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The development of databases to handle the vast amount of molecular biological data is thus a fundamental task of bioinformatics. This chapter introduces some basic concepts related to databases, in particular, the types, designs, and architectures of biological databases.

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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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Why biological databases ? • Exponential growth in biological data. • Data (genomic sequences, 3D structures, 2D gel analysis, MS analysis, Microarrays...) are no longer published in a conventional manner, but directly submitted to databases. • Essential tools for biological research.

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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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Introduction to Databases: From Biological to Spatio-Temporal Peter Revesz. Developments in modern computers and applications have resulted in a shift away from relational databases into a chaotic proliferation of specialized database systems. As many of these advanced databases are unable to communicate with each other, database ...

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Limitations of biological databases Though crucial for data organization and storage, the challenges affiliated with biological databases are manifold. Firstly, integration of database content is complicated by inconsistencies at the ontological level; non-standardized nomenclature thus requires a workaround, necessitating map-pings.

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Primary databases (also known as data repositories) are highly organised, user-friendly gateways to the huge amount of biological data produced by researchers around the world. The primary databases were first developed for the storage of experimentally determined DNA and protein sequences in the 1980s and 90s.

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