

Dusp10 Cas9-KO Strategy

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Project Overview

Project Name

Dusp10

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Dusp10* gene. The schematic diagram is as follows:



- The *Dusp10* gene has 2 transcripts. According to the structure of *Dusp10* gene, exon2 of *Dusp10-201* (ENSMUST00000048655.7) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Dusp10* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice homozygous for a knock-out allele display alterations in both innate and adaptive immune responses.
- The *Dusp10* gene is located on the Chr1. 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)

Dusp10 dual specificity phosphatase 10 [Mus musculus (house mouse)]

Gene ID: 63953, updated on 31-Jan-2019

Summary



Official Symbol	Dusp10 provided by MGI
Official Full Name	dual specificity phosphatase 10 provided by MGI
Primary source	MGI:MGI:1927070
See related	Ensembl:ENSMUSG00000039384
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	2610306G15Rik, A158871, MKP-5, MKP5
Expression	Broad expression in thymus adult (RPKM 37.9), testis adult (RPKM 13.3) and 21 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

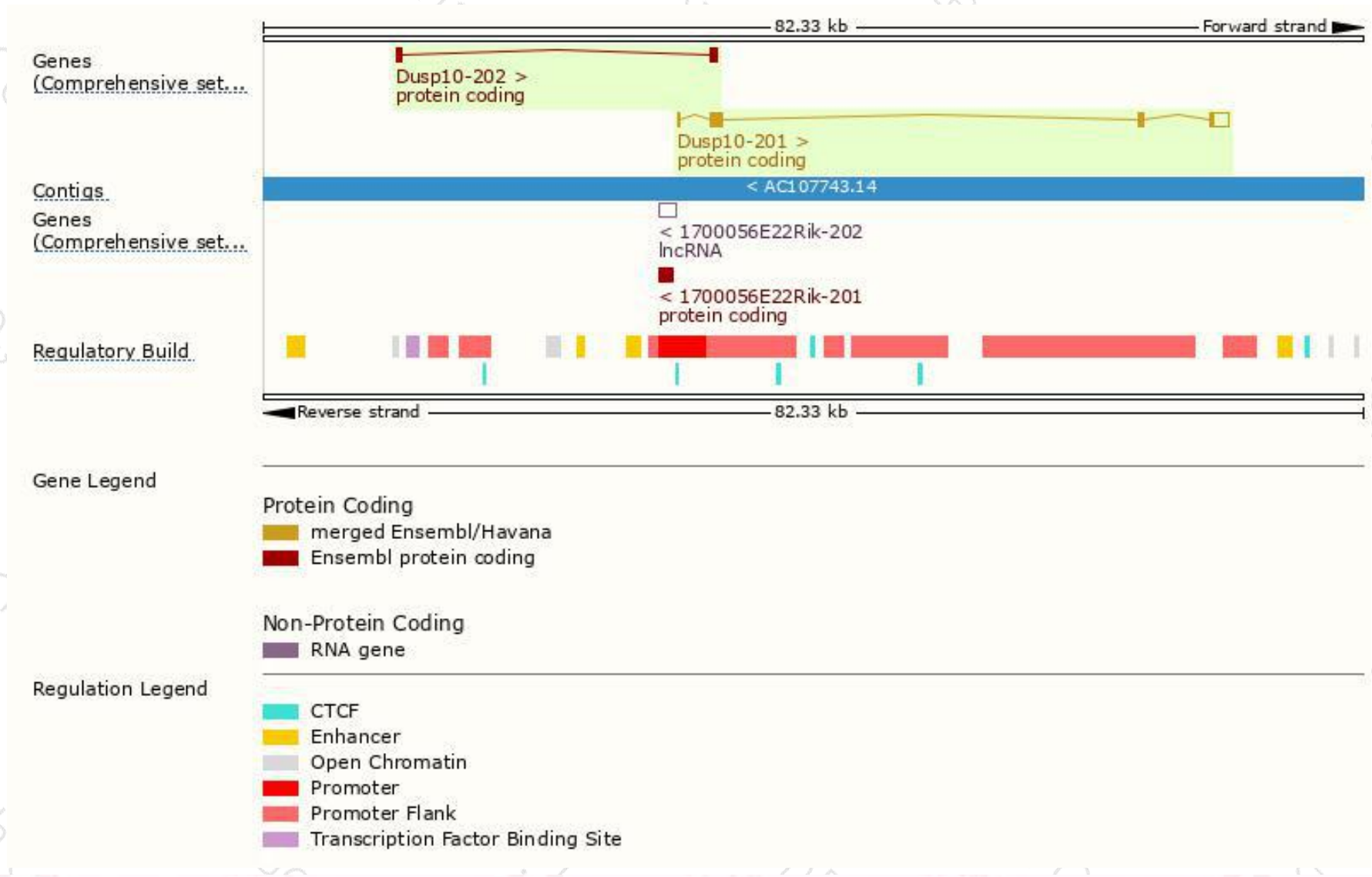
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Dusp10-201	ENSMUST00000048655.7	2792	483aa	Protein coding	CCDS15591	Q9ESS0	TSL:1 GENCODE basic APPRIS P1
Dusp10-202	ENSMUST00000139839.1	724	145aa	Protein coding	-	D3Z2S1	CDS 3' incomplete TSL:3

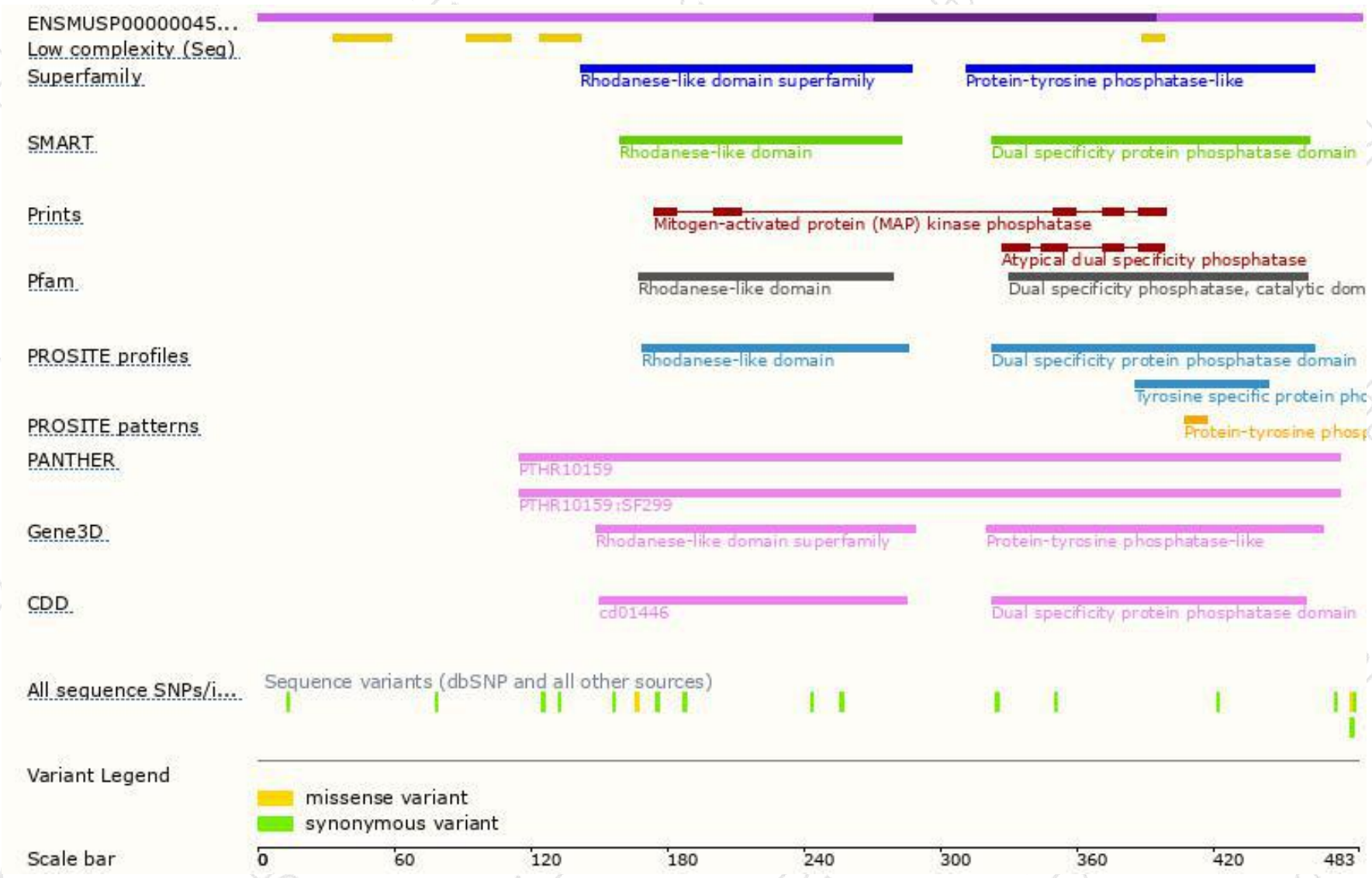
The strategy is based on the design of *Dusp10-201* transcript,The transcription is shown below



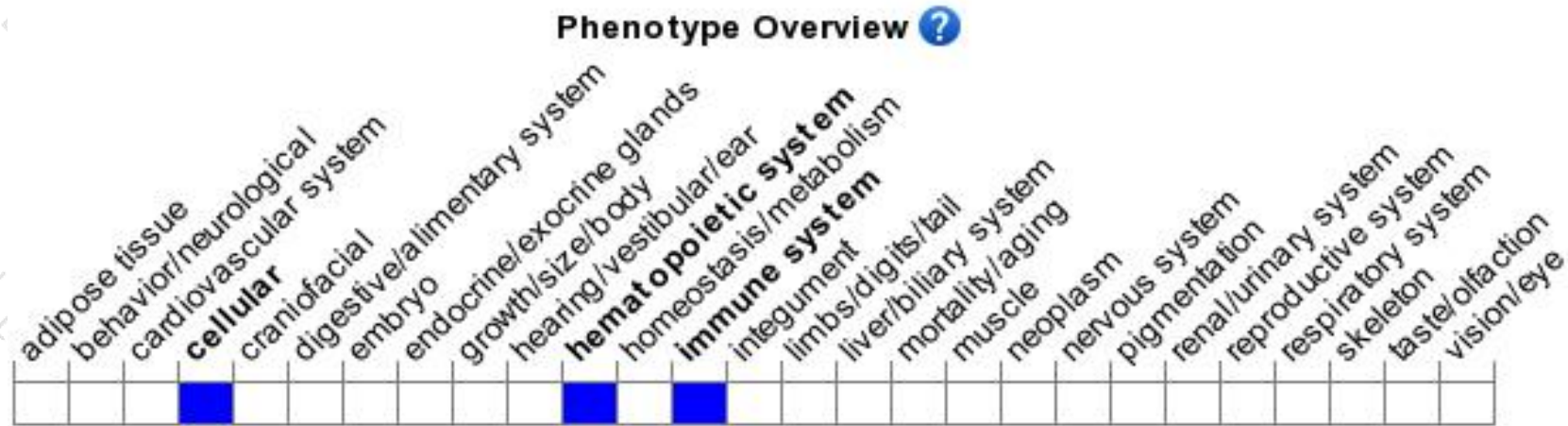
Genomic location distribution



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 display alterations in both innate and adaptive immune responses.

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

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