

Ipp Cas9-KO Strategy

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Project Overview

Project Name

Ipp

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Ipp* gene. The schematic diagram is as follows:



- The *Ipp* gene has 2 transcripts. According to the structure of *Ipp* gene, exon3-exon8 of *Ipp*-202 (ENSMUST00000106479.7) transcript is recommended as the knockout region. The region contains 1238bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Ipp* gene. The brief process is as follows: CRISPR/Cas9 system will

- The *Ipp* gene is located on the Chr4. If the knockout mice are crossed with other mice strains to obtain double gene positive homozygous mouse offspring, please avoid the two genes on the same chromosome.
- This Strategy is designed based on genetic information in existing databases. Due to the complexity of biological processes, all risk of the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)

Ipp IAP promoted placental gene [Mus musculus (house mouse)]

Gene ID: 16351, updated on 31-Jan-2019

Summary



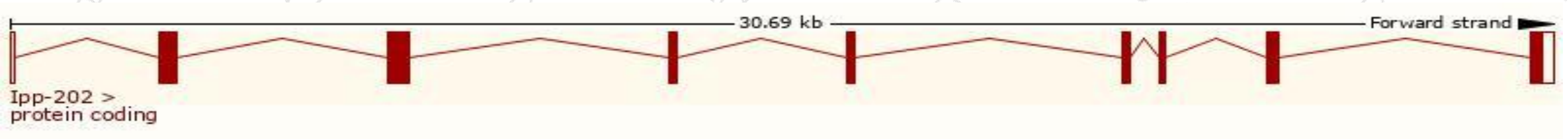
Official Symbol	Ipp provided by MGI
Official Full Name	IAP promoted placental gene provided by MGI
Primary source	MGI:MGI:96581
See related	Ensembl:ENSMUSG00000028696
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	D4Jhu8, Mipp
Expression	Ubiquitous expression in liver E14 (RPKM 3.6), liver E14.5 (RPKM 2.9) and 26 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

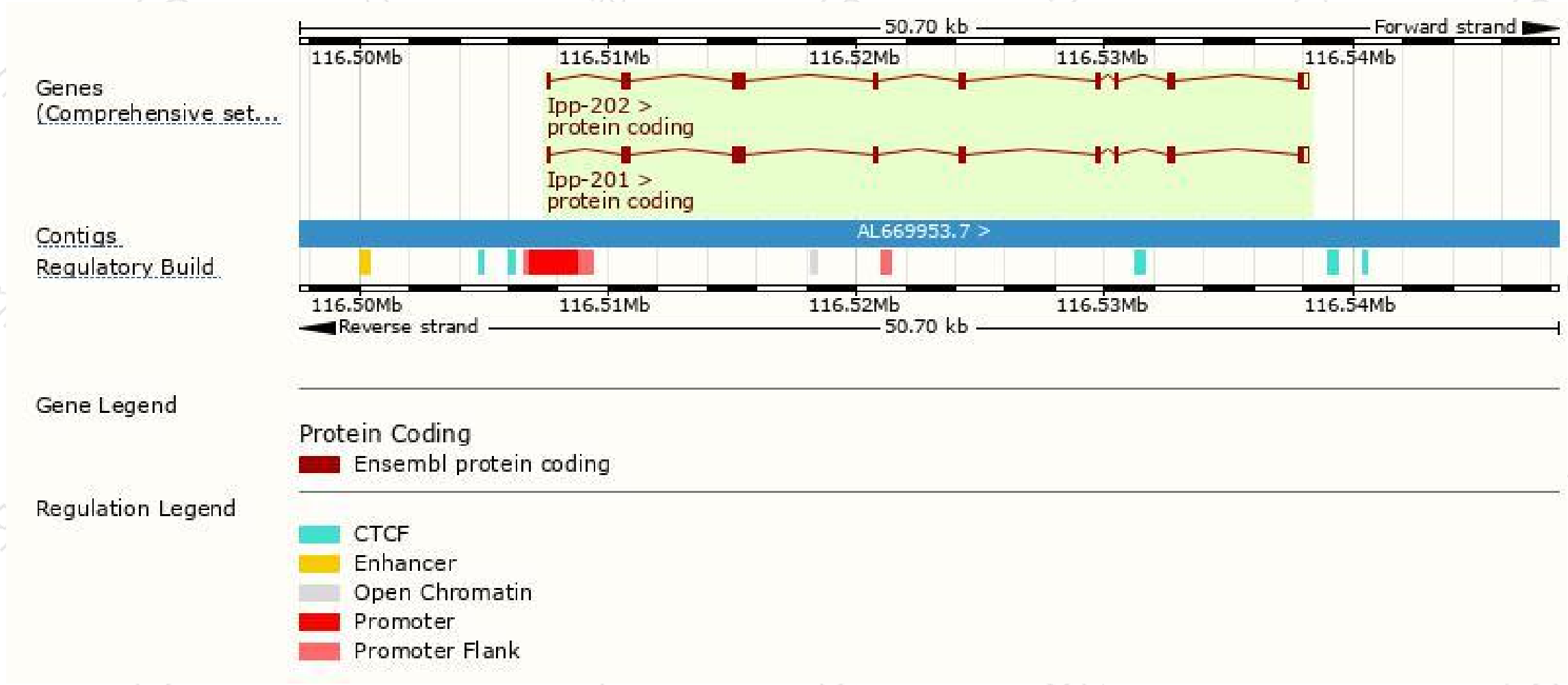
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ipp-202	ENSMUST00000106479.7	2118	584aa	Protein coding	CCDS38851	P28575	TSL:1 GENCODE basic APPRIS P1
Ipp-201	ENSMUST00000030461.4	2115	584aa	Protein coding	CCDS38851	P28575	TSL:5 GENCODE basic APPRIS P1

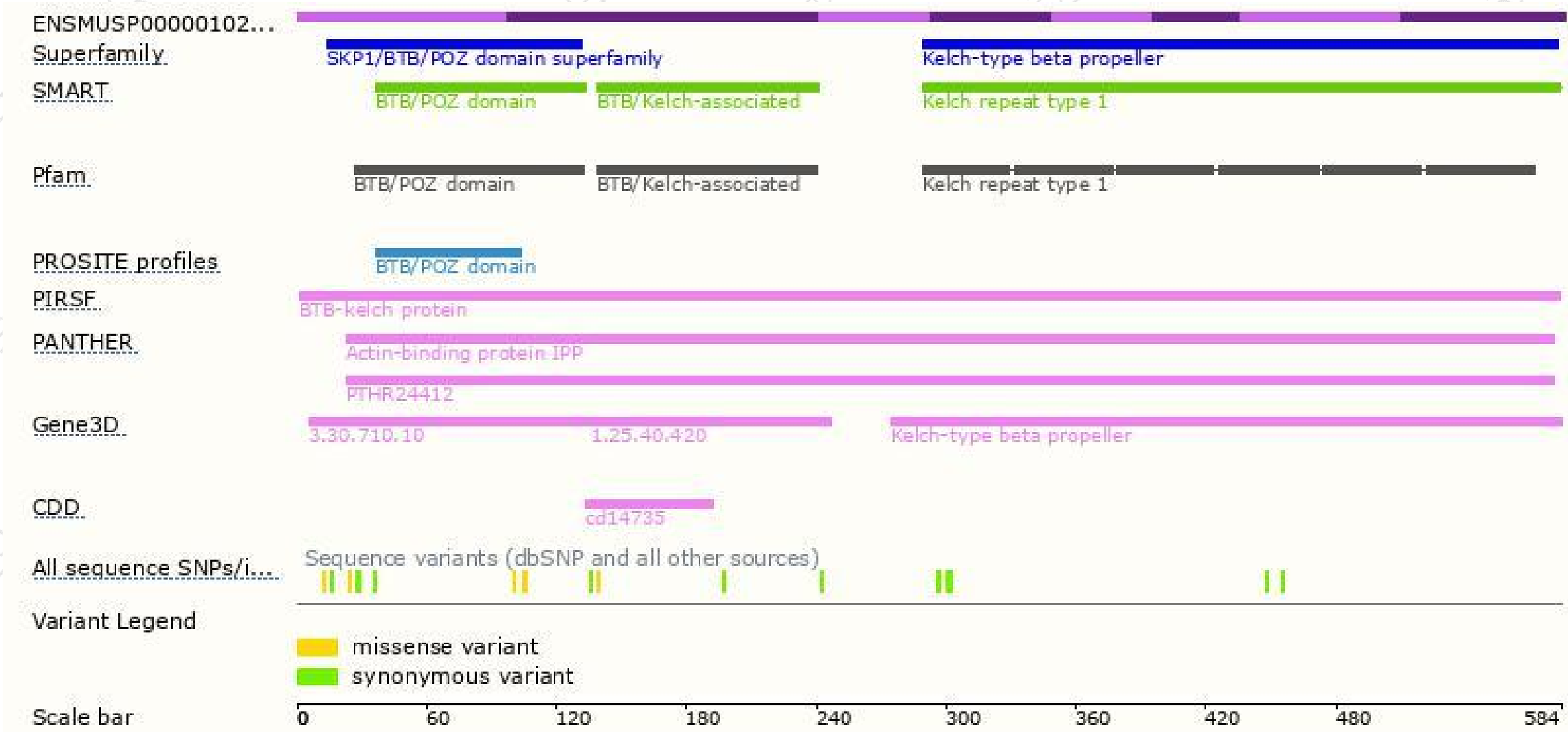
The strategy is based on the design of *Ipp-202* transcript,The transcription is shown below



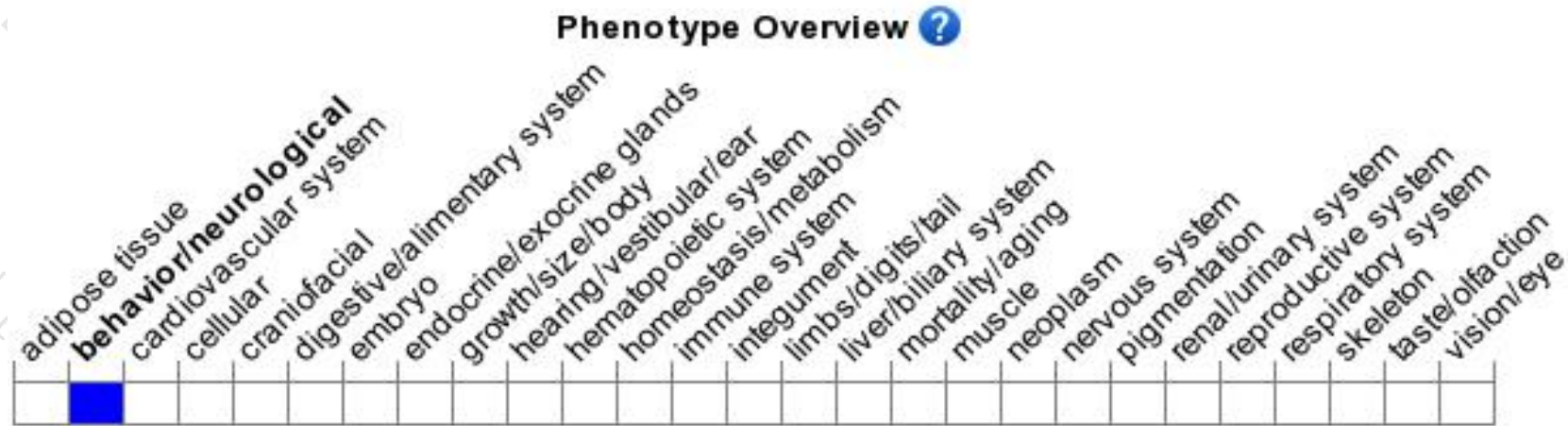
Genomic location distribution



Protein domain



Mouse phenotype description(MGI)



Phenotypes affected by the gene are marked in blue. Data quoted from MGI database(<http://www.informatics.jax.org/>).

If you have any questions, you are welcome to inquire.

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