

# Global Next-Generation Sequencing (NGS) Market Report and Forecast to 2021

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#### **Abstracts**

Next-Generation Sequencing (NGS) Report by Material, Application, and Geography – Global Forecast to 2021 is a professional and comprehensive research report on the world's major regional market conditions, focusing on the main regions (North America, Europe and Asia-Pacific) and the main countries (United States, Germany, United Kingdom, Japan, South Korea and China).

In this report, the global Next-Generation Sequencing (NGS) market is valued at USD XX million in 2017 and is projected to reach USD XX million by the end of 2021, growing at a CAGR of XX% during the period 2017 to 2021.

The report firstly introduced the Next-Generation Sequencing (NGS) basics: definitions, classifications, applications and market overview; product specifications; manufacturing processes; cost structures, raw materials and so on. Then it analyzed the world's main region market conditions, including the product price, profit, capacity, production, supply, demand and market growth rate and forecast etc. In the end, the report introduced new project SWOT analysis, investment feasibility analysis, and investment return analysis.

The major players profiled in this report include:
Beijing Genomics Institute
CapitalBio Corporation
Berry Genomics Co. Ltd
Illumina
Thermo Fisher Scientific
Roche
Qiagen



The end users/applications and product categories analysis:

On the basis of product, this report displays the sales volume, revenue (Million USD), product price, market share and growth rate of each type, primarily split into-Targeted Sequencing

Exome Sequencing

Transcriptome Sequencing

On the basis on the end users/applications, this report focuses on the status and outlook for major applications/end users, sales volume, market share and growth rate of Next-Generation Sequencing (NGS) for each application, including-Genome Sequencing
Genome Resequencing
DNA-protein Interactions (ChIP-sequencing)



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Targeted Sequencing

Exome Sequencing

Transcriptome Sequencing

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