



Clinical Exome Solutions for Rare Diseases

Whole Exome Sequencing (WES) is the most effective tool for diagnosing complicated and multi-system rare diseases. WES searches through all coding regions of currently identified genes, yielding a high chance of correctly diagnosing patients with a complex phenotype. Novogene offers comprehensive and cost-effective clinical exome solutions that meet the needs of patients and your business.

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Why Partner With Us?



High-quality WES services, bioinformatic analysis, and interpretation.



Fast turnaround time at a highly competitive price.



Customized data file types and reports.

Who Benefits?



Hospitals

Transform patient care by providing affordable genetic diagnostic services to your patients.



Diagnostic Laboratories

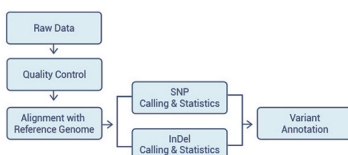
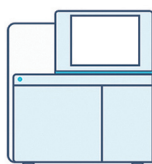
Implement highly complex genetic testing without expensive equipment purchases, validations, or any other associated costs.



Bioinformatics Services

Be a one-stop shop and expand your product offerings with our wet-lab capabilities, providing sequencing data quality you can trust.

Explore Clinical Exome Solutions at Novogene



Whole Exome Sequencing

DNA Extraction
Library Prep
Sequencing

- Performed at Novogene CAP-accredited or CLIA-certified laboratories.
- Data quality with high on-target reads percentage and coverage uniformity.
- Fast Turnaround Time.

Bioinformatic Platform

Data Quality Control
Alignment
Variant Calling & Annotation

- Knowledge-driven NGS analysis, interpretation, and reporting platform based on the GeneCards Suite Knowledge base from >120 data sources and Novogene's in-house database.
- Rapid diagnosis of causal mutations, and prioritization of variants based on their association with patients' phenotypes.
- Evaluation of variants with respect to pathogenicity and causality, which are categorized according to ACMG guidelines.

Interpretation

Variant Evaluation
& Assertion
Test Report & Sign-off

- Report includes primary findings related to patients' phenotypes, medical actionable secondary findings, as well as carrier status for autosomal recessive disorders.
- Clinical interpretation is provided following international best-practice guidelines.

Service Features



Sample Types:

Blood, Saliva, Buccal Swab, or DNA



Exome Capture:

Agilent SureSelect Human All Exon V6



Sequencing Platform:

Illumina NovaSeq 6000



Turnaround Time:

10 Days (from sample receipt to FASTQ)



CLIA-validated WES Performance Metrics

Mean Sequencing Coverage 103X

Average Raw Data 13.75G

Base Pairs Covered at $\geq 20x$ $\geq 94\%$

Coverage Uniformity $\geq 90\%$

Repeatability $\geq 98\%$ for SNVs
 $\geq 88\%$ for Indels

Sensitivity $\geq 98\%$ for SNVs
 $\geq 90\%$ for Indels

Specificity $\geq 99\%$ for SNVs
 $\geq 98\%$ for Indels

Mapping Rate $\geq 97\%$

Custom Deliverables

File / Report Types	Sequencing Only	Sequencing + Analysis	Sequencing + Analysis + Interpretation
FASTQ Files	✓	✓	✓
Data QC Report	✓	✓	✓
BAM Files	✗	✓	✓
VCF Files	✗	✓	✓
Medical Report	✗	✗	✓



Data Releasing Through AWS



Free Data Uploading Service



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