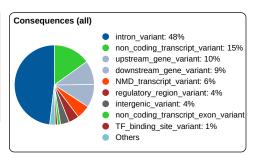
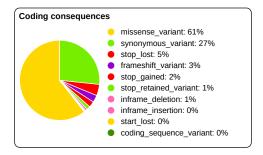
Variant Effect Predictor results

Job details

Summary statistics

Category	Count
Variants processed	29507
Variants filtered out	0
Novel / existing variants	28736 (97.4) / 771 (2.6)
Overlapped genes	3947
Overlapped transcripts	15705
Overlapped regulatory features	4498





Results preview

Navigation (per variant)		Filters			
Page: 1 of 5902	Show: 1 5 10 50 All variants	Uploaded variant	▼ is ▼	defined	Add
		Download			

All: <u>VCF VEP TXT</u>
BioMart: <u>Variants</u> <u>Genes</u>

Show/hide	columns (24	hidden)							Scr see moi
Uploaded variant	Location	Allele	Consequence	Symbol	Gene	Feature type	Feature	Biotype	Exi: » vari
	1:14653- 14653	Т	downstream_gene_variant	DDX11L1	ENSG00000223972	Transcript	ENST00000450305.2	transcribed_unprocessed_pseudogene	<u>rs626</u>
	1:14653- 14653	Т	downstream_gene_variant	DDX11L1	ENSG00000223972	Transcript	ENST00000456328.2	IncRNA	<u>rs626</u>
	1:14653- 14653	Т	intron_variant, non_coding_transcript_variant	WASH7P	ENSG00000227232	Transcript	ENST00000488147.1	unprocessed_pseudogene	<u>rs626</u>
	1:14653- 14653	Т	downstream_gene_variant	MIR6859-1	ENSG00000278267	Transcript	ENST00000619216.1	miRNA	<u>rs626</u>
	<u>1:55299-</u> <u>55299</u>	Т	downstream_gene_variant	OR4G4P	ENSG00000268020	Transcript	ENST00000606857.1	unprocessed_pseudogene	<u>rs103</u>
	<u>1:55299-</u> <u>55299</u>	Т	upstream_gene_variant	OR4G11P	ENSG00000240361	Transcript	ENST00000642116.1	IncRNA	<u>rs103</u>
	<u>1:566186-</u> <u>566186</u>	С	intergenic_variant	-	-	-	-	-	-
	1:568709- 568709	G	intergenic variant	-	-	-	-	-	-
	<u>1:601077-</u> <u>601077</u>	Т	upstream_gene_variant	AL669831.3	ENSG00000230021	Transcript	ENST00000357876.6	IncRNA	-
	<u>1:601077-</u> <u>601077</u>	Т	intron_variant, non_coding_transcript_variant	AL669831.3	ENSG00000230021	Transcript	ENST00000419394.2	IncRNA	-
	1:601077- 601077	Т	intron_variant, non_coding_transcript_variant	AL669831.3	ENSG00000230021	Transcript	ENST00000440196.3	IncRNA	-
	<u>1:601077-</u> <u>601077</u>	Т	downstream_gene_variant	AL669831.3	ENSG00000230021	Transcript	ENST00000440200.5	IncRNA	-
	<u>1:601077-</u> <u>601077</u>	Т	intron_variant, non_coding_transcript_variant	AL669831.3	ENSG00000230021	Transcript	ENST00000634337.2	IncRNA	-

Variant Effect Predictor results - Homo sapiens - Ensembl genome browser 97

Uploaded variant	Location	Allele	Consequence	Symbol	Gene	Feature type	Feature	Biotype	Exist varia
	<u>1:601077-</u> <u>601077</u>	Т	intron_variant, non_coding_transcript_variant	AL669831.3	ENSG00000230021	Transcript	ENST00000634833.2	IncRNA	-
	<u>1:601077-</u> <u>601077</u>	Т	intron_variant, non_coding_transcript_variant	AL669831.3	ENSG00000230021	Transcript	ENST00000635509.2	IncRNA	-
	<u>1:601077-</u> <u>601077</u>	Т	intron_variant, non_coding_transcript_variant	AL669831.3	ENSG00000230021	Transcript	ENST00000641296.1	IncRNA	-
	<u>1:601077-</u> <u>601077</u>	Т	intron_variant, non_coding_transcript_variant	AL669831.3	ENSG00000230021	Transcript	ENST00000648019.1	IncRNA	-
4									•

Page: 4 1 of 5902 | **Show:** $\underline{1}$ 5 $\underline{10}$ 50 All variants

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