

Long Read Sequencing Market By Application (Research Applications, Clinical Applications), By Technique (Single-Molecule Real-Time (SMRT) Sequencing, Nanopore Sequencing, Synthetic Long Read Sequencing), By Product (Instruments, Consumables, Services), By End User (Academic and Government Research Institutes, Pharmaceutical and Biotech Companies, Hospitals and Clinics, Others): Global Opportunity Analysis and Industry Forecast, 2023-2032

Market Report | 2024-03-01 | 395 pages | Allied Market Research

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Report description:

The long read sequencing market was valued at \$724.43 million in 2022 and is estimated to reach \$5,163.41 million by 2032, exhibiting a CAGR of 21.7% from 2023 to 2032. Long read sequencing is a genomic technique that reads longer stretches of DNA compared to traditional methods. It employs innovative technologies like single-molecule sequencing to generate DNA sequences spanning thousands to millions of base pairs. Platforms such as Pacific Biosciences and Oxford Nanopore Technologies utilize techniques such as nanopore sequencing to achieve extended read lengths. Long read sequencing facilitates comprehensive analysis of complex genomic regions, repetitive sequences, and structural variations. Its applications include de novo genome assembly, resolving complex genomic structures, understanding genetic diseases, and exploring microbial diversity. Overall, long read sequencing revolutionizes genomic research by providing detailed and accurate genomic information.□

The rise in the prevalence of genetic disorders and cancer is indeed a significant driver for the increased demand for long read sequencing technologies. Genetic disorders and cancer are inherently complex diseases influenced by multiple genetic factors, including mutations, structural variations, and genomic rearrangements. Long read sequencing offers the ability to comprehensively analyze the entire genome, enabling researchers and clinicians to identify rare and novel genetic variants associated with these diseases more accurately. For instance, the World Health Organization (WHO) estimated that approximately

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400,000 children develop cancer every year worldwide, emphasizing the urgent need for advanced genomic technologies to better understand the genetic basis of cancer and develop targeted therapies. By providing insights into the underlying genetic mechanisms of disease, long read sequencing plays a crucial role in advancing precision medicine initiatives and improving clinical outcomes for patients with genetic disorders and cancer. □

In addition, with the decreasing cost of sequencing and increasing awareness of the importance of genomics in various domains, there's a growing demand for comprehensive genomic insights. Long read sequencing provides a deeper understanding of genomic structure and function, allowing researchers to uncover complex genetic variations and better understand the genetic basis of diseases, traits, and evolutionary processes. Thus, the rise in genetic and cancer disorders and the wide application of long read sequencing in the disease diagnosis is potentially driving the market growth. □

However, the lack of skilled professionals in the field of long read sequencing is expected to hinder market growth. Operating and interpreting data from long read sequencing platforms require specialized expertise in molecular biology, bioinformatics, and genomics. The shortage of adequately trained personnel may limit the adoption and utilization of long read sequencing technologies, delaying their implementation in research, clinical diagnostics, and industrial applications. □

The long read sequencing market is segmented on the basis of application, technique, product, end user, and region. By application, it is bifurcated into research application and clinical application. By technique, the market is categorized into single-molecule real-time sequencing (SMRT), nanopore sequencing and synthetic long read sequencing. On the basis of product, it is segmented into instruments, consumables, and services. The instrument segment is further classified into system and software. By end user, it is segregated into academic and government research institutes, pharmaceutical and biotechnological companies, hospitals and clinics, and others. Region wise, the market is analyzed across North America (U.S., Canada, and Mexico), Europe (Germany, France, UK, Italy, Spain, and rest of Europe), Asia-Pacific (Japan, China, India, Australia, South Korea, and rest of Asia-Pacific), and LAMEA (Brazil, South Africa, Saudi Arabia, and rest of LAMEA). □

Major key players that operate in the global long read sequencing market are Agilent Technologies, Inc. BaseClear BV, Cantata Bio, FG Technologies, □Grandomics, □MicrobesNG, Novogene co., Ltd., Oxford Nanopore Technologies, Pacific Biosciences Inc., CD Genomics, □Danaher Corporation, BGI Genomics, □St Vincent's Hospital Australia, Takara Holdings Inc., Thermo Fisher Scientific, Inc., Element Biosciences, and Illumina, Inc. Key players operating in the market have adopted product launch, acquisition, partnership and collaboration as their key strategy to expand their product portfolio. □

Key Benefits for Stakeholders

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This report provides a quantitative analysis of the market segments, current trends, estimations, and dynamics of the long read sequencing market analysis from 2022 to 2032 to identify the prevailing long read sequencing market opportunities.

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The market research is offered along with information related to key drivers, restraints, and opportunities.

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Porter's five forces analysis highlights the potency of buyers and suppliers to enable stakeholders make profit-oriented business decisions and strengthen their supplier-buyer network.

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In-depth analysis of the long read sequencing market segmentation assists to determine the prevailing market opportunities.

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Major countries in each region are mapped according to their revenue contribution to the global market.

-

Market player positioning facilitates benchmarking and provides a clear understanding of the present position of the market players.

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The report includes the analysis of the regional as well as global long read sequencing market trends, key players, market segments, application areas, and market growth strategies.

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- Regulatory Guidelines
- Additional company profiles with specific to client's interest
- Additional country or region analysis- market size and forecast
- Expanded list for Company Profiles
- Historic market data
- Key player details (including location, contact details, supplier/vendor network etc. in excel format)

Key Market Segments

By Application

- Research Applications
- Clinical Applications

By Technique

- Nanopore Sequencing
- Synthetic Long Read Sequencing
- Single-Molecule Real-Time (SMRT) Sequencing

By Product

- Instruments
- Type
- System
- Software
- Consumables
- Services

By End User

- Academic and Government Research Institutes
- Pharmaceutical and Biotech Companies
- Hospitals and Clinics
- Others

By Region

- North America
- U.S.
- Canada
- Mexico
- Europe
- Germany
- France

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- UK
- Italy
- Spain
- Rest of Europe
- Asia-Pacific
- Japan
- China
- India
- Australia
- South Korea
- Rest of Asia-Pacific
- LAMEA
- Brazil
- Saudi Arabia
- South Africa
- Rest of LAMEA
- Key Market Players
- Cantata Bio
- Illumina, Inc.
- Danaher Corporation
- Element Biosciences
- FG Technologies
- MicrobesNG
- Oxford Nanopore Technologies plc.
- CD Genomics
- BGI Genomics
- St Vincents Hospital Melbourne
- Agilent Technologies, Inc.
- TAKARA HOLDINGS INC.
- BaseClear B.V.
- GrandOmics
- Novogene co., ltd.
- PacBio
- Thermo Fisher Scientific Inc.[]

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