Clinical Exome Sequencing, Proband Only (with Mitochondrial Genome)

Test Code: EXMTO
Turnaround time: 6 weeks
CPT Codes: 81415 x1, 81460 x1

Condition Description

What is the Medical EmExome?

The human exome is the complete coding (exonic) region of the genome. It is estimated to encompass approximately 1-2% of the genome, yet contains approximately 85% of disease-causing pathogenic variants. The Medical EmExome is expertly curated to target genes known or suspected to cause disease. The design provides >96% coverage of 19,000 genes, with a mean read depth of 100X. For the ~5,400 disease-associated genes analyzed, we typically get coverage >98%.

This testing includes sequencing of the mitochondrial genome.

Will a particular gene be covered on the EmExome?

The Exome Coverage Tool can be used to view typical depth of sequence coverage obtained by exome sequencing performed by EGL. To access this tool, please click here. If a gene of specific interest does not have consistent (100%) coverage, please ask us about Sanger fill-in.

Will EGL release raw exome data?

Yes, upon request.

Will EGL re-analyze data?

Yes, upon request.

What options are available for Medical EmExome testing?

EXOMT - Medical EmExome: Clinical Exome Sequencing, Trios
EXOME - Medical EmExome: Clinical Exome Sequencing, Proband Only
EXOMA - Medical EmExome: Clinical Exome Sequencing, Additional Family Member (EXOMT should be order first or at the same time)
EXMTT – Clinical Exome Sequencing, Family Trios (with Mitochondria Genome)
EXMTO – Clinical Exome Sequencing, Proband Only (with Mitochondria Genome)
EXMX3 – Expedited Medical EmExome: Clinical Exome Sequencing, Trios
EXMX5 – Expedited Medical EmExome: Clinical Exome Sequencing, Proband Only
EXO3DD - Medical EmExome Array: Deletion/Duplication Analysis
EXINT - Interpretation Only (Exome or Genome)
EXOMR - Medical EmExome: Clinical Exome Reanalysis
EX3UP – Medical EmExome: Trios Upgrade (Reanalysis of a proband-only exome as a trio; call for details)

Indications

This test is indicated for individuals with a complex or ambiguous phenotype or for individuals with clinical features of a genetic disorder for whom previous testing has been non-diagnostic.

Methodology

Medical EmExome is performed on genomic DNA using in solution hybridization to enrich for the exome. These targeted regions are then sequenced using next-generation sequencing technology at an average coverage of 100X in the target regions. This sequencing typically provides >96% coverage of the ~19,100 genes in the exome at >20X. Intronic variants within 10 nucleotides from the exon/intron boundaries are analyzed, unless prohibited by the complexity of the sequence. The DNA sequence is mapped to and analyzed in comparison with the published human genome build UCSC hg19 reference sequence. The targeted coding exons and splice junctions of genes associated with disease are assessed for the depth of coverage and data quality threshold values. EGL has developed an EmExome bioinformatics analysis pipeline to compare sequence changes in the individual being tested to the reference sequence. High-quality variants which pass EGL’s quality filters are not confirmed by Sanger sequencing.

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Reportable variants that do not pass the quality filters are confirmed using bidirectional Sanger sequence analysis.

The genes encoded by the mitochondrial genome are PCR amplified using 46 pairs of overlapping primers. The PCR products are sequenced in the forward and reverse directions. Sequences are compared to revised Cambridge reference sequence (rCRS). Low levels of heteroplasmy may not be detected.

### Detection

Based on published studies, WES is expected to provide a diagnosis in 20-30% of the cases for rare and ultra-rare disorders.

### Specimen Requirements

Submit only 1 of the following specimen types

* Preferred specimen type: Whole Blood

**Type: Whole Blood**

Specimen Requirements:

In EDTA (purple top) or ACD (yellow top) tube: 5-10 ml for all ages

Specimen Collection and Shipping: Refrigerate until time of shipment. Ship sample within 5 days of collection at room temperature with overnight delivery.

Specimen Collection and Shipping: Do not expose specimen to heat or direct sunlight. Keep the specimen dry. Ship sample at room temperature with overnight delivery.

**Type: Saliva**

Specimen Requirements:

NGSwab is accepted for this testing. NGSwab utilizes Oragene's OCD-100 kit to collect high quality DNA from saliva. For more information, click [here](#)

Specimen Collection and Shipping: Do not expose specimen to heat or direct sunlight. Keep the specimen dry. Ship sample at room temperature with overnight delivery.

**Special Instructions**

Please submit medical records or clinic summary notes, and a signed consent form when ordering exome testing. Failure to receive clinic notes or a signed consent form may result in a delay in testing.