



BASICS OF BIOLOGICAL CHEMISTRY

Assignment

January 2015 Finals

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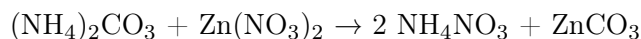
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1 Questions from Prof. J. Vanderleyden – dr. H. Steenackers

1.1 Chemical reaction equation

Consider the following reaction: $(\text{NH}_4)_2\text{CO}_3 + \text{Zn}(\text{NO}_3)_2 \rightarrow \text{NH}_4\text{NO}_3 + \text{ZnCO}_3$

a.) Balance the equation



b.) Reactants and products

Q: Name all reactants and reaction products.

A:

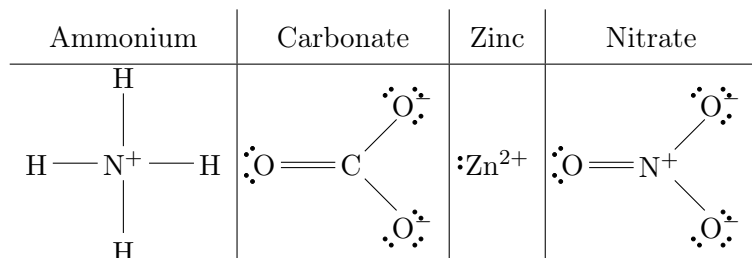
- $(\text{NH}_4)_2\text{CO}_3$: Ammonium carbonate
- $\text{Zn}(\text{NO}_3)_2$: Zinc nitrate
- NH_4NO_3 : Ammonium nitrate
- ZnCO_3 : Zinc carbonate

c.) Lewis structure, VESPR

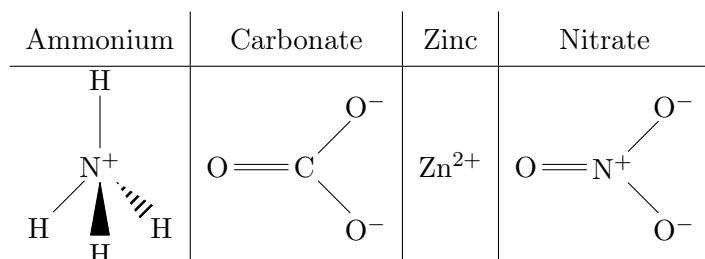
Q: Construct the Lewis structures of the polyatomic ions you recognize and predict their molecular structure using the VSEPR theory.

A:

- Lewis Structure of the ions:



- Molecular structure prediction:



d.) Oxidation states

Q: Determine the oxidation state of all the atoms in all the compounds. Is this an oxidation-reduction reaction?

A: Ammonium Carbonate and Zinc Nitrate (the reactants) are very soluble in water and will thus move freely. The Zinc and Carbonate ions will then precipitate.

- Zn has an oxidation state of 2: $\text{Zn} \rightarrow \text{Zn}^{2+} + 2 \text{e}^-$
- (CO_3) has an oxidation state of 4: $(\text{CO}_3)^{2-} + 2 \text{e}^- \rightarrow \text{CO}_3$

e.) Mass

Q: How many grams of ZnCO_3 can be prepared from 400g $\text{Zn}(\text{NO}_3)_2$ by using sufficient $(\text{NH}_4)_2\text{CO}_3$?

A: Let's start by computing the molecular weight of the 2 reactants:

Molecular weight of $\text{Zn}(\text{NO}_3)_2$ 189.36 g/mol

Molecular weight of $(\text{NH}_4)_2\text{CO}_3$ 96.09 g/mol

Given that there are 400g of $\text{Zn}(\text{NO}_3)_2$, we can calculate the number of moles of reactant (and ignore that of $(\text{NH}_4)_2\text{CO}_3$ since it is in excess):

Moles of $\text{Zn}(\text{NO}_3)_2$ $400 \text{ g} / 189.36 \text{ g/mol} = 2.11237 \text{ moles}$

From this last figure, we can infer that the number of moles of ZnCO_3 will be 2.11237. Given the molecular mass of ZnCO_3 , we can compute the amount of ZnCO_3 produced to be: $2.11237 \text{ mol} * 125.3889 \text{ g/mol} = 264.8678 \text{ g}$.

1.2 DNA sequence analysis

The following diagram shows part of a template DNA strand, with sections X,Y and Z being the exons of a gene:

```
5'                               3'
GTA GGT TGT ATC GAT GGT CAT
---
X       Y       Z
```

a.) DNA Replication

Q: What is the corresponding sequence on the new daughter strand made from the given parent strand during replication?

A: Given the principle of base pairing, we can determine the daughter sequence to be (here in the 3' to 5' direction):

```
5'                               3'
GTA GGT TGT ATC GAT GGT CAT
CAT CCA ACA TAG CTA CCA GTA
3'                               5'
```

b.) Translated Protein

Q: What polypeptide sequence will be synthesized from the given template DNA? Give a short overview of the different processes (and enzymes) involved in the synthesis of polypeptides from template DNA. Where in the cell do these processes take place?

A: The synthesized polypeptide will consist of the amino acids **Val-Cys-Ile-His**. Since it is mentioned that exons are present, we can assume the translation will take place with the eukaryotic machinery. It will consist of the following stages:

- Transcription: In the nucleus, the RNA Polymerase II will be recruited and will bind to the promoter of the gene. It will produce, by moving in the 5' to 3' direction, a pre messenger RNA which will be identical to the DNA template sequence (with the exception that Uracyl will be used instead of Thymine, and also the addition of a 5' CAP). That messenger RNA will then be processed by the spliceosome, which will remove the introns, and a Poly-A tail will also be added at the 3' end of the mRNA. The mRNA is then ready to go outside of the nucleus to be translated.
- Translation: the mRNA leaves the nucleus and passes through the reticulated ER where it will be captured by a ribosome that will either bind to the ER or not. It will then start scanning for a start codon in the mRNA. From that point on, the synthesis of a polypeptide will be accomplished by reading 3 base pairs at a time and pairing these 3 with the correct tRNA. After that, the polypeptide will either be processed further and sent to the golgi apparatus, or will remain in the cytosol.

c.) Mutated exon

Q: What polypeptide sequence will be synthesized if the ATC in exon Y is mutated to TTC? What polypeptide sequence will be synthesized if the ATC in exon Y is mutated to ATG? Which of those substitution mutations is likely to be more harmful? Why?

A: Here are the new sequences with mutated exons:

- TGTATC -> TGT TTC : the resulting polypeptide will be **Val-Phe-His**
- -> ATG : the resulting polypeptide will be **Val-Met-His**

The second

d.) Interactions with antibiotics

Q: Which steps in polypeptide synthesis are affected by resp. the macrolide antibiotics and the tetracycline antibiotics?

A:

e.) Comparison of error rates

Q: The error rate in RNA synthesis is much higher than the error rate of DNA replication. What is the origin of this difference? Motivate why this is not a serious problem.

A:

1.3 tRNA 3D-Structure

Q: All tRNA molecules have a particular 3D-structure. Which functional groups and which chemical bonds/interactions contribute to this particular structure? Why is this particular structure of importance for the biological function?

A:

2 Questions from Prof. B. Sels

2.1 Biopolymer organisation

Q: The course and the textbook systematically organize four important biopolymers mainly according to their chemical structure. Attempt a complete reorganization of the various biopolymer structures (and subfamilies!) according to the following three physiological functions: energy, structure, and communication. Explain the physiological function of each biopolymer type with regard to its chemical structure and/or physical properties.

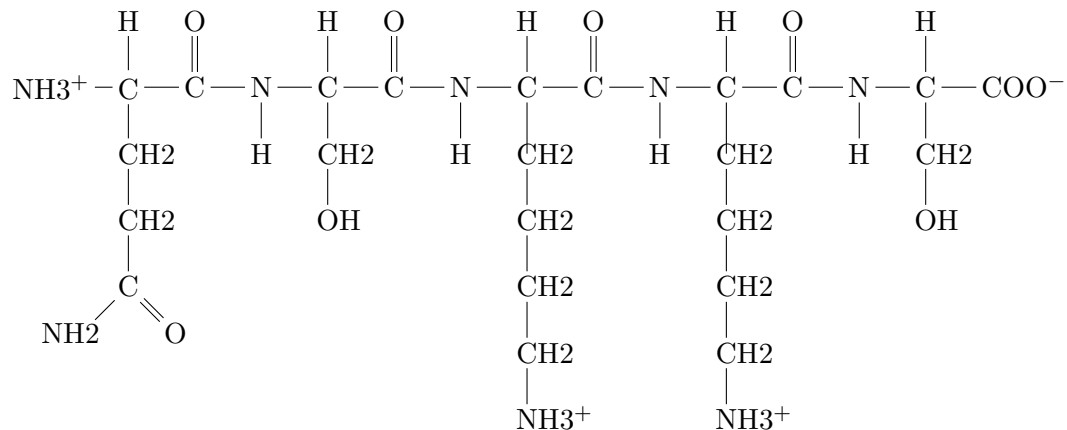
A:

2.2 Chemical structure of proteins and proteins separation

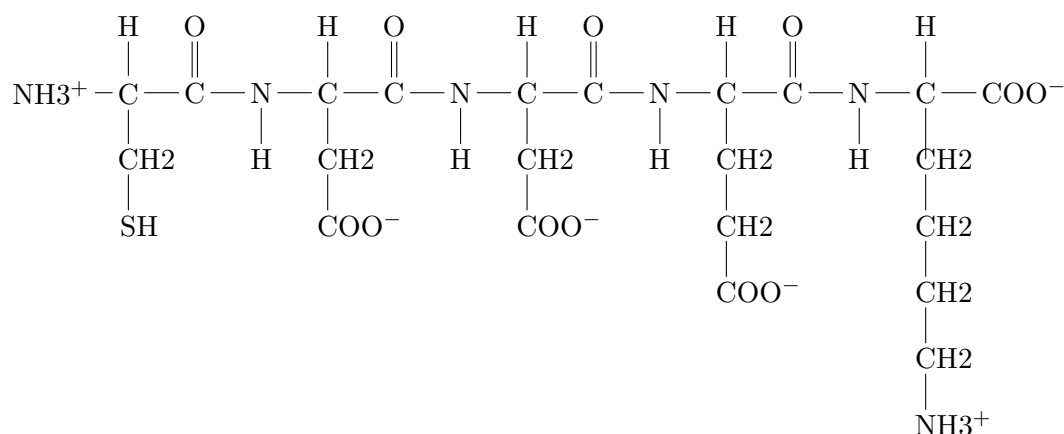
Q: Draw the chemical structure of the following two oligopeptide structures, a) Gln-Ser-Lys-Lys-Ser and b) Cys-Asp-Asp-Glu-Lys, determine its net charge in physiological conditions. How would you separate the two peptides ?

A: These are the chemical structures of:

- Gln-Ser-Lys-Lys-Ser



- Cys-Asp-Asp-Glu-Lys



Under physiological conditions (ie, pH around 7.35), these would be the net charge on each polypeptide:

- Gln-Ser-Lys-Lys-Ser: net charge is +2
 $\text{NH}_3^+ - \text{Gln} - \text{Ser} - \text{Lys}^+ - \text{Lys}^+ - \text{Ser} - \text{COO}^-$
- Cys-Asp-Asp-Glu-Lys: net charge is -2
 $\text{NH}_3^+ - \text{Cys} - \text{Asp}^- - \text{Asp}^- - \text{Glu}^- - \text{Lys}^+ - \text{COO}^-$

Separation of both proteins can thus be achieved by ion exchange chromatography since they both have a distinct charge.

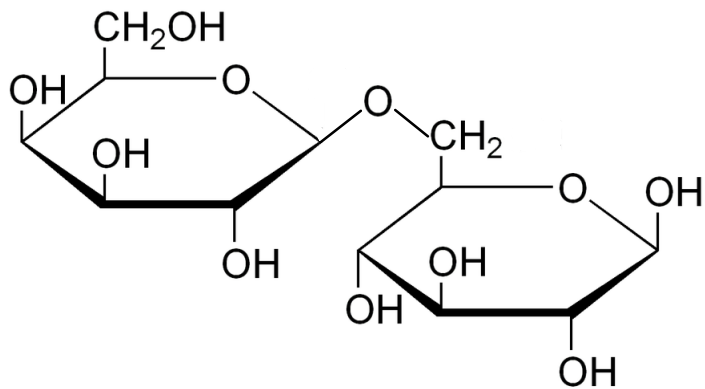
2.3 Chemical structure of disaccharides

Q: Draw the chemical structure of the following disaccharides: a) the β -anomer of $\alpha(1 \rightarrow 6)$ galactoglucose and b) $\beta, \alpha(1 \rightarrow 2)$ glucofructose.

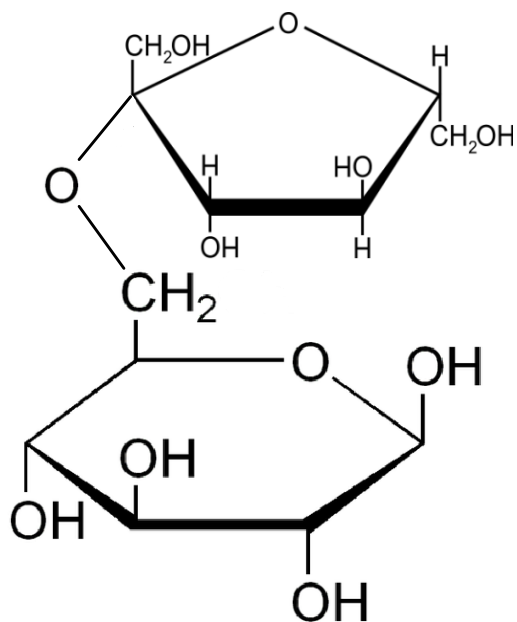
A: These are the chemical structure of:

- β -anomer of $\alpha(1 \rightarrow 6)$ galactoglucose

Beta anomers have a cis relationship between the CH_2OH group on the C_1 and the OH group on the C_6 . This helps us determine the structure of the monosaccharides galactose and glucose. The polymerisation is achieved through an α binding between the C_6 of the Galactose, and the C_1 of the glucose molecule, giving the following molecular structure:

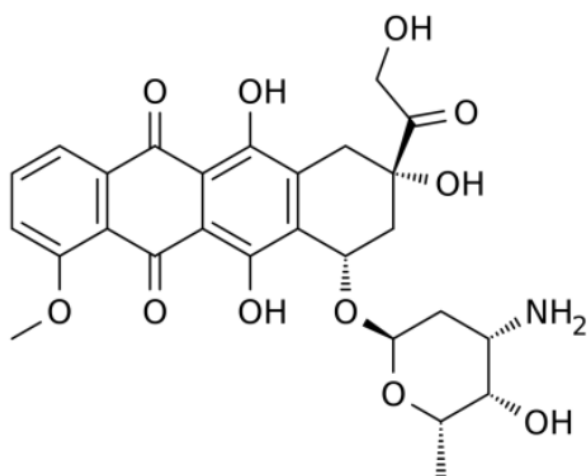


- $\beta, \alpha(1 \rightarrow 2)$ glucofructose



3 Questions from Prof. D. De Vos

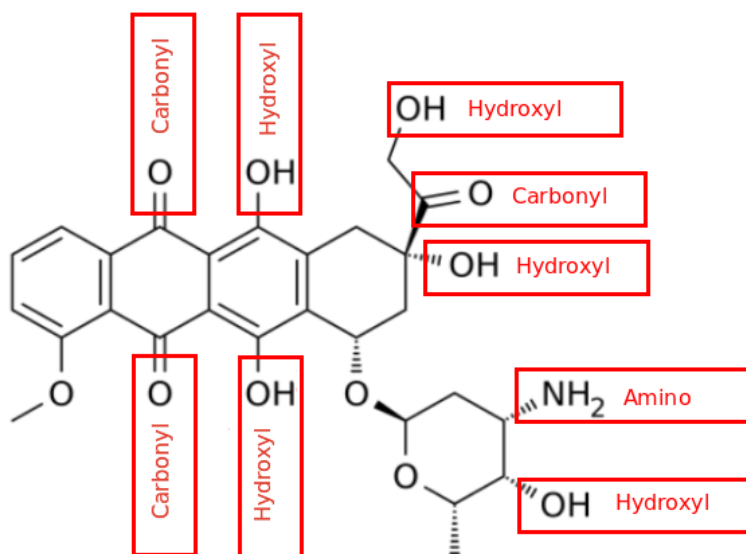
Considering the following molecule:



3.1 Functional groups

Q: Name all functional groups

A: See annotated figure below



3.2 Water and oil solubility factors

Q: Indicate which groups make the molecule rather water-soluble than oil-soluble

A: The following groups can partake in hydrogen bonds with water molecules and increase the solubility of the molecule in water :

- Hydroxyl groups (5 of them)
- Carbonyl groups (3 of them)

- Amino group (1 present)