Mid-semester examination (50 marks; 2 hours)

Question 1. Explain the following (pointed and short answers) (10 marks)
1.1. Minimum labeling vs. Saturation labeling
1.2. IPG strips vs. tube gels
1.3. Forward vs. reverse proteomics
1.4. On-gel electrophoresis vs. Off-gel electrophoresis
1.5. Ion exchange <i>vs.</i> gel exclusion chromatography
1.6. Soft vs. hard ionization
1.7. Equilibration I vs. II
1.8. "b" vs. "y" ions

- **1.9.** Proteomics *vs.* protein chemistry
- **1.10.** Passive rehydration vs. Active rehydration

Question 2. Identify the problem as demonstrated in these 2D gels and describe your suggestions to overcome the problem in each gel image (1.5+1=2.5 X 4=10 marks)

	2D gel image	Problem (s)	Your recommendation (how to overcome the problem)
i			
ii			
iii			
iv			

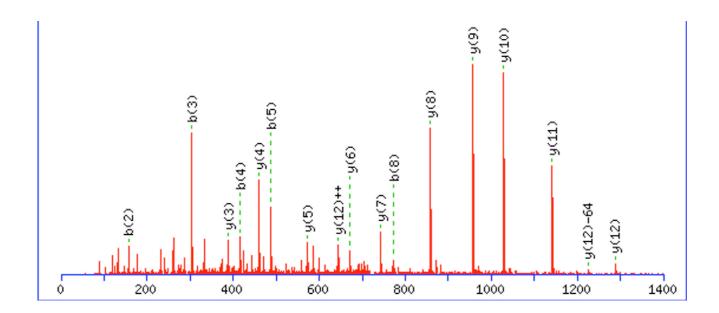
Question 3. $(2 \times 6 = 12 \text{ marks})$

Glutathione peroxidase 3 (Gpx3) is an important antioxidant enzyme in *Saccharomyces cerevisiae*. A study compared the wild type *S. cerevisiae* strain with its Gpx3 mutant strain by using two dimensional gel electrophoresis on 4-7 pI, 24 cm IPG strips. 2D gel images were analyzed using software and 6 spots were selected for further electrospray ionization quadrupole time-of-flight tandem mass spectrometry (ESI-Q-TOF MS/MS) analysis. Raw MS data was processed by MASCOT search program (Version 2.2, Matrix Science) and used for protein identification. Details of proteins identified by MS/MS is listed in table.

Analyse this data in Table and determine what will be your evaluation for data (proteins A-F) on a scale of 1-5 (1 – Best data; 2 – Good data; 3 – OK but acceptable; 4 – Doubtful, there is further need to look at raw data; 5 – Reject). Please fill in your option (only one option for each) in given space. Also, write-down your rationale for choosing which option is correct.

Protei n	Species	pI	Accession No.	Peptid e match	MOWS E score	Fold chang e	Your answer (1-5 scale)
A	Saccharomyces pombe	5.9	gi 6715442	1	13	-3.1	
В	Saccharomyces cerevisiae	6.3	gi 195650293	13	200	+2.3	
С	Saccharomyces cerevisiae	7.0	gi 195639750	7	25	+1.7	
D	Bacillus subtilis	5.5	gi 3036946	1	17	+2.6	
E	Saccharomyces cerevisiae	8.6	gi 13096169	2	39	-2.6	
F	Saccharomyces cerevisiae	5.1	gi 194701222	3	60	-1.9	

Question 4. In MS spectrum provided to you, please write-down the peptide sequence and also show your steps in answer book. You are free to use calculator (8 marks).



Question 5. Please write the answer of following questions ($2X5 = 10 \text{ marks}$)
5.1. In reverse phase chromatography what are mobile and stationary phases?
5.2. What is the extraction solution used for in-gel digestion procedure?
5.3. The processes by which an ion is selected for MS/MS analysis is known as?
5.4. In multidimensional separation, size exclusion chromatography followed by reversed phase is less used due to
5.5. The reflector in TOF analyzer is used for which purpose?