

Hsh2d Cas9-KO Strategy

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Project Overview

Project Name

Hsh2d

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Hsh2d* gene. The schematic diagram is as follows:



- The *Hsh2d* gene has 2 transcripts. According to the structure of *Hsh2d* gene, exon3-exon5 of *Hsh2d-201* (ENSMUST00000072097.13) transcript is recommended as the knockout region. The region contains 376bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Hsh2d* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice homozygous for a knock-out allele exhibit enhanced IL-2 production, increased T cell proliferation in response to TCR/CD28 stimulation, splenomegaly, and an increased frequency of activated T cells.
- The *Hsh2d* gene is located on the Chr8. 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)

Hsh2d hematopoietic SH2 domain containing [Mus musculus (house mouse)]

Gene ID: 209488, updated on 31-Jan-2019

Summary



Official Symbol	Hsh2d provided by MGI
Official Full Name	hematopoietic SH2 domain containing provided by MGI
Primary source	MGI:MGI:2676364
See related	Ensembl:ENSMUSG00000062007
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	ALX, Hsh2
Expression	Biased expression in spleen adult (RPKM 12.1), thymus adult (RPKM 5.5) and 9 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

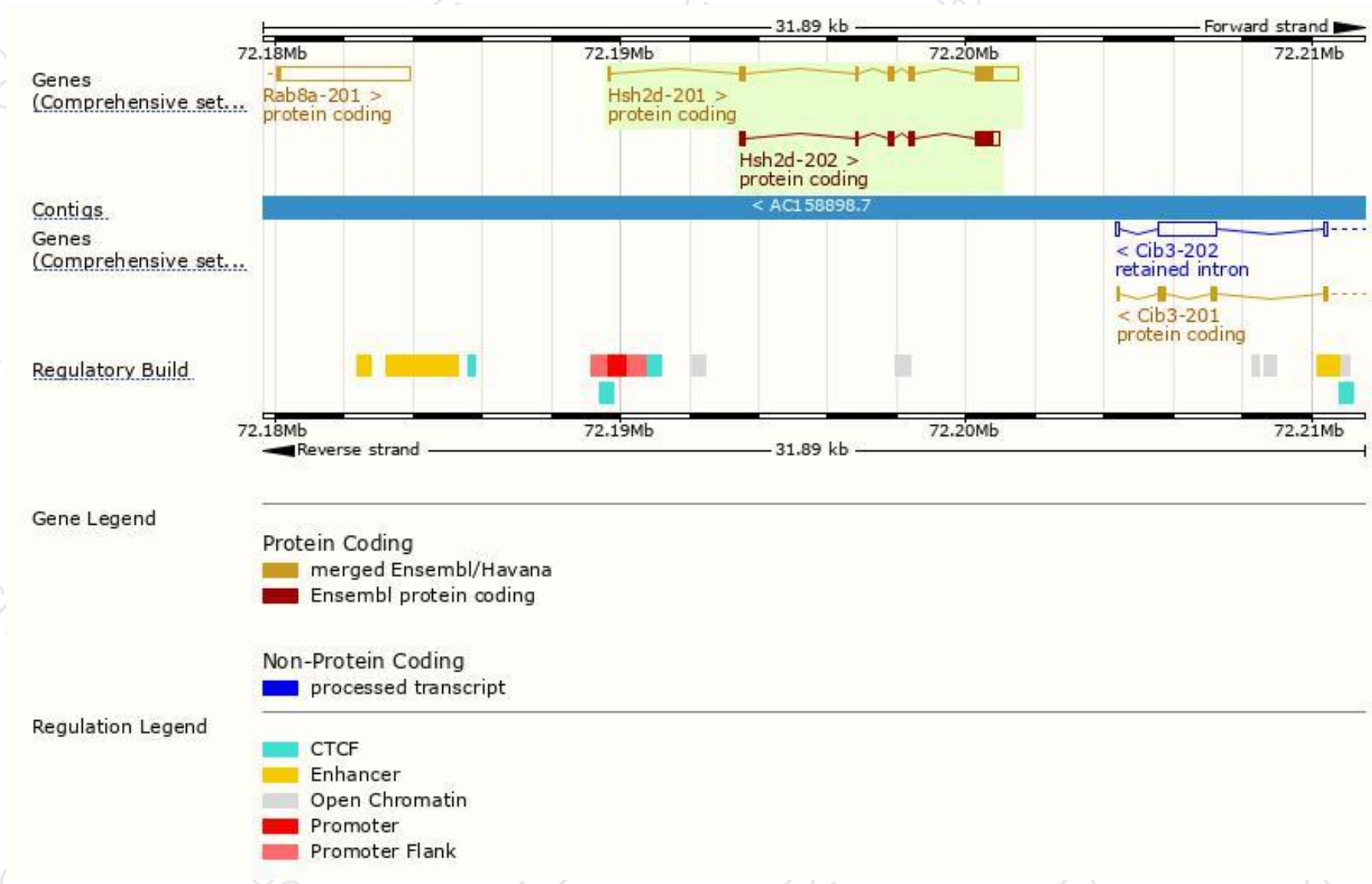
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Hsh2d-201	ENSMUST00000072097.13	1824	334aa	Protein coding	CCDS22410	Q6VYH9	TSL:1 Gencode basic APPRIS P1
Hsh2d-202	ENSMUST00000165324.1	1238	334aa	Protein coding	CCDS22410	Q6VYH9	TSL:1 Gencode basic APPRIS P1

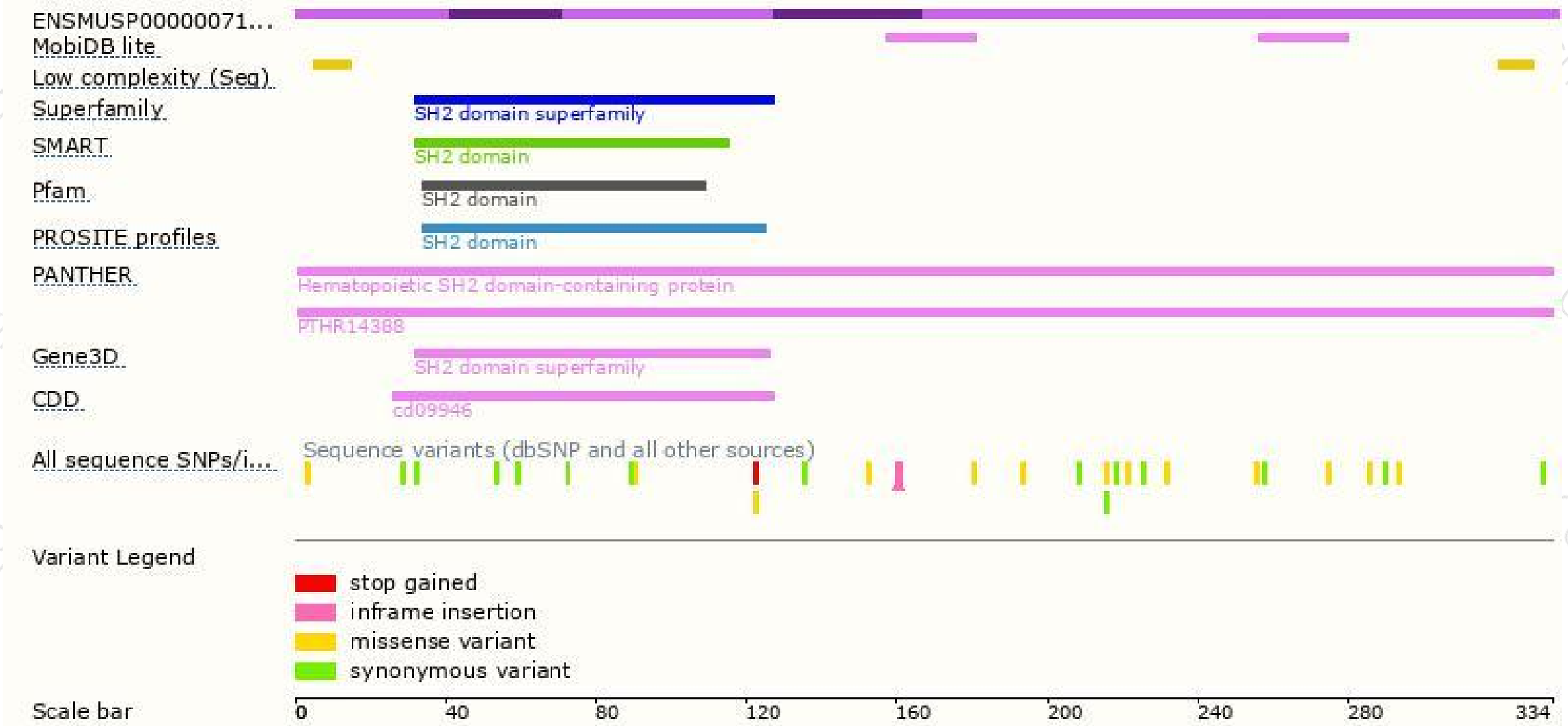
The strategy is based on the design of *Hsh2d-201* transcript,The transcription is shown below



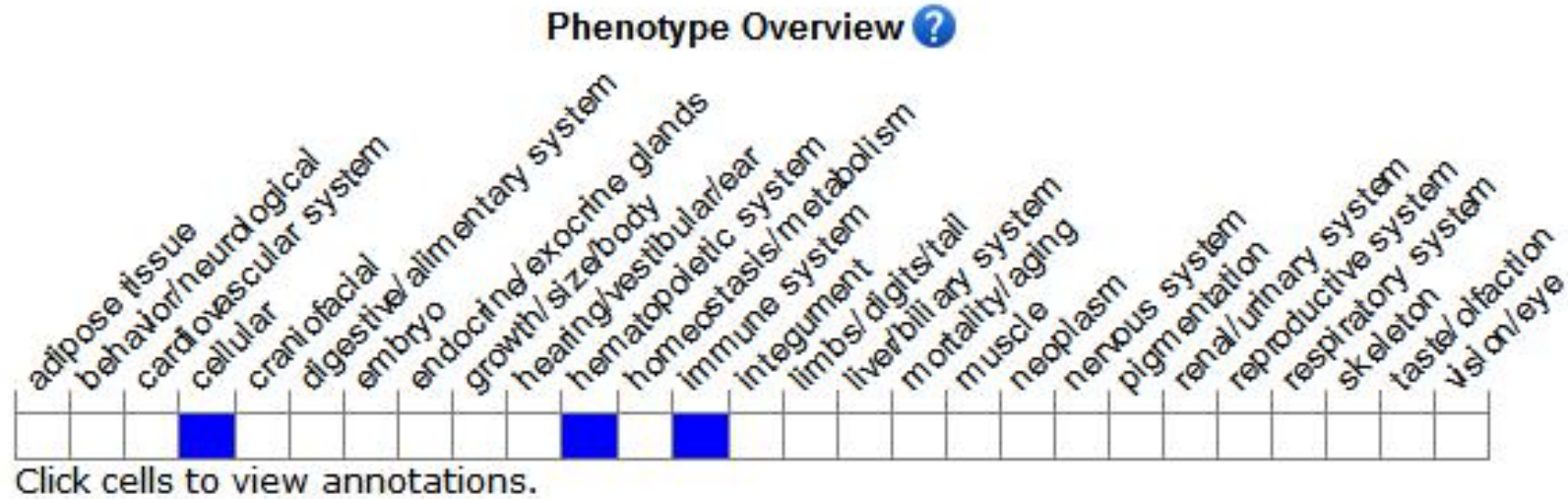
Genomic location distribution



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/>).

According to the existing MGI data, Mice homozygous for a knock-out allele exhibit enhanced IL-2 production, increased T cell proliferation in response to TCR/CD28 stimulation, splenomegaly, and an increased frequency of activated T cells.

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

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