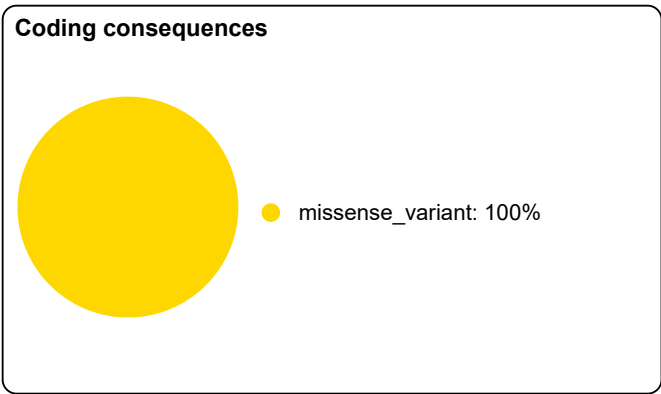
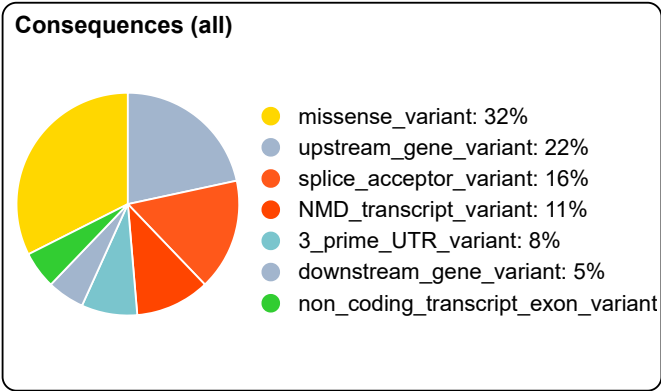


Variant Effect Predictor results

Job details

Summary statistics

Category	Count
Variants processed	2
Variants filtered out	0
Novel / existing variants	0 (0.0) / 2 (100.0)
Overlapped genes	3
Overlapped transcripts	33
Overlapped regulatory features	0



Results preview

Navigation (per variant)

New job

Page: 1 of 1 | Show: 1 All variants

Filters

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Uploaded variant

is

defined

Add

All: VCF VEP TXT

BioMart: Variants Genes

Show/hide columns (28 hidden)

Uploaded variant	Location	Allele	Consequence	Symbol	Gene	Feature type	Featu
.	1:65619927-65619927	T	splice_acceptor_variant	LEPR	ENSG00000116678	Transcript	ENST
.	1:65619927-65619927	T	splice_acceptor_variant	LEPR	ENSG00000116678	Transcript	ENST
.	1:65619927-65619927	T	splice_acceptor_variant	LEPR	ENSG00000116678	Transcript	ENST
.	1:65619927-65619927	T	splice_acceptor_variant	LEPR	ENSG00000116678	Transcript	ENST

Uploaded variant	Location	Allele	Consequence	Symbol	Gene	Feature type	Featu
.	1:65619927-65619927	T	splice acceptor variant	LEPR	ENSG00000116678	Transcript	ENST
.	1:65619927-65619927	T	upstream gene variant	LEPR	ENSG00000116678	Transcript	ENST
.	1:65619927-65619927	T	splice acceptor variant	LEPR	ENSG00000116678	Transcript	ENST
.	X:71121046-71121046	T	missense variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	missense variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	missense variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	downstream gene variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	upstream gene variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	3_prime UTR variant, NMD transcript variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	missense variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	upstream gene variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	missense variant, NMD transcript variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	non coding transcript exon variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	3_prime UTR variant, NMD transcript variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	upstream gene variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	missense variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	missense variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	non coding transcript exon variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	upstream gene variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	upstream gene variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	upstream gene variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	missense variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	missense variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	3_prime UTR variant, NMD transcript variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	missense variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	upstream gene variant	MED12	ENSG00000184634	Transcript	ENST
.	X:71121046-71121046	T	downstream gene variant	-	ENSG00000228427	Transcript	ENST
.	X:71121046-71121046	T	missense variant	MED12	ENSG00000184634	Transcript	ENST

9/1/26, 18:06

Variant Effect Predictor results - Homo_sapiens - Ensembl genome browser 115

Uploaded variant	Location	Allele	Consequence	Symbol	Gene	Feature type	Featu
.	X:71121046-71121046	T	missense_variant	MED12	ENSG00000184634	Transcript	ENST

Page: 1 of 1

 | Show: [1](#) [All](#) variants

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