

## NOTES: CH 5, part 2: Proteins and Nucleic Acids

### 5.4 - Proteins have many structures, resulting in a wide range of functions

- Proteins account for more than 50% of the dry mass of most cells
- Protein functions include:
  - \_\_\_\_\_
  - \_\_\_\_\_
  - \_\_\_\_\_ / \_\_\_\_\_
  - \_\_\_\_\_
  - \_\_\_\_\_
  - catalysis of reactions (\_\_\_\_\_)
- an enzyme is a type of protein that acts as a catalyst, \_\_\_\_\_

### PROTEINS:

- consist of 1 or more polypeptide chains folded and coiled into specific conformations
- ***Polypeptide chains*** = \_\_\_\_\_
  - arranged in \_\_\_\_\_
  - linked by \_\_\_\_\_
  - range in length from a few to 1000+

### AMINO ACIDS:

- \_\_\_\_\_
- structure of an amino acid (SKETCH):
  - \_\_\_\_\_
  - \_\_\_\_\_
  - \_\_\_\_\_
  - \_\_\_\_\_
  - variable R group (side chain)

- Cells use \_\_\_\_\_ to make thousands of proteins
- linked together by \_\_\_\_\_ (links the carboxyl group of 1 amino acid to the amino group of another; requires dehydration / condensation)

### PROTEIN STRUCTURE:

- a protein's \_\_\_\_\_ (3-D shape)

### Four levels of protein structure:

#### 1) Primary structure:

- \_\_\_\_\_ (forms the "backbone" of a protein)
  - determined by \_\_\_\_\_
  - a slight change can affect a protein's conformation and function (e.g. sickle-cell hemoglobin)
  - can be sequenced in the laboratory (insulin)

## 2) Secondary structure:

- \_\_\_\_\_ of a protein's polypeptide backbone
- stabilized by H-bonds between peptide linkages \_\_\_\_\_ (NOT the amino acid side chains)

**ALPHA HELIX** = \_\_\_\_\_ stabilized by H-bonding between every 4th peptide bond (3.6 amino acid/turn)

**BETA PLEATED SHEET** = sheet of antiparallel chains folded into " \_\_\_\_\_ "

## 3) Tertiary Structure:

- irregular contortions of protein due to bonding \_\_\_\_\_ (R groups) \_\_\_\_\_

## 4) Quaternary Structure:

- association of 2 or more protein subunits to form a single functioning molecule (i.e. \_\_\_\_\_ and \_\_\_\_\_)

## Protein Form and Function

- A functional protein consists of one or more polypeptides twisted, folded, and coiled into a unique shape
- The sequence of amino acids determines a protein's three-dimensional formation
- **A** \_\_\_\_\_

## Protein Conformation

- determined by physical and chemical environmental conditions
- **DENATURATION**: process that alters a protein's native conformation and hence its biological activity
  - \_\_\_\_\_
  - \_\_\_\_\_
  - chemical agents that disrupt H-bonding
  - transfer to an organic solvent
  - \_\_\_\_\_

## **Protein Denaturation:**

- **A denatured protein is misshapen and therefore biologically inactive**

## The Protein-Folding Problem

- It is hard to predict a protein's conformation from its primary structure
- Most proteins probably go through several states on their way to a stable conformation
- Chaperonins are protein molecules that assist the proper folding of other proteins

## 5.5 - NUCLEIC ACIDS!!

### **5.5 - Nucleic acids** \_\_\_\_\_

- The amino acid sequence of a polypeptide is programmed by a unit of inheritance called a \_\_\_\_\_
- \_\_\_\_\_, a nucleic acid

• **Two types of nucleic acids:**

1) DNA

2) RNA

**The Roles of Nucleic Acids**

- There are two types of nucleic acids:

- \_\_\_\_\_ (DNA)

- \_\_\_\_\_ (RNA)

- DNA directs synthesis of messenger RNA (mRNA) and, through mRNA, controls protein synthesis
- Protein synthesis occurs in \_\_\_\_\_

1. **DNA = Deoxyribonucleic acid**

- encodes the instructions for amino acid sequences of proteins
- is copied and passed from one generation of cells to another

2. **RNA = Ribonucleic acid**

- functions in the actual \_\_\_\_\_ coded for by DNA
- carries the encoded information to the ribosomes; carries the amino acids to the ribosome; a major component of ribosomes

\_\_\_\_\_ → \_\_\_\_\_ → \_\_\_\_\_

**Structure of Nucleic Acids**

- polymers of monomers called \_\_\_\_\_
- Each nucleotide consists of:

Sketch one nucleotide in the space below:

1. **Pentose** (5-carbon sugar)

- \_\_\_\_\_  
- \_\_\_\_\_

2. **Phosphate group** (attached to #5 carbon on sugar)

3. **Nitrogenous base**

- purines (double ring)
- pyrimidines (single ring)

- The portion of a nucleotide without the phosphate group is called a \_\_\_\_\_
- nucleotides are joined together by \_\_\_\_\_ (between phosphate of one nucleotide and the sugar of the next)
- results in a backbone with a repeating pattern of sugar-phosphate-sugar-phosphate...

**The DNA Double Helix**

- A DNA molecule has two polynucleotides spiraling around an imaginary axis, forming a \_\_\_\_\_
- One DNA molecule includes many genes
- The nitrogenous bases in DNA form hydrogen bonds in a complementary fashion:

\_\_\_\_\_ , \_\_\_\_\_

## DNA & Proteins as Tape Measures of Evolution

- genes and their products (proteins) document the hereditary background of an organism
- linear sequences of DNA are passed \_\_\_\_\_; 2 siblings have greater similarity in their DNA than do unrelated individuals...
- it follows, that **2 closely related species** would \_\_\_\_\_  
& protein sequences than 2 distantly related species would...
- that is the case!!!
- example: **the  $\beta$  chain of human hemoglobin**:
- this chain contains **146 amino acids**
  - humans & gorillas \_\_\_\_\_
  - humans & frogs \_\_\_\_\_
- ***Molecular biology has added a new “tape measure” with which we can study evolutionary relationships!!***