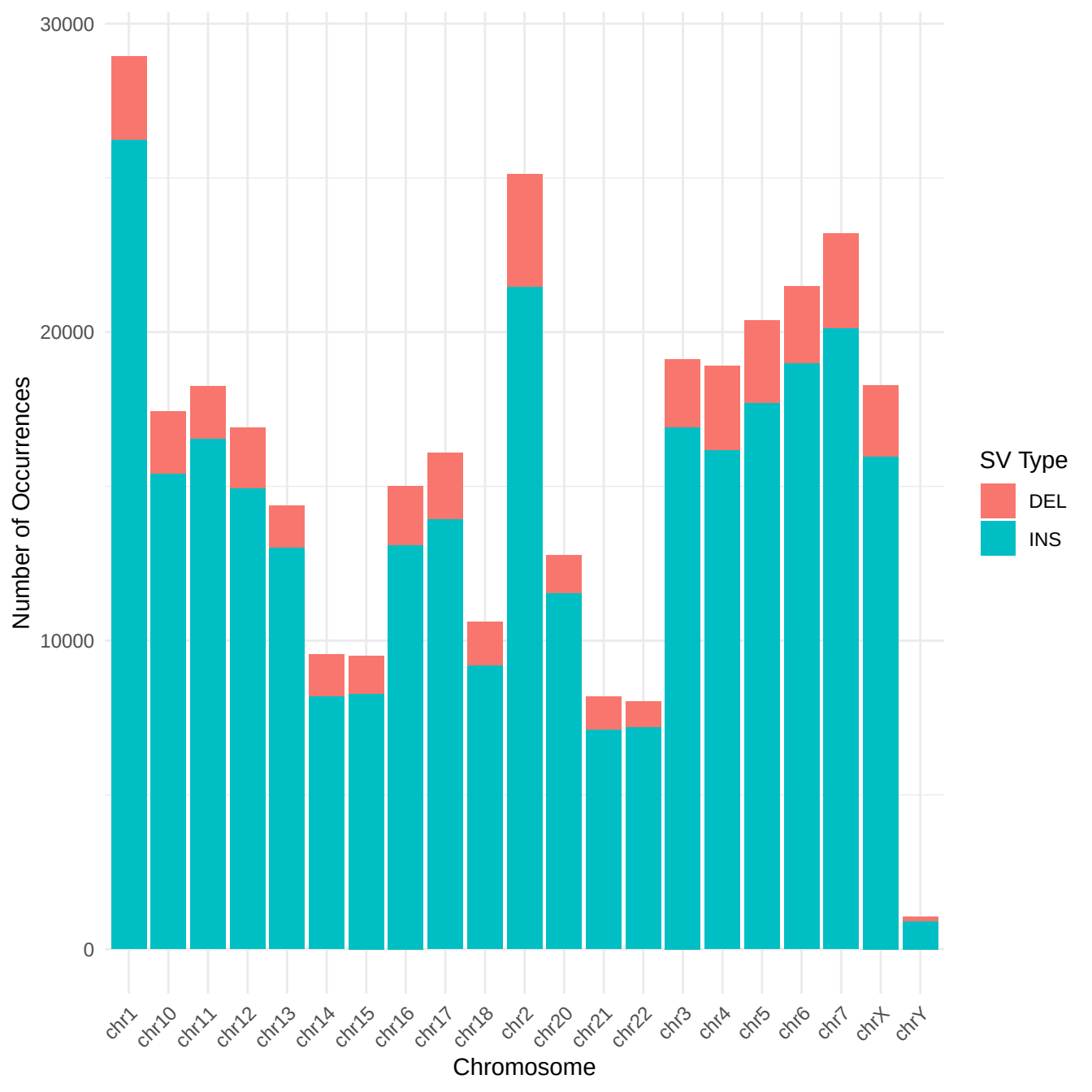
30000
20000
10000
0
Number of Occurrences
chr1
chr10
chr11
chr12
chr13
chr14
chr15
chr16
chr17
chr18
chr2
chr20
chr21
chr22
chr3
chr4
chr5
chr6
chr7
chrX
chrY
Chromosome
SV Type
DEL
INS

Genetic Report for Sample HG00732

Sample HG00732 is at risk for Hepatocellular carcinoma|Small cell lung carcinoma|Lung cancer due to an identified structural variant at the genomic coordinate chr3:195786528. 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 Human Phenotype Ontology:HP:0001402,Human Phenotype Ontology:HP:0002899,Human Phenotype Ontology:HP:0003007,Human Phenotype Ontology:HP:0006750,MONDO:MONDO:0007256,MedGen:C2239176,OMIM:114550,Or CT:187769009,SNOMED CT:25370001|Human Phenotype Ontology:HP:0030357,MONDO:MONDO:0008433,MeSH:D055752,MedGen:C0149925,C CT:363358000.

Genetic Report for Sample HG00733

Sample HG00733 is at risk for Hepatocellular carcinoma|Small cell lung carcinoma|Lung cancer due to an identified structural variant at the genomic coordinate chr3:195786528. 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 Human Phenotype Ontology:HP:0001402,Human Phenotype Ontology:HP:0002899,Human Phenotype Ontology:HP:0003007,Human Phenotype Ontology:HP:0006750,MONDO:MONDO:0007256,MedGen:C2239176,OMIM:114550,Or CT:187769009,SNOMED CT:25370001|Human Phenotype Ontology:HP:0030357,MONDO:MONDO:0008433,MeSH:D055752,MedGen:C0149925,C CT:363358000.

Genetic Report for Sample HG01175

Sample HG01175 is at risk for Hepatocellular carcinoma|Lung cancer due to an identified structural variant at the genomic coordinate chr3:195786528. 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 Human Phenotype Ontology:HP:0001402,Human Phenotype Ontology:HP:0002899,Human Phenotype Ontology:HP:0003007,Human Phenotype Ontology:HP:0006750,MONDO:MONDO:0007256,MedGen:C2239176,OMIM:114550,OrCT:187769009,SNOMED CT:25370001|MONDO:MONDO:0008903,MedGen:C0242379,OMIM:211980,SNOMED CT:363358000.

Genetic Report for Sample HG02145

Sample HG02145 is at risk for Hepatocellular carcinoma|Small cell lung carcinoma|Lung cancer due to an identified structural variant at the genomic coordinate chr3:195786528. 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 Human Phenotype Ontology:HP:0001402,Human Phenotype Ontology:HP:0002899,Human Phenotype Ontology:HP:0003007,Human Phenotype Ontology:HP:0006750,MONDO:MONDO:0007256,MedGen:C2239176,OMIM:114550,Or CT:187769009,SNOMED CT:25370001|Human Phenotype Ontology:HP:0030357,MONDO:MONDO:0008433,MeSH:D055752,MedGen:C0149925,C CT:363358000.

Genetic Report for Sample HG03540

Sample HG03540 is at risk for Hepatocellular carcinoma|Lung cancer due to an identified structural variant at the genomic coordinate chr3:195786528. 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 Human Phenotype Ontology:HP:0001402,Human Phenotype Ontology:HP:0002899,Human Phenotype Ontology:HP:0003007,Human Phenotype Ontology:HP:0006750,MONDO:MONDO:0007256,MedGen:C2239176,OMIM:114550,Or CT:187769009,SNOMED CT:25370001|MONDO:MONDO:0008903,MedGen:C0242379,OMIM:211980,SNOMED CT:363358000.

Genetic Report for Sample NA21309

Sample NA21309 is at risk for Lung cancer due to an identified structural variant at the genomic coordinate chr3:195782436. 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 MONDO:MONDO:0008903,MedGen:C0242379,OMIM:211980,SNOMED CT:363358000

Genetic Report for Sample ei:HG00733

Sample ei:HG00733 is at risk for Hepatocellular carcinoma|Small cell lung carcinoma|Lung cancer due to an identified structural variant at the genomic coordinate chr3:195786528. 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 Human Phenotype Ontology:HP:0001402,Human Phenotype Ontology:HP:0002899,Human Phenotype Ontology:HP:0003007,Human Phenotype Ontology:HP:0006750,MONDO:MONDO:0007256,MedGen:C2239176,OMIM:114550,Or CT:187769009,SNOMED CT:25370001|Human Phenotype Ontology:HP:0030357,MONDO:MONDO:0008433,MeSH:D055752,MedGen:C0149925,C CT:363358000.

Genetic Report for Sample li:HG00733

Sample li:HG00733 is at risk for Hepatocellular carcinoma|Small cell lung carcinoma|Lung cancer due to an identified structural variant at the genomic coordinate chr3:195786528. 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 Human Phenotype Ontology:HP:0001402,Human Phenotype Ontology:HP:0002899,Human Phenotype Ontology:HP:0003007,Human Phenotype Ontology:HP:0006750,MONDO:MONDO:0007256,MedGen:C2239176,OMIM:114550,Or CT:187769009,SNOMED CT:25370001|Human Phenotype Ontology:HP:0030357,MONDO:MONDO:0008433,MeSH:D055752,MedGen:C0149925,C CT:363358000.