

Shahjalal University of Science & Technology, Sylhet Department of Biochemistry and Molecular Biology

B. Sc. (Hons) 4th Year 1st Semester: Midterm-III

Course No.: BMB -422 Course Title: Genomics and Proteomics

Marks-10, Time-1Hour

1.	Match	the functions	with the	databases	that dea	ls with -
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10x0.3 = 3.0

- i. Gene ii. Bioprojects.iii. GRC įv. dbSNP/v. Genbank yi. OMIM yii. PubMed yiii. Bookself ix. CDD x. dbEST
- a. A collection of genomics, functional genomics and genetics studies and links to their resulting database
- b. A collection of biomedical books
- c. A collection sequence alignment and profiles representing protein domains conserves in evolution
- d. Includes single nucleotide variation, microsatellites and small scale insertion and deletion
- NIH genetic sequence database
- A. Contains short single pass reads of transcrips sequences
- g. Maintains responsibility for the human and mouse references genome
- h. Database of genes focusing on genome that have been completely sequenced
- i. A database of human genes and genetic disorder
- j. A database citations and abstracts for biomedical literature

2. Write down the full name of the acronym-

10x0.2=2.0

a. OMIM b. NCBI iii. EST iv. SNP v. GRC vi. BLAST vii. EMBL viii. STRING IX. PDB X. SIB

3. Write the functions of the following tools-

5x0.3=1.5

- a. Nucleotide blast
- b. Protein blast
- c. blastx
- d. tblastn
- e. tblastx

4. Briefly write down the functions of the following database (any three)

3x0.5=1.5

- a. STRING
- b. I-tasser
- c. PDB
- d. Cytoscape

5. Explain the statements-

2x1.0=2.0

- a. 2-D gel electrophoresis provides benefits over traditional electrophoresis
- b. The application of proteomics is immense and emerging