

Comparative Genomics Services Section Home

History

The roots of comparative genomics can be traced back to Charles Darwin s theory of evolution in the mid-19th century. However, the practical application of comparing genomes gained traction in the late 20th century with advancements in DNA sequencing technologies. The Human Genome Project, completed in 2003, marked a significant milestone, laying the foundation for comparative studies of the human genome with other species.

Evolution Till Date

Comparative genomics has witnessed rapid evolution due to technological advancements. Early comparisons focused on identifying conserved regions among genomes, shedding light on shared genes and regulatory elements. As sequencing costs decreased and computational tools improved, researchers expanded their analyses to include whole-genome comparisons, identifying structural variations, gene families, and functional elements.

Pharmaceuticals

Identifying conserved drug targets across species for drug development. 2.

Biotechnology

Developing genetically modified organisms with desirable traits. 4.

Evolutionary Biology

Studying speciation, adaptation, and evolutionary relationships. 6.

Conservation Biology

Assessing genetic diversity to inform conservation strategies. 8.

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Structural Biology

Understanding protein structures and functions through comparative analyses. 10.

Metagenomics

Analyzing genetic content in environmental samples to study microbial communities. 12.

Bioinformatics

Developing computational tools for genome comparisons and analysis. 14.

Population Genetics

Studying genetic diversity within and among populations. 16.

Immunology

Investigating immune system genes and responses across species. 18.

Parasitology

Understanding host-parasite interactions and coevolution. 20.

Drug Resistance

Studying genetic mechanisms of antibiotic and antiviral resistance. 22.

Cancer Research

Comparing cancer genomes to identify driver mutations and therapeutic targets. 24.

Immunotherapy

Identifying potential immunotherapy targets through genome comparisons.

Single-Cell Genomics

Studying gene expression at the single-cell level across species.

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2.

Epitranscriptomics

Exploring RNA modifications and their evolutionary implications. 4.

Functional Annotation

Uncovering the functions of non-coding elements in the genome. 6.

Comparative Metabolomics

Linking genomic information with metabolic pathways. 8.

Environmental Genomics

Studying how genomes adapt to changing environments. 10.

Synthetic Evolution

Engineering genomes for desired traits through comparative insights. 12.

Phylogenomics

Integrating genomic and phenotypic data to reconstruct evolutionary history. 14.

Coevolution Studies

Analyzing genetic adaptations between interacting species. 16.

Evolutionary Medicine

Applying evolutionary insights to medical research and treatments. 18.

Ethical Considerations

Addressing ethical implications of genomic research and data sharing. 20.