

Gene Prediction Tools Market Size, Share & Trends Analysis Report By Type (Software, Services), By Method (Empirical Methods, Ab Initio Methods), By Application, By End-use, By Region, And Segment Forecasts, 2024 - 2030

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Abstracts

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Gene Prediction Tools Market Size & Trends

The global gene prediction tools market size was estimated at USD 139.8 million in 2023 and is projected t%li%grow at a CAGR of 18.57% from 2024 t%li%2030. This can be attributed t%li%the increasing R&D in biotechnology and pharmaceuticals, technological advancements in genomics, the growing demand for personalized medicine, and the rising prevalence of genetic disorders. Moreover, growth in government initiatives and funding, and the integration of AI and ML are driving the adoption of gene prediction tools. For instance, in December 2022, the UK government announced t%li%invest around USD 190 million in cutting-edge genomics research t%li%establish an advanced genomic healthcare system.

The growing prevalence of genetic disorders and the need for early diagnosis and intervention are driving the demand for gene prediction tools. Accurate gene prediction can facilitate the identification of genetic mutations associated with various diseases, enabling early diagnosis and improved patient outcomes. For instance, genetic disorders such as cystic fibrosis, muscular dystrophy, and certain types of cancer can be better managed with early and precise genetic identification.

The rapid progress in genomic technologies, such as next-generation sequencing



(NGS) and bioinformatics, has revolutionized gene prediction. These advancements have enabled more accurate and efficient identification of genes, driving the adoption of gene prediction tools. For instance, the advent of high-throughput sequencing technologies allows for comprehensive analysis of entire genomes, providing vast amounts of data that can be analyzed using sophisticated gene prediction algorithms. By identifying specific genetic variations, healthcare providers can customize treatments for conditions like cancer, cardiovascular diseases, and rare genetic disorders.

Government initiatives and increased funding for genomic research are als%li%significant drivers of the gene prediction tools market. Governments and research institutions worldwide are investing in genomic projects t%li%advance scientific understanding and improve public health. For instance, the National Institutes of Health (NIH) in the U.S. has funded numerous genomic research initiatives, including the Human Genome Project and the All of Us Research Program. In February 2024, the Australian Government announced the investment of USD 500.1 million in genomic research under the Genomics Health Futures Mission t%li%enhance disease testing and diagnosis, provide personalized treatment options, and improve health outcomes while reducing unnecessary interventions and health costs. Such initiatives provide substantial support for the development and adoption of gene prediction tools, fostering market growth.

Global Gene Prediction Tools Market Report Segmentation

This report forecasts revenue growth at global, regional, and country levels and provides an analysis of the latest industry trends in each of the sub-segments from 2018 t%li%2030. For this study, Grand View Research has segmented the global gene prediction tools market report based on type, method, application, end-use, and region.

Type Outlook (Revenue, USD Million, 2018 - 2030)

Software

Services

Method Outlook (Revenue, USD Million, 2018 - 2030)

Empirical Methods



Ab Initi%li%Methods

Others
Application Outlook (Revenue, USD Million, 2018 - 2030)
Drug Discovery & Development
Diagnostics Development
Others
End-use Outlook (Revenue, USD Million, 2018 - 2030)
Academic & Research
Industrial
Regional Outlook (Revenue, USD Million, 2018 - 2030)
North America
U.S.
Canada
Mexico
Europe
UK
Germany
France
Italy
Spain



Denmark
Sweden
Norway
Asia Pacific
Japan
China
India
Australia
South Korea
Thailand
Latin America
Brazil
Argentina
MEA
South Africa
Saudi Arabia
UAE
Kuwait



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