

Highlights

- Industry leader in Whole Exome Sequencing produced >200,000 Exomes to Date
- Optimized products drive variant discovery in focused disease areas

Leveraging vast experience in the production and analysis of human whole exome sequence data, our research offerings represent the cumulative output of the Broad Institute’s knowledge, maximizing specific utility for variant discovery in specific disease areas. Careful analysis of sample inputs, library construction methods, and coverage deliverables has resulted in a set of offerings to provide optimal data to drive scientific discovery.

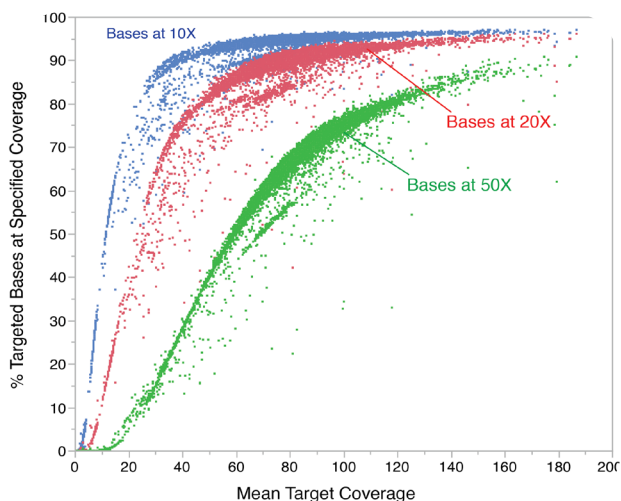
The Broad Institute Genomic Service’s research Whole Exome Sequencing products leverage our experience in producing >200,000 human exomes to date. Using a content bait set containing 38Mb target territory, this content includes all of our previous exome content plus additional coding content that brings the total coverage of the RefSeq and GENCODE v12 databases to >98%. Our coverage deliverables look closer than mean target coverage, providing valuable insight into the actual percentage of targets achieving a minimum guaranteed coverage.

What’s Included

- Sample Receipt, QC & Plating & Sample Preparation
- Illumina HiSeq 2500 or 4000 Sequencing
- Sample Fidelity QC (96 SNP fingerprinting)

Data Delivery and Access

- De-Multiplexed, aggregated Picard BAM file with contamination-checked, and summary metrics
- Data accessed via secure online digital transfer
- Germline Variant Call Format (VCF) and Somatic Mutation Variant Format (MAF) files are also available



Distribution of coverage levels for targeted bases for representative samples sequenced to ~10X, ~20X, and ~50X mean target coverage

Product Details

Product Name	Utility	Input DNA Amount	Compatible Extraction Products	Library Construction	Read Length	Coverage Deliverable	Equivalent MTC
Standard Coverage Germline Exome	non-cancer, including population genomics, rare and common disease cohort studies	100ng	Blood, *saliva, slides, cell pellets or buffy-coat	Transposase Base	2 x 150 bp	85% targeted bases at >20X	~60-80X
Standard Coverage Human WES *Available in Express (28 Day TAT)	control for case-control tumor/normal & trio/somatic analysis	100ng	Fresh frozen tissue, blood, *saliva, cell pellets, or buffy coat	Ligation Based	2 x 76bp	80% targeted bases at >20X	~50-80X
Deep Coverage Human WES *Available in Express (28 Day TAT)	case/proband for case-control tumor/normal & trio/somatic analysis	100ng	Fresh frozen tissue, blood, *saliva, cell pellets, or buffy coat	Ligation Based	2 x 76bp	85% targeted bases >50X	~120-150X

*refer to our application notes for special considerations when working with these material types, www.genomics.broadinstitute.org/products/nucleic-acid-extractions