

- Explain heuristic alignment algorithm in details.
- b) Explain PAM substitution Metrics of sequence alignment in details.

OR

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- **8.** a) Explain Needleman Wunsch algorithm in details.
 - b) Explain BLOSUM substitution metrics of sequence alignments in details.
- 9. a) Explain the primary databases with the help of examples.
 - b) Discuss about searching and retrieval system from www.

OR

- a) Discuss about PDB structural databases.
 - Explain the secondary databases based on
 - i) Swissprot
 - ii) PIR

a)

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10.

12.

b)

- iii) KEGG.
- 11. a) Discuss how biochemical databases are utilized as an expertise for managing the full data 6 life cycle.
 - b) Explain the biochemical databases extension by metabolic surveys.

OR

v.solveout

- Write short note on :
 - i) EXGESCY
 - ii) BRENDA
 - iii) WIT.

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