

TOPIC VIII: CELL PHYSIOLOGY II

Learning Outcomes: Upon completion of Topic VIII (8), you should be able to

- a) Define the terms metabolism, anabolism and catabolism
- b) Define the terms transcription and translation.
- c) Briefly describe the process of transcription.
- d) Briefly describe the process of translation.
- e) Define the terms codon and anticodon.
- f) Explain the roles of tRNA and mRNA in protein synthesis.
- g) Describe the possible fates of proteins after their synthesis.
- h) Describe the characteristics of enzymes.
- i) Explain the role of ATP in the cell.
- j) Define the term cellular respiration.
- k) Explain the process by which glucose is converted through metabolic pathways to carbon dioxide and water (e.g. glycolysis, Krebs cycle and electron transport chain).
- l) Briefly describe the pathways by which fats and proteins can be used to synthesize ATP in a cell.

A) Overview

- Cells = living building blocks of body
- chemical processes in body (performed by cells) = metabolism
 - anabolism = building molecules
 - catabolism = breakdown of molecules

B) Basic Cell Processes:**1) Protein Synthesis:**

- proteins can be enzymes, structural etc
- process of protein synthesis involves:
 - a) DNA Transcription
 - DNA → mRNA (messenger RNA)
 - steps:
 - i) DNA uncoils at site of gene to expose gene base sequence
 - ii) RNA strand is formed, using code on DNA template to add complementary RNA nucleotides
 - e.g. DNA Template **A T C G C A**
 - mRNA **U A G C G U***
 - * in RNA – uracil (not thymine)

b) Translation

- mRNA \Rightarrow proteins

- steps:

- i) mRNA associates with ribosomes

- 3 bases on mRNA = base triplet = a codon (e.g. CUU, ACG, etc)

- each codon codes for 1 amino acid (aa's can have more than one codon)

- ii) transfer RNA (tRNA) with specific aa and anticodon

- (complimentary bases to codon) binds to matching RNA

- iii) peptide bond is formed to attach aa to growing peptide chain

- so overall: sequence of DNA bases determines mRNA sequence \rightarrow determines aa sequence for protein

c) After translation:

- proteins produced on free ribosomes - released into cytosol or go to nucleus or mitochondria

- proteins produced by ribosomes on ER (RER) - released into ER lumen

- modified by addition of sugars (glycoprotein)

- transferred to Golgi complex in vesicles

- Golgi complex:

- i) further modifies protein by

- carb changes/additions (act as an address to send them to the right destination)

- lipid additions

- ii) packages modified proteins (in vesicles) and

- sends to destination: cell membrane, secretion, lysosomes

- § Lysosomes = membrane-bound organelles containing digestive enzymes (acid environment)

- Enzymes: (names end in ase)

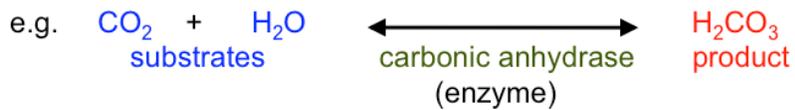
- increase reaction rates

- not used up in reaction

- very sensitive to pH, temperature

- synthesis controlled by cell (proteins) – cell can vary activity

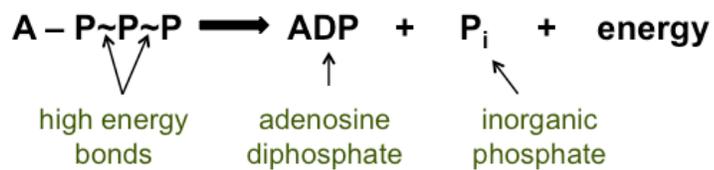
- very specific



* this reaction can go in both directions – some can't

2. ATP Catabolism

- Adenosine Triphosphate (ATP)
 - adenine + ribose (= adenosine) + 3 phosphates
 - energy stored in phosphate bonds (covalent)



- energy produced used for reactions (protein synthesis, active membrane transport, muscle contraction etc)

- little ATP is stored, so constantly have to make it

3) Cellular Respiration:

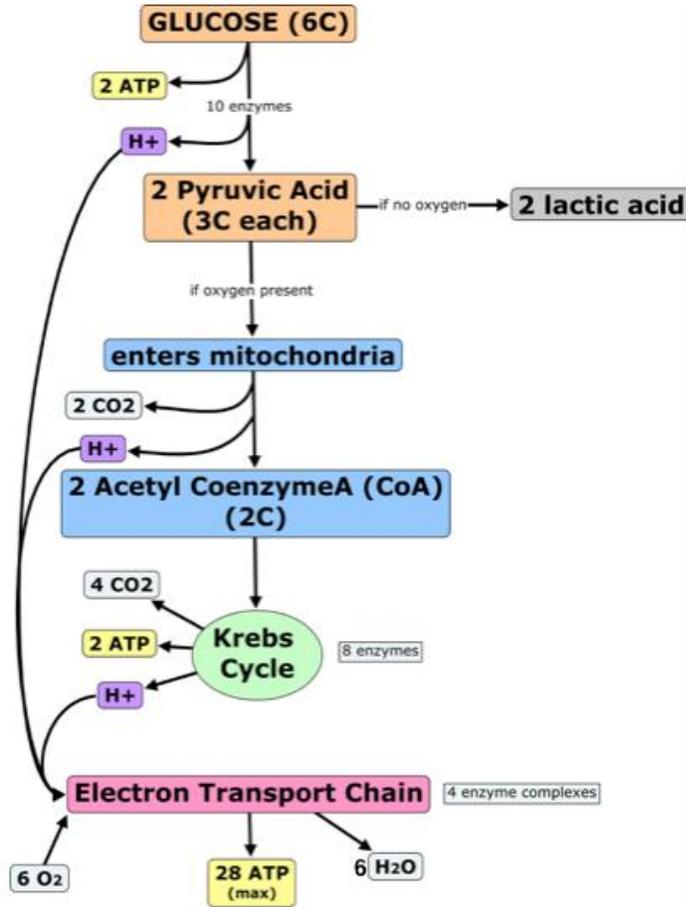
- production of ATP using glucose
 - glucose enters most cells by facilitated transport (may be ↑ in some cells by insulin)

- overall:



- steps:

- a) glycolysis (anaerobic) in cytosol
- b) enters mitochondria – becomes aerobic
- c) Krebs Cycle
- d) Electron Transport Chain



Glycolysis (glucose to pyruvic acid)

- In the body:

a) glycogen → glucose → ATP

b) proteins

- some aa can be converted to pyruvic acid or enter Krebs cycle
- depending on body's need may form new glucose (liver, kidney) or ATP (most cells)

c) fats

- primary storage form of energy in the body (triglycerides)
- broken down to form acetyl CoA → ATP