{#patient}

Whole Exome Sequencing Report

Patient Information

- **Name:**
- **Sex/Age:**
- **Date Received:**
- **Indication:**

Sequencing Platform

Whole Exome Sequencing on the Illumina Novaseq 6000 NGS Platform

Clinical Indication:

Likely Pathogenic Variant Detected Related to the Clinical Phenotype

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Key Findings

genome

Gene&Transcript	Location	Variant	Zygosity	Inhertance	ClinicalSignificance
{gene.geneAndTranscript}					{d.clinicalSignificance

Genetic test results are reported based on the recommendations of American College of Medical Genetics

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Variant Interpretation & Clinical Correlation

****:

The submitted sample showsvariant inof gene().

Variant Characteristics:

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Variant Evidence

Gene Impact

- **Gene:**

- **Effect:**

genoox

- **Exon:**
- **Transcript:**
- **Protein:**
- **Coding:**

Based on above evidence, this variant (::) is classified as.

OMIM Phenotype:

Recommendations

1.
2.
3.

Methodology

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Genoox

Limitations/Disclaimer

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