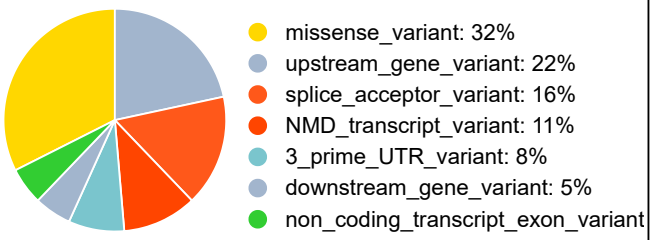
Variant Effect Predictor results

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

Consequences (all)



Coding consequences

Results preview

Navigation (per variant) New job

Page: 1 of 1 | **Show:** 1 All variants

Filters

Uploaded variant | is | defined | Add

Download

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

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

New job