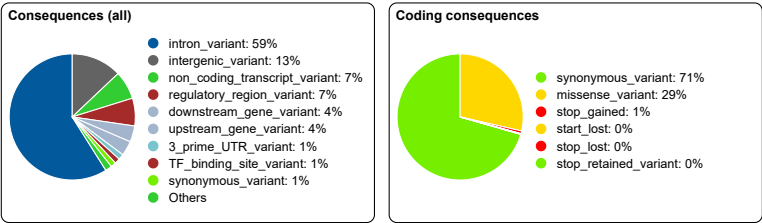


Variant Effect Predictor results

Job details

Summary statistics

Category	Count
Variants processed	164638
Variants filtered out	0
Novel / existing variants	76489 (46.5) / 88149 (53.5)
Overlapped genes	16780
Overlapped transcripts	51898
Overlapped regulatory features	33643



Results preview

Navigation (per variant)

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Filters

Uploaded variant is defined

Download

All: VCF VEP TXT BioMart: Variants Genes

New job

Show/hide columns (38 hidden)

Uploaded variant	Location	Allele	Consequence	Symbol	Gene	Feature type	Feature	Biotype	Existing variant	Feature strand	ENSP	GIVEN REF	USED REF	BAM EDIT	Motif name
rs3131972	1:752721-752721	G	upstream_gene_variant	FAM87B	400728	Transcript	NR_103536.1	lncRNA	rs3131972	1	-	A	A	-	Scroll to see more columns
rs3131972	1:752721-752721	G	regulatory_region_variant	-	-	RegulatoryFeature	ENSR00000918393	promoter	rs3131972	-	-	-	-	-	»
rs3131972	1:752721-752721	G	TF_binding_site_variant	-	-	MotifFeature	ENSM00800836535	-	rs3131972	1	-	-	-	-	»
rs11240777	1:798959-798959	G	downstream_gene_variant	FAM41C	284593	Transcript	NR_027055.1	lncRNA	-	-1	-	A	A	OK	-
rs11240777	1:798959-798959	G	downstream_gene_variant	LINC01128	643837	Transcript	NR_047519.1	lncRNA	-	1	-	A	A	-	-
rs11240777	1:798959-798959	G	downstream_gene_variant	LINC01128	643837	Transcript	NR_047521.1	lncRNA	-	1	-	A	A	-	-
rs11240777	1:798959-798959	G	downstream_gene_variant	LINC01128	643837	Transcript	NR_047523.1	lncRNA	-	1	-	A	A	-	-
rs11240777	1:798959-798959	G	downstream_gene_variant	LINC01128	643837	Transcript	NR_047524.1	lncRNA	-	1	-	A	A	-	-
rs11240777	1:798959-798959	G	downstream_gene_variant	LINC01128	643837	Transcript	NR_047525.1	lncRNA	-	1	-	A	A	-	-
rs13303106	1:891945-891945	G	intron_variant	NOC2L	26155	Transcript	NM_015658.4	protein_coding	rs13303106	-1	NP_056473.3	A	A	-	-
rs13303106	1:891945-891945	G	upstream_gene_variant	KLHL17	339451	Transcript	NM_198317.3	protein_coding	rs13303106	1	NP_938073.1	A	A	-	-
rs13303010	1:894573-894573	G	intron_variant	NOC2L	26155	Transcript	NM_015658.4	protein_coding	-	-1	NP_056473.3	A	A	-	-
rs13303010	1:894573-894573	G	upstream_gene_variant	KLHL17	339451	Transcript	NM_198317.3	protein_coding	-	1	NP_938073.1	A	A	-	-
rs13303010	1:894573-894573	G	regulatory_region_variant	-	-	RegulatoryFeature	ENSR00000000111	promoter	-	-	-	-	-	-	-
rs6696281	1:903104-903104	T	intron_variant	PLEKHN1	84069	Transcript	NM_001160184.2	protein_coding	rs6696281	1	NP_001153656.1	C	C	-	-
rs6696281	1:903104-903104	T	intron_variant	PLEKHN1	84069	Transcript	NM_001367552.1	protein_coding	rs6696281	1	NP_001354481.1	C	C	-	-
rs6696281	1:903104-903104	T	intron_variant	PLEKHN1	84069	Transcript	NM_032129.3	protein_coding	rs6696281	1	NP_115505.2	C	C	-	-
rs6696281	1:903104-903104	T	downstream_gene_variant	KLHL17	339451	Transcript	NM_198317.3	protein_coding	rs6696281	1	NP_938073.1	C	C	-	-

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New job

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