



Mahindra University Hyderabad  
École Centrale School of Engineering  
End-semester Regular Examination  
(Batch 2022)

UM

Program: B. Tech

Branch: CAM

Year: I

Semester: II

Subject: Chemistry II (Fractal) (CH 1204)

Date: 12/06/2023

Time Duration: 3.0 Hours

Start Time: 10.00 AM

Max. Marks: 100

Instructions

1. Answer all questions.
2. No clarifications will be entertained during the examination.
3. If any information is missing, make appropriate assumptions and proceed.

**PART-A**

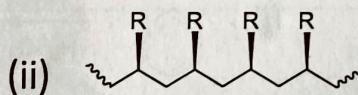
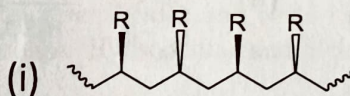
Q1. (Marks: [4+4+2=10])

~~(A)~~ What do you understand by following types of polymers?

(i) homopolymers (ii) copolymers

~~(B)~~ Describe addition (or chain growth) polymerization method?

~~(C)~~ Identify isotactic polymer and syndiotactic polymer from the following structures.

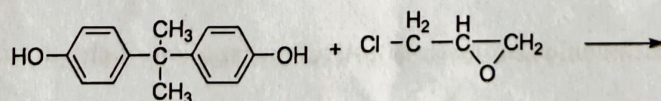


Q2. (Marks: [4+3+3=10])

~~(A)~~ Show the mechanism for (i) initiation (ii) propagation, and (iii) termination steps of free radical polymerization.

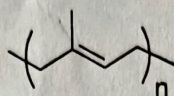
(B) What kind of epoxy resin product is obtained by following polymerization reaction?

?



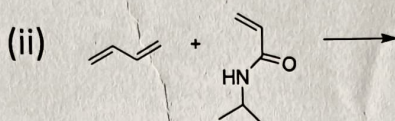
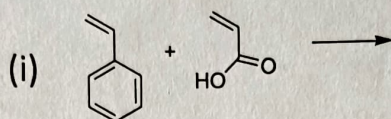


- (C) Show the vulcanization process in rubber by taking an example of following natural polymer  
? poly(isoprene).



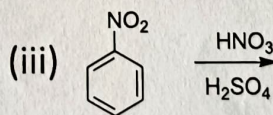
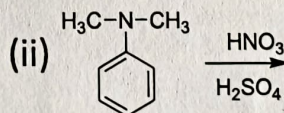
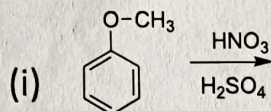
Q3. (Marks: [4+3+3=10])

- (A) Draw the structure of the polymer products in the following reaction.



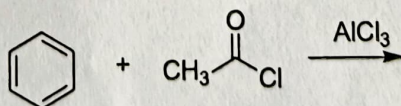
- (B) What is called (i) electrophiles, and (ii) nucleophiles?

- (C) Draw the structures of products for the following nitration reactions.



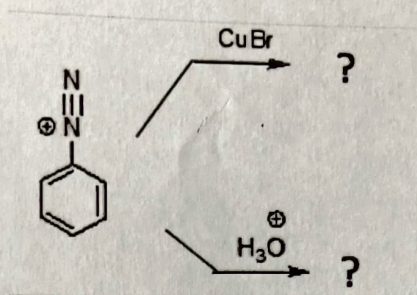
Q4. (Marks: [3+2+5=10])

- (A) Identify the name of following acylation reaction, and find out its product.

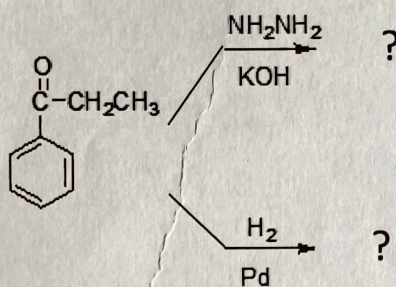


- (B) The following nucleophilic aromatic substitution reaction of the diazonium ions will lead to which product?





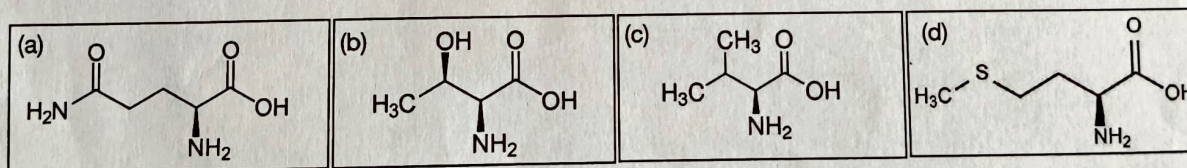
(C) Identify the name of the reaction and the product structure for the following.



## PART-B

Q5. ( Marks: [8+2=10])

- (i) Identify the amino acid residues given below, and write their names using the three-letter code. Classify them into two groups: Hydrophobic and hydrophilic.



- (ii) Give examples of each chemical interactions that stabilize protein structures: (a) Covalent, and (b) Non-covalent

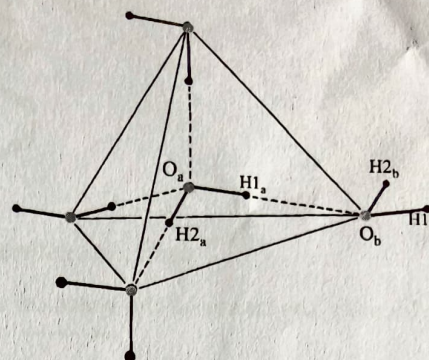
Q6. ( Marks: [5+5=10])

Draw the structures of the following peptides, and label torsion angles Phi ( $\Phi$ ) and Psi ( $\Psi$ ): (a) Gly-Ala, and (b) Ala-Gly



**Q7. ( Marks: [3+2+10+5=20])**

Here a three-dimensional tetrahedral network of water molecules is shown. Oxygen atoms (shown in grey) are situated at the centre and at the vertices of a regular tetrahedron. Answer the following questions:



- Write down the expressions for the two types of nonbonded interaction potentials between two oxygen atoms  $H1_a$  and  $O_b$ .
- Can you imagine how many such van der Waals,  $V_{ab}^{vdW}$  and electrostatic,  $V_{ab}^{ele}$  interaction terms will arise for two interacting water molecules?
- Derive the force ( $z$ -direction) acting on atoms interacting via Lennard-Jones potential.
- Now, calculate the Lennard-Jones force ( $z$ -direction) acting on  $H1_a$  and  $O_b$  atoms separated by  $2.7 \text{ \AA}$ . [Given:  $\epsilon_{OH} = -0.1 \text{ kcal/mol}$ ,  $\sigma_{OH} = 2.7 \text{ \AA}$ ]

**Q8. ( Marks: 10)**

Consider a system of distinguishable particles with five microstates with energies  $0, \epsilon, 2\epsilon, 2\epsilon$  and  $3\epsilon$  in equilibrium with a reservoir at temperature  $T = 3000 \text{ K}$ . Find the partition function of the system. [Given:  $\epsilon = 1.0 \text{ eV}$ , Boltzmann constant ( $k_B$ ) =  $1.381 \times 10^{-23} \text{ J K}^{-1}$ ,  $1 \text{ eV} = 1.602 \times 10^{-19} \text{ J}$ ]

**Q9. ( Marks: [5+5=10])**

- Match the following:

Protein structures	Structural elements
Primary	Loops, turns
Secondary	Backbone
Tertiary	Monomer, dimer
Quaternary	$\alpha$ -helices, $\beta$ -sheets

- Fill in the blanks with appropriate answer.
  - There are \_\_\_\_\_ different naturally occurring amino acids.
  - Properties of amino acids are determined by their specific \_\_\_\_\_.
  - Time-step in a typical MD simulation is \_\_\_\_\_.
  - Velocity Verlet integration algorithm is \_\_\_\_\_.
  - In microcanonical ensemble each microstate exists with \_\_\_\_\_.