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Biological Databases Biological databases can be broadly classified into sequence and structure databases. Sequence databases are applicable to both nucleic acid sequences and protein sequences, whereas structure database is applicable to only Proteins. Databases in general can be classified in to

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Hence, there is a natural demand for diverse and powerful bioinformatic databases for managing this staggering volume of biological data. This chapter introduces the readers to some basic concepts...

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This chapter introduces the readers to some basic concepts about biological databases including their types and designs, as well as information retrieval procedures. This chapter basically focuses on important biological databases like GenBank.

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A biological database is a large, organized body of persistent data, usually associated with computerized software designed to update, query, and retrieve components of the data stored within the system. A simple database might be a single file containing many records, each of which includes the same set of information.

Biological Databases- Types and Importance ...

Biological databases emerged as a response to the huge data generated by low-cost DNA sequencing technologies. One of the first databases to emerge was GenBank, which is a collection of all available protein and DNA sequences. It is maintained by the National Institutes of Health (NIH) and the National Center for Biotechnology Information (NCBI).

Biological Databases: An Overview and Future Perspective ...

Databases introduced include genome and sequence databases such as GenBank and Ensembl, as well as protein databases such as PDB and UniProt. Databases related to RNA, sequence variation, pathways and interactions, metagenomics, and epigenomics are also presented.

605.652 - Biological Databases and Database Tools | Johns ...

Biological databases are libraries of life sciences information, collected from scientific experiments, published literature, high-throughput experiment technology, and computational analysis. They contain information from research areas including genomics, proteomics, metabolomics, microarray gene expression, and phylogenetics.

Biological database - Wikipedia

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