

Genetic Report for Sample HG00512

Sample HG00512 likely has Disease Z due to the identified structural variant at the genomic coordinate chr1:90082. The identified structural variant in the genome of this individual is pathogenic based on previous research. More information about this genetic variant can be found in the ClinVar database using the ClinVar reference ID 91011.

Genetic Report for Sample HG02055

Sample HG02055 likely has Disease X due to the identified structural variant at the genomic coordinate chr1:10746. The identified structural variant in the genome of this individual is pathogenic based on previous research. More information about this genetic variant can be found in the ClinVar database using the ClinVar reference ID 1234.

Genetic Report for Sample HG03540

Sample HG03540 likely has Disease Y due to the identified structural variant at the genomic coordinate chr1:10626. The identified structural variant in the genome of this individual is likely pathogenic based on previous research. More information about this genetic variant can be found in the ClinVar database using the ClinVar reference ID 5678.

Genetic Report for Sample PGP1

Sample PGP1 likely has Disease Z due to the identified structural variant at the genomic coordinate chr1:90082. The identified structural variant in the genome of this individual is pathogenic based on previous research. More information about this genetic variant can be found in the ClinVar database using the ClinVar reference ID 91011.