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1. National Centre for Biotechnology Information was developed by
National Institute of Health (ANS)

2. A Computer-based organization of sequence and structural data of biomolecules is called
Database (ANS)

3. Searching human genome, chromosomes, genes, genomic sequences similar to a given protein sequence, all are activities that can be done in
ENSEMBL (ANS)

4. One of the following is among the purpose of establishing Data Library by European Molecular Biology Laboratory
collecting, organizing and distributing data on nucleotide sequence and other information related to them (ANS)

5.Meta PredictProtein section is a program for predicting protein
secondary structure (ANS)

6. An alignment of a protein sequence to the 3D structure of another protein is called
THREADING (ANS)

7. The statistical method used to access the robustness of phylogenetic trees produced by various tree-building methods is the
BOOTSTRAP (ANS)

8. A text-based search engine for bioinformatics databases, at NCBI is
ENTREZ (ANS)

9. Two sequences can be locally aligned by using
BLAST2SEQUENCES (ANS)

10. Database analysis are of two categories, these are;
Genomic analysis and Proteomics analysis (ANS)

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