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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

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might be a single file containing many records, each of which includes the same set of information.

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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 ...

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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.

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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,

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metabolomics, microarray gene
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