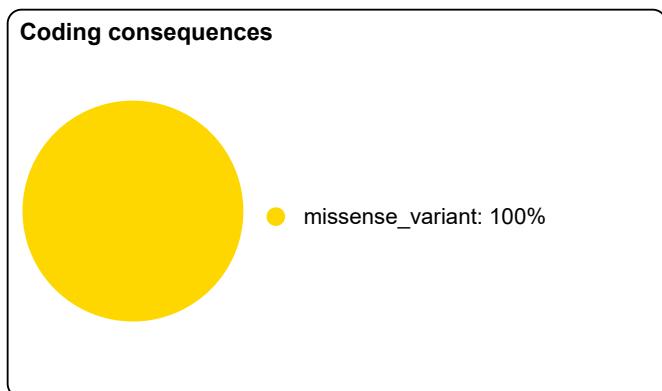
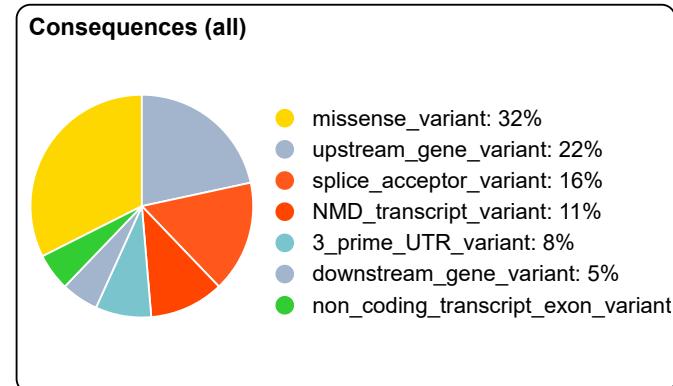


## Variant Effect Predictor results

### 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**

Uploaded variant  is  defined  All: [VCF](#) [VEP](#) [TXT](#)  
BioMart: [Variants](#) [Genes](#)

Show/hide columns (28 hidden)

Uploaded variant	Location	Allele	Consequence	Symbol	Gene	Feature type	Feature
.	1:65619927-	T <a href="#">65619927</a>	splice_acceptor_variant	LEPR	<a href="#">ENSG00000116678</a>	Transcript	<a href="#">ENST</a>
.	1:65619927-	T <a href="#">65619927</a>	splice_acceptor_variant	LEPR	<a href="#">ENSG00000116678</a>	Transcript	<a href="#">ENST</a>
.	1:65619927-	T <a href="#">65619927</a>	splice_acceptor_variant	LEPR	<a href="#">ENSG00000116678</a>	Transcript	<a href="#">ENST</a>
.	1:65619927-	T <a href="#">65619927</a>	splice_acceptor_variant	LEPR	<a href="#">ENSG00000116678</a>	Transcript	<a href="#">ENST</a>

Uploaded variant	Location	Allele	Consequence	Symbol	Gene	Feature type	Feature
	<a href="#">1:65619927-</a>	T <a href="#">65619927</a>	splice_acceptor_variant	LEPR	<a href="#">ENSG00000116678</a>	Transcript	<a href="#">ENST</a>
	<a href="#">1:65619927-</a>	T <a href="#">65619927</a>	upstream_gene_variant	LEPR	<a href="#">ENSG00000116678</a>	Transcript	<a href="#">ENST</a>
	<a href="#">1:65619927-</a>	T <a href="#">65619927</a>	splice_acceptor_variant	LEPR	<a href="#">ENSG00000116678</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	missense_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	missense_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	missense_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	downstream_gene_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	upstream_gene_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	3_prime_UTR_variant, NMD_transcript_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	missense_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	upstream_gene_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	missense_variant, NMD_transcript_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	non_coding_transcript_exon_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	3_prime_UTR_variant, NMD_transcript_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	upstream_gene_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	missense_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	missense_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	non_coding_transcript_exon_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	upstream_gene_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	upstream_gene_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	upstream_gene_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	missense_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	missense_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	3_prime_UTR_variant, NMD_transcript_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	missense_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	upstream_gene_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	downstream_gene_variant	-	<a href="#">ENSG00000228427</a>	Transcript	<a href="#">ENST</a>
	<a href="#">X:71121046-</a>	T <a href="#">71121046</a>	missense_variant	MED12	<a href="#">ENSG00000184634</a>	Transcript	<a href="#">ENST</a>

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

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New job

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