Nevogene

Whole Exome Sequencing

Novogene Whole Exome Sequencing (WES) is a sequencing method that employs high-throughput sequencing of exon regions of more than 20,000 genes per individual, that are enriched through sequence capture technology. WES targets all protein-coding regions (~1% of the whole genome) responsible for 85% of known disease-causing variants. It provides molecular diagnosis of genetic diseases and enables the exploration of novel mutations and new pathogenic mechanisms.



Sensitivity $\geq 92\%$ for SNVs $\geq 92\%$ for InDels

Test Performance

Mean Sequencing Coverage

 $>100 \times$

Average Raw Data

≥12 Gb

Base Pairs Covered at ≥20×

>93%

Repeatability

 $\geq 95\%$ for SNVs $\geq 95\%$ for InDels

≥99%

Mapping Rate



Specificity $\geq 99\%$ for SNVs $\geq 99\%$ for InDels

2 weeks from sample receival to report.

Whole blood in an EDTA purple-top, Buccal Swab, Saliva, or gDNA.

ISO17025.

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