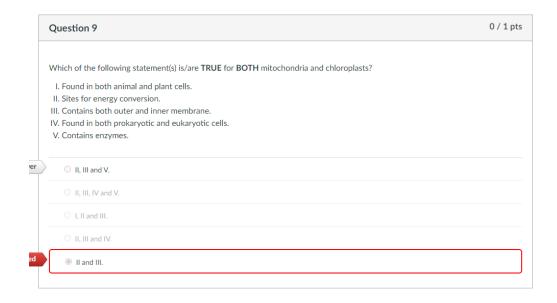
- · nucleotide found in nucleus
- rough ER protein synthesis and modification
- · smooth ER lipid and steroid synthesis



• lysosomes - carry acidic enzymes - invaders fed into them to destroy them

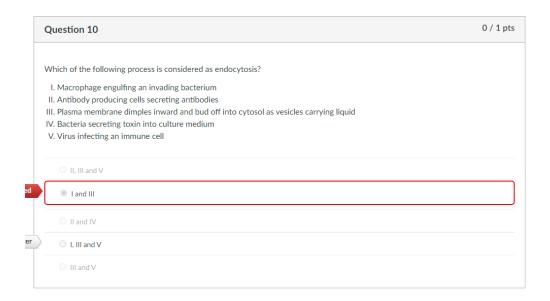


- conformational change needed for both active transport and facilitated diffusion
- active transport against concentration gradient
- · diffusion follows concentration gradient
- ATP (energy) needed for active transport to work but not facilitated diffusion



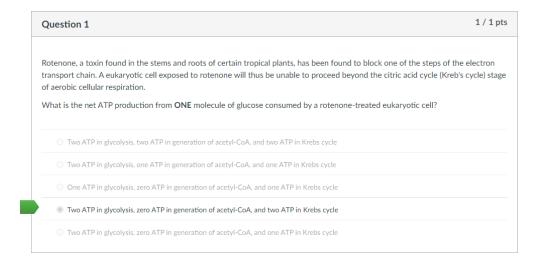
- · chloroplast only found in plant cells
- mitochondria and chloroplast membrane-bound organelles, cant be found in prokaryotic cells

 both contains enzymes (for activation energy) to carry out the energy conversion process

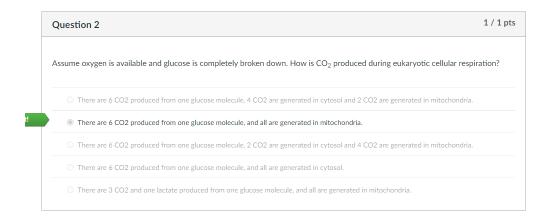


- endocytosis something entered the cell
- exocytosis something exited the cell

Energy of Life

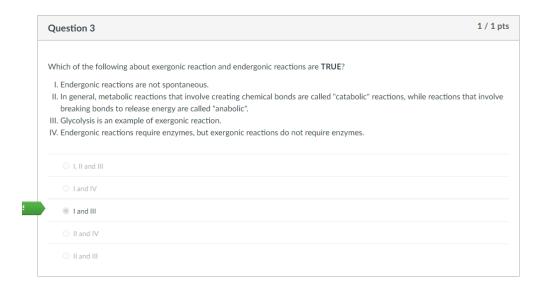


- net ATP production
- krebs cycle gives 1 ATP per cycle, 2 pyruvate = 2 ATP

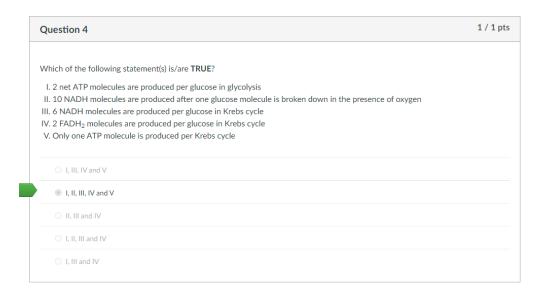


• in mitochondria

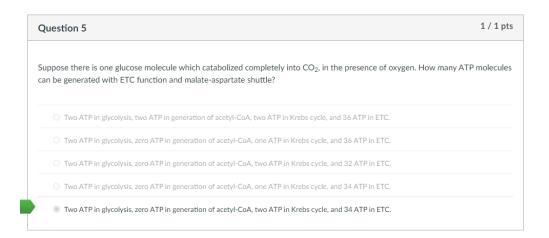
- 2 CO2 during acetyl coA
- 4 CO2 from kreb cycle



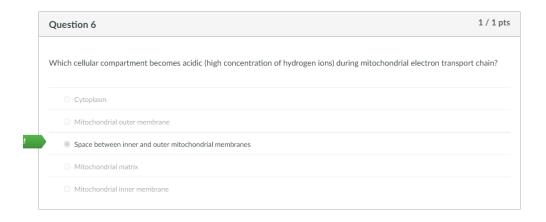
- endergonic reactions not spontaneous requires energy to start
- · catabolic reactions break down bonds
- anabolic create bonds
- glycolysis break down glucose to release energy (exergonic)
- both endergonic and exergonic require enzymes to lower activation energy to start reaction



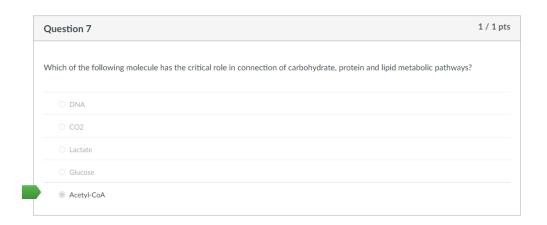
output of breaking down glucose for energy



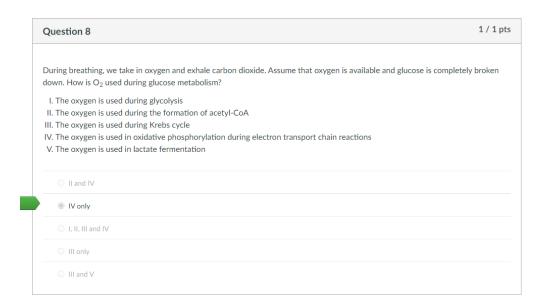
- malate-aspartate shuttle gives 38 ATP (34 in ETC)
- Glycerol-phosphate gives 36 ATP (32 in ETC)



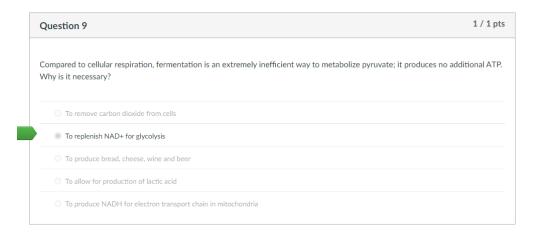
- · electron transport chain happens in intermembrane spaces in mitochondria
 - H+ high in concentration during this phase



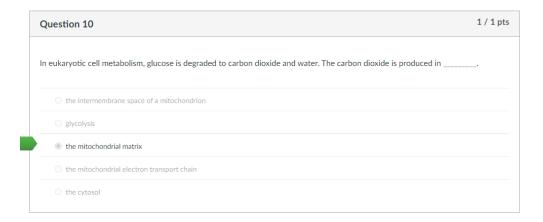
 carbs, protein, lipids can be converted into acetyl coA for energy conversion



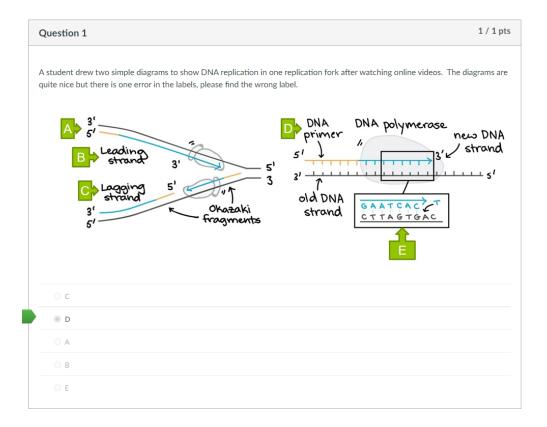
- oxidative phosphorylation
 - o phosphorylation ADP + P = ATP
 - oxidation H+(from NADH and FADH) + O = H2O



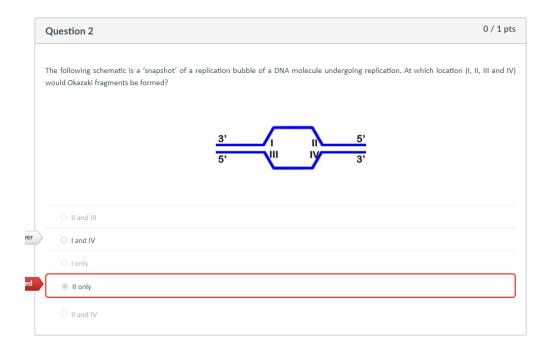
• glycolysis requires a lot of NAD+



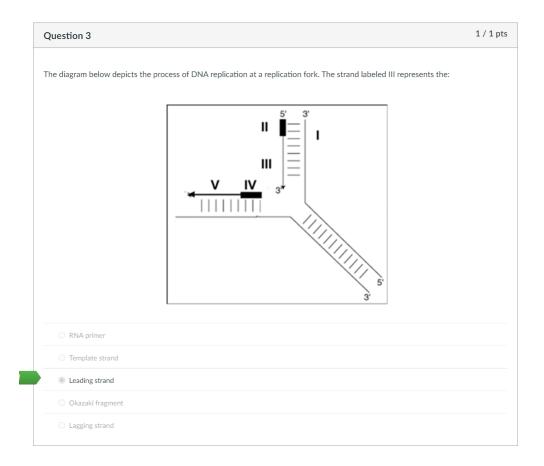
DNA and Heredity



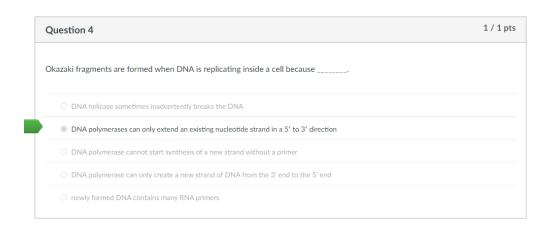
• D should be RNA primer



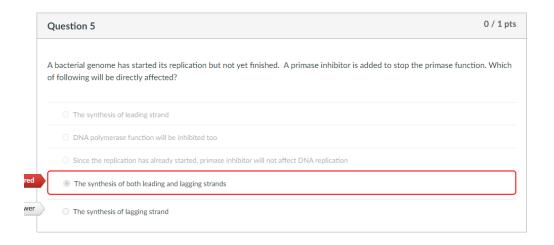
• look at direction of primer



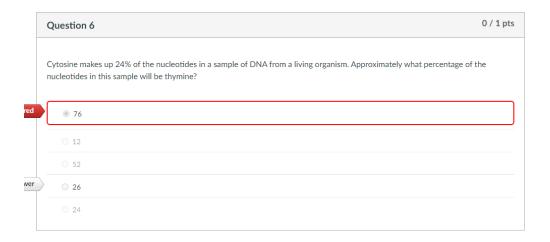
- III leading strand
- V okazaki fragment



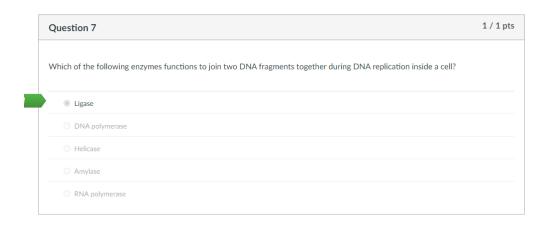
newly formed DNA contains many RNA primers - result of okazaki fragments



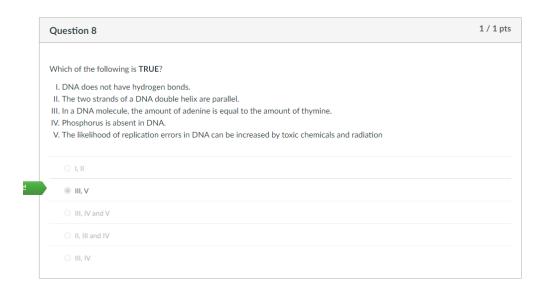
- lagging strand requires RNA primer multiple times for making okazaki fragments
- leading strand only need RNA primer once as it is continuous



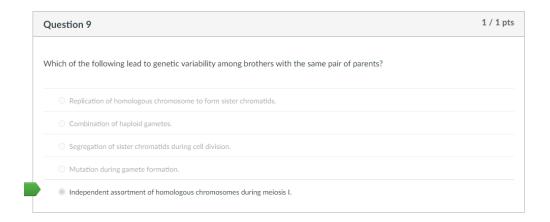
- C 24%, G 24% → C and G 48%
- A and T $52\% \rightarrow A / T 26\%$



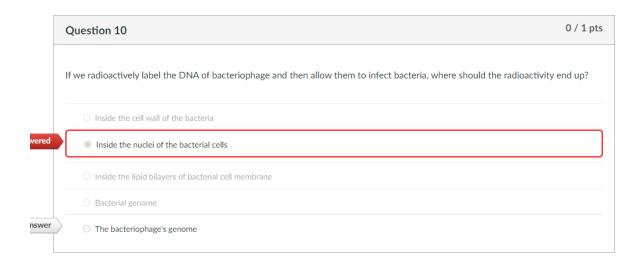
• ligase joins DNA fragments together by forming covalent bonds



- DNA strands are antiparallel
- Phosphorus (P) is present in the backbone of DNA

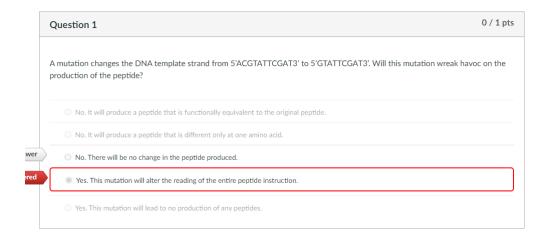


- same set of chromosomes can be combined differently different characteristics exhibited
- replication and segregation of sister chromatids are essential for cell division

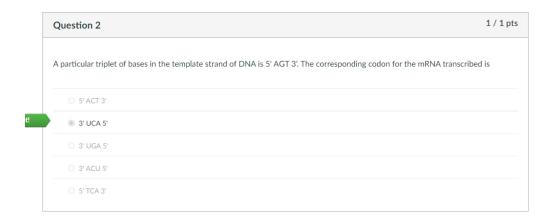


nuclei is another name for nucleus

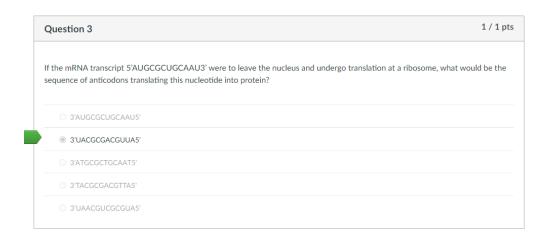
Gene Expression



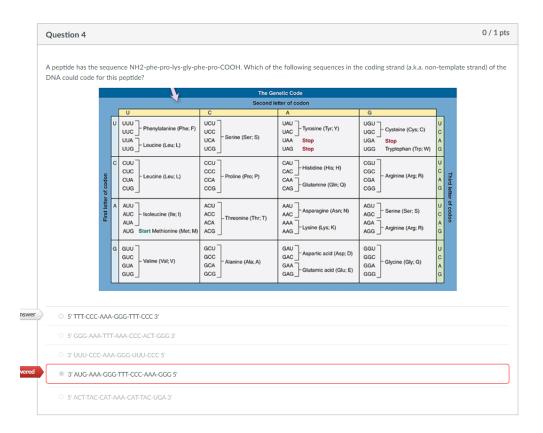
- transcription starts from 3' end, so resulting mRNA will start from 5'
- only the last 2 nucleotide of the template strand is gone, will not affect the reading of mRNA (at least the front part)
 - thus will not affect the peptide produced



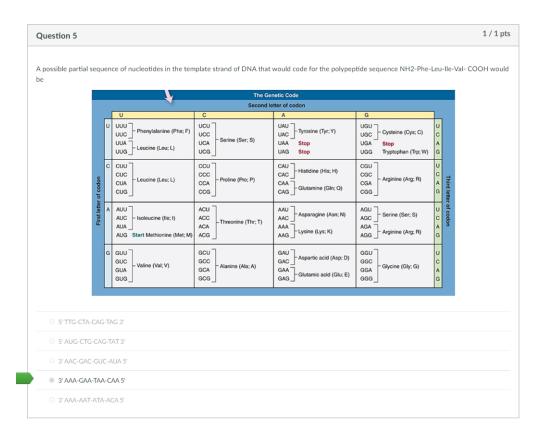
- codon will be antiparallel, so 3' to 5'
- RNA complementary nucleotides A-U, G-C
 - o note that A is paired with U and not T



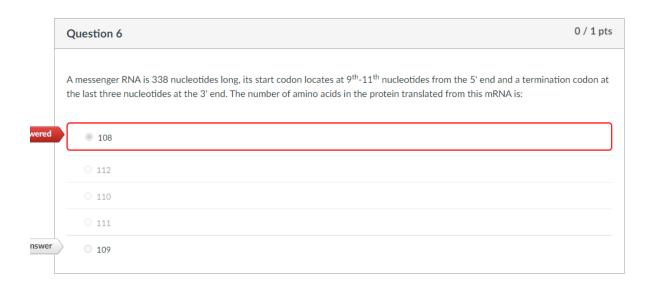
• anticodon complementary to mRNA



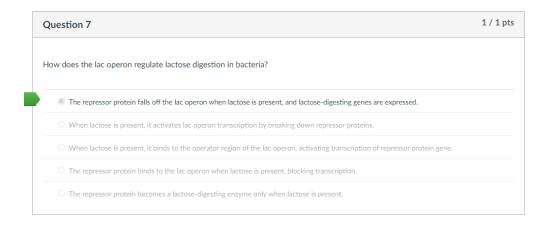
- phe UUU coded
- only TTT is complementary to AAA (complement of UUU)



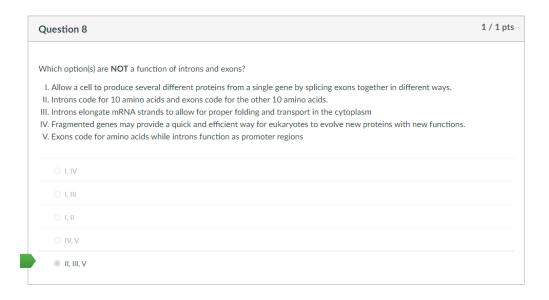
- NH2 5' end
- COOH 3' end



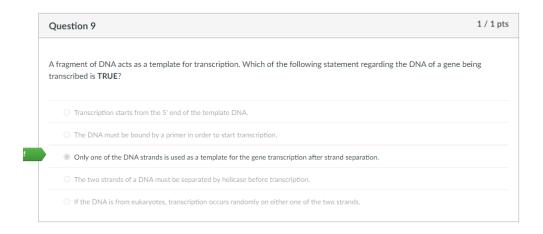
- 338 8 (before start codon) 3 (stop codon) 327
- 327 / 3 (3 nucleotide for 1 amino acid) = 109 amino acids



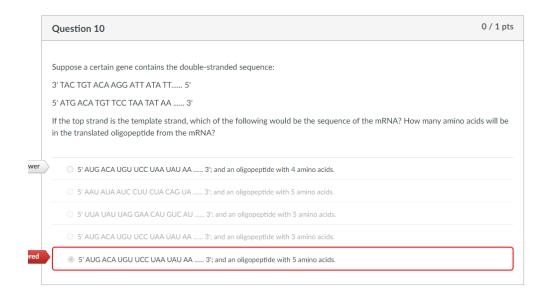
 repressor binds to lactose when lactose is present, allowing the DNA polymerase to work on lac operon to make the protein to digest lactose



- intron doesnt code for any amino acids
- intron doesnt elongate mRNA strands
- promoter region infront of where the gene for protein code starts
 - a region of DNA upstream of a gene where relevant proteins (such as RNA polymerase and transcription factors) bind to initiate transcription of that gene

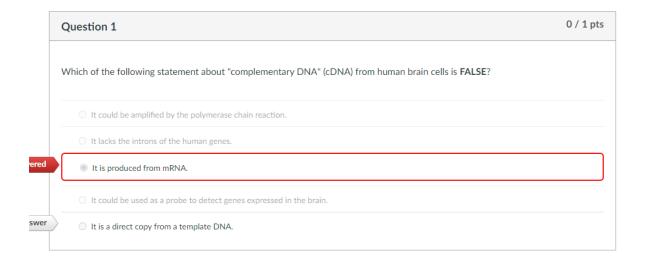


- transcription doesnt need helicase as RNA polymerase is able to unzip the DNA strands
- · transcription starts from the 3' end

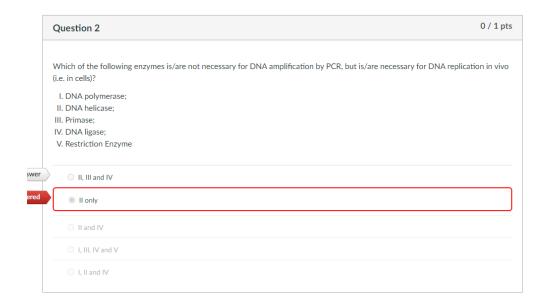


- UAA is stop codon, so only 4 amino acids will be created
 - o from AUG to UCC

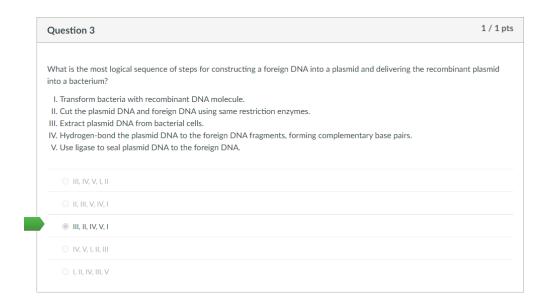
Biotechnology



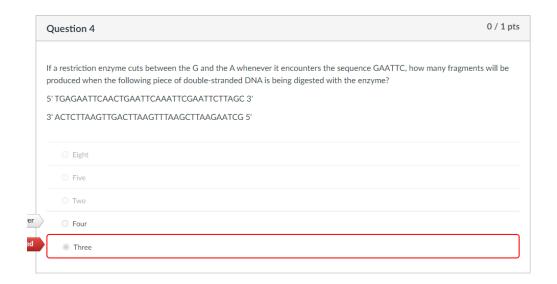
- cDNA is a direct copy OF template DNA, not FROM template DNA
- made from mature mRNA (free of introns)



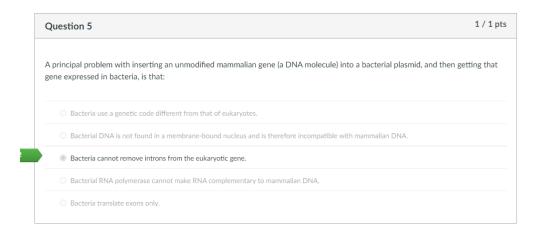
- polymerase needed for both replication and amplification
- helicase to open/unzip the DNA strands
 - needed for replication
 - not needed for amplification heat is used instead
- primase to generate primer
 - forward and reverse primer used in amplification, dont need primase
- ligase to join DNA fragments (okazaki fragments)
 - replication lagging strand generates okazaki fragments, need ligase to join them
 - amplification no need to join DNA fragments
- restriction enzyme used for gene editing, cleaves DNA at restriction sites to extract the gene there
 - not required in both replication and amplification



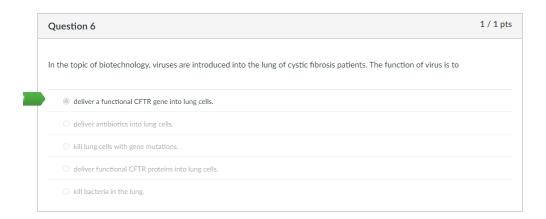
take out vector (3) → edit vector (2, 4, 5) → insert into target (1)



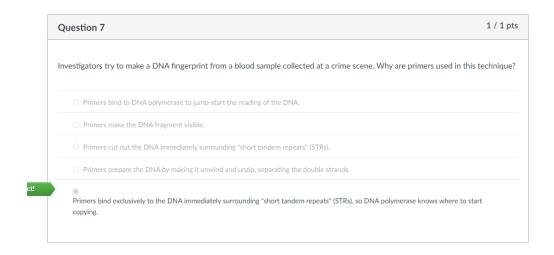
- read qn they asking for number of fragments;~;
- cut 3 times → 4 fragments



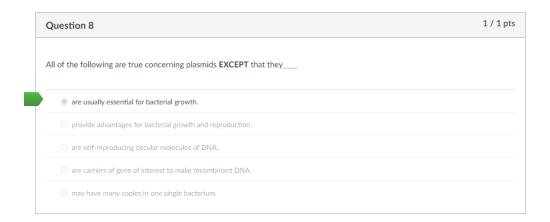
bacterial cant remove intron



• gene therapy - using virus to introduce normal/functional genes



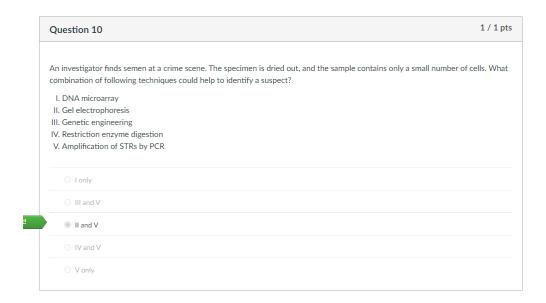
- specified primers designed to bind to regions around STRs to amplify them
- primers bind to DNA and not DNA polymerase



 plasmids are nonessential for growth but can be beneficial (e.g. antibiotic resistance)



- find the one with the highest number of complementary pairs
 - my answer is missing one G, between CC and AA



• gel electrophoresis - identify DNA present