

NPTEL

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Courses » Proteins and Gel-Based Proteomics



Announcements

Course

Ask a Question

Progress

Unit 3 - Week-1: Basics of amino acids and proteins



Course outline

How to access the portal

Introduction to Gel-Based Proteomics

Week-1: Basics of amino acids and proteins

- L1. Introduction to amino acids
- L2. Introduction to proteins
- L3. Protein folding & misfolding
- L4. Protein purification techniques
- L5. Introduction to proteomics
- Lab Session1.1:
 Demonstration
 of proteinprotein
 interactions
 using label-free
- Week-1AssignmentAnswer Key

biosensors

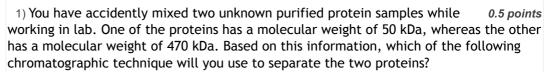
Quiz : Week-1 Assignment-1

Week-2: Gelbased proteomics and sample preparation

Week-3: Twodimensional gel

Week-1 Assignment-1

The due date for submitting this assignment has passed. Due on 2016-03-30, 23:45 IS ... As per our records you have not submitted this assignment.



- Reverse phase chromatography
- High performance liquid chromatography
- Gel permeation chromatography
- Ion-exchange chromatography

No, the answer is incorrect.

Score: 0

Accepted Answers:

Gel permeation chromatography

- 2) Which of the following method(s) can be used to determine the N-terminal 0.5 points sequence of proteins?
 - Sanger method
 - Edman reaction
 - Mass spectrometry
 - All of the above

No, the answer is incorrect.

Score: 0

Accepted Answers:

All of the above

- 3) Which of these amino acids has a good buffering capacity at physiological 0.5 points pH?
 - Histidine
 - Cysteine
 - Tryptophan
 - Proline

No, the answer is incorrect.

Score: 0

Accepted Answers:

Histidine

Proteins and Gel-Based Proteomics - - Unit 3 - Week-1: Basics of amino acids and proteins

4) Which of the following charged group(s) will be present in Valine at a pH of 0.5 points 2? ○ -NH³⁺ -C00⁻ -NH²⁺ a and b No, the answer is incorrect. Score: 0 **Accepted Answers:** -NH³⁺ 5) Which of the following statement is NOT true for protein folding? Once the protein is unfolded it is impossible for it to go back to its native state Protein folding of a protein is dictated by its primary structure No protein adopts only one conformation Protein folding is a cooperative process No, the answer is incorrect. Score: 0 **Accepted Answers:** Once the protein is unfolded it is impossible for it to go back to its native state 6) Which of the following amino acid pairs have 2 chiral centres? 0.5 points Proline, Tyrosine Phenylalanine, Glycine Serine, Cysteine Isoleucine, Threonine No, the answer is incorrect. Score: 0 **Accepted Answers:** Isoleucine, Threonine 7) You have a serum sample mixture from which you want to separate IgG. You 0.5 points know that IgG has very high affinity for Protein A. Which of these chromatography technique would you use to separate IgG from the complex mixture? Affinity chromatography Gel permeation chromatography Cation exchange chromatography Anion exchange chromatography No, the answer is incorrect. Score: 0 **Accepted Answers:** Affinity chromatography 8) Which of the following is a clinical goal of proteomics? 0.5 points Biomarker Discovery Disease monitoring Identification of therapeutic targets All of above No, the answer is incorrect. Score: 0 **Accepted Answers:** All of above 9) Which of the following reagent is used for disruption of disulphide bonds in 0.5 points proteins?

Proteins and Gel-Based Proteomics Unit 3 - Week-1: Basics of amino acids and proteins	
SDS	
O DTT	
O Urea	
O IAA	
No, the answer is incorrect. Score: 0	
Accepted Answers: DTT	f
10)Why is the peptide bond, which is thermodynamically unstable, resistant to 0.5 pospontaneous hydrolysis?	oints Y
Because of partial double character	
 Because of zwitter-ionic nature of amino acids 	Ľ
Presence of side chains	:
None of the above	in
No, the answer is incorrect. Score: 0	8+
Accepted Answers: Because of partial double character	
11)Which of the following amino acid does not possess a chiral carbon centre? 0.5 pc	oints
Histidine	
Cysteine	
Glycine	
Tryptophan	
No, the answer is incorrect. Score: 0	
Accepted Answers: Glycine	
12)You have to perform binding studies of Hemoglobin with different ligands 0.5 pc but the protein solution has impurities like dichromate ions, salts etc. These impurit may interfere with your binding studies and may give you false results. Which of the following chromatographic technique would be the preferred choice of method for protein purification?	ies
Reverse phase chromatography	
High performance liquid chromatography	
Gel permeation chromatography	
 Ion-exchange chromatography 	
No, the answer is incorrect. Score: 0	
Accepted Answers: Gel permeation chromatography	
13)Which of the following amino acids is most likely to be present in the inner 0.5 per core region of proteins?	oints
Lysine	
○ Valine	
Glutamic acid	
Asparagine	
No, the answer is incorrect. Score: 0	
Accepted Answers: Valine	

Proteins and Gel-Based Proteomics Unit 3 - Week-1: Basics of amino acids and protein 14)Which of the following amino acid would you expect in relatively large number in histones? Remember that histones are highly alkaline proteins with pl values.	0.5 points
Lysine Methionine Tyrosine Proline	
No, the answer is incorrect. Score: 0	f
Accepted Answers: Lysine	5
15)Which of these amino acid can undergo phosphorylation in its side chain group?	0.5 poi
ProlineThreonineAlanineMethionine	in g-
No, the answer is incorrect. Score: 0	
Accepted Answers: Threonine	
16)Which of the following statements is TRUE for the two label-free interaction technologies e.g. Surface Plasmon Resonance (SPR) and Bio-layer Interferometry (BLI)? (Note: Question 16-20 are Lab session based questions)	0.5 points
 SPR has fluidic system whereas BLI does not BLI has fluidic system whereas SPR does not Both techniques have fluidic system None of them has fluidic system 	
No, the answer is incorrect. Score: 0	
Accepted Answers: SPR has fluidic system whereas BLI does not	
17]n Bio-layer Interferometry (BLI) when there is an interaction between two molecules, how does the shape of curve changes? It	0.5 points
goes down. rises up. becomes linear. becomes hyperbolic.	
No, the answer is incorrect. Score: 0	
Accepted Answers: rises up.	
18)nteraction studies using label-free platform are advantageous over other techniques because they?	0.5 points
 Overcome the issue of non-specific binding in labels Involve real time approach Provide kinetic information All of the above 	
No, the answer is incorrect. Score: 0	
Accented Answers	

Proteins and Gel-Based Proteomics - - Unit 3 - Week-1: Basics of amino acids and proteins All of the above 19) The basic principle of Bio-layer Interferometry (BLI) technique is based on 0.5 points which of the following phenomenon? Diffraction Polarisation Refraction Interference No, the answer is incorrect. Score: 0 **Accepted Answers:** Interference 20Complex proteomics studies unravelling the physiological pathways are very 0.5 poil crucial if we are mapping protein-protein interaction networks. Which of the following method is NOT used for protein-protein interaction studies? Surface Plasmon Resonance (SPR) Bio-layer Interferometry (BLI) Protein microarray SDS-PAGE No, the answer is incorrect. Score: 0 **Accepted Answers:**

Previous Page

SDS-PAGE

End

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