

Dusp5 Cas9-KO Strategy

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Reviewer:

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Project Overview

Project Name

Dusp5

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Dusp5* gene. The schematic diagram is as follows:



- The *Dusp5* gene has 1 transcript. According to the structure of *Dusp5* gene, exon2 of *Dusp5-201* (ENSMUST00000038287.6) transcript is recommended as the knockout region. The region contains 149bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Dusp5* gene. The brief process is as follows: CRISPR/Cas9 system v

- According to the existing MGI data, Mice homozygous for a knock-out allele exhibit increased proliferation and apoptosis and altered metabolic profiles in T cells. Mice homozygous for other null alleles exhibit altered eosinophilic response to parasitic infection and increased susceptibility to induced tumors.
- The 5-terminal regulation of *Gm50394* may be affected.
- The *Dusp5* gene is located on the Chr19. 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)

Dusp5 dual specificity phosphatase 5 [*Mus musculus* (house mouse)]

Gene ID: 240672, updated on 28-Dec-2019

Summary

Official Symbol Dusp5 provided by [MGI](#)
Official Full Name dual specificity phosphatase 5 provided by [MGI](#)
Primary source [MGI:MGI:2685183](#)
See related [Ensembl:ENSMUSG00000034765](#)
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 Gm337
Expression Broad expression in thymus adult (RPKM 25.5), duodenum adult (RPKM 24.5) and 19 other tissues [See more](#)
Orthologs [human](#) [all](#)

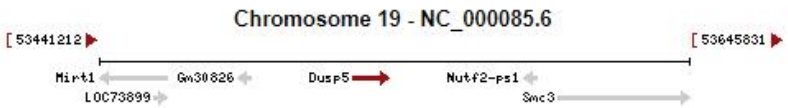
Genomic context

Location: 19; 19 D2

See Dusp5 in [Genome Data Viewer](#)

Exon count: 4

Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	19	NC_000085.6 (53529145..53541809)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	19	NC_000085.5 (53603808..53615812)

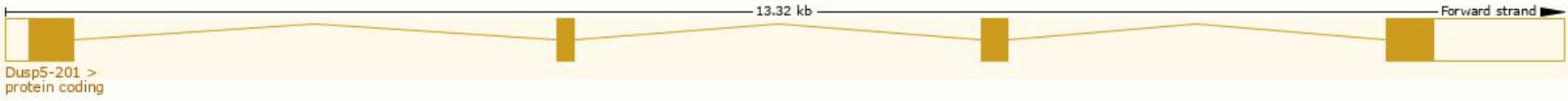


Transcript information (Ensembl)

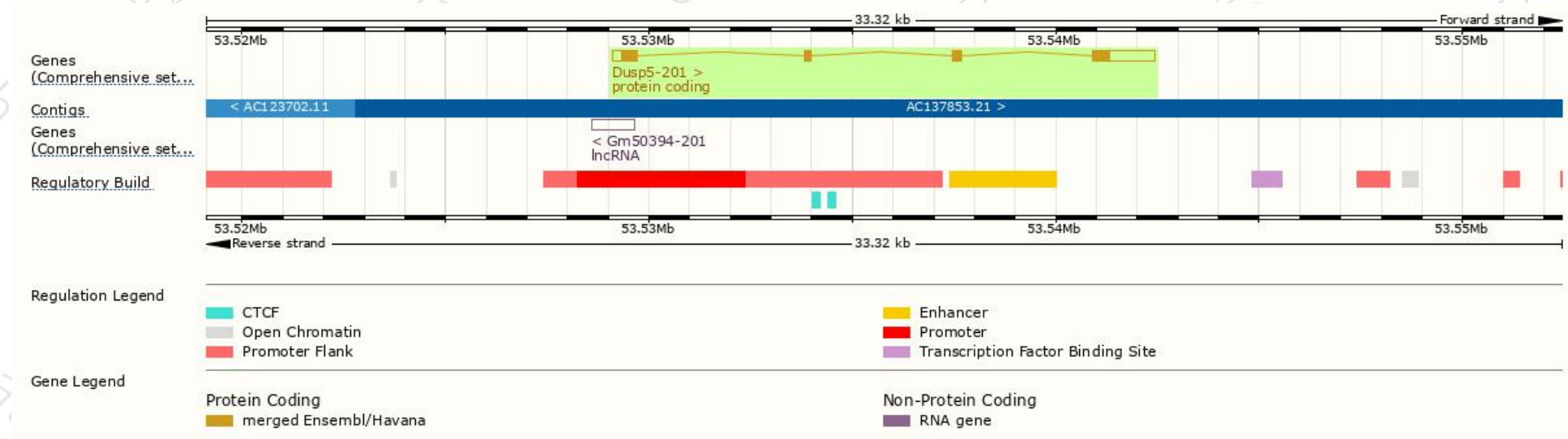
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Dusp5-201	ENSMUST00000038287.6	2473	384aa	Protein coding	CCDS38024	Q1HL35	TSL:5 Gencode basic APPRIS P1

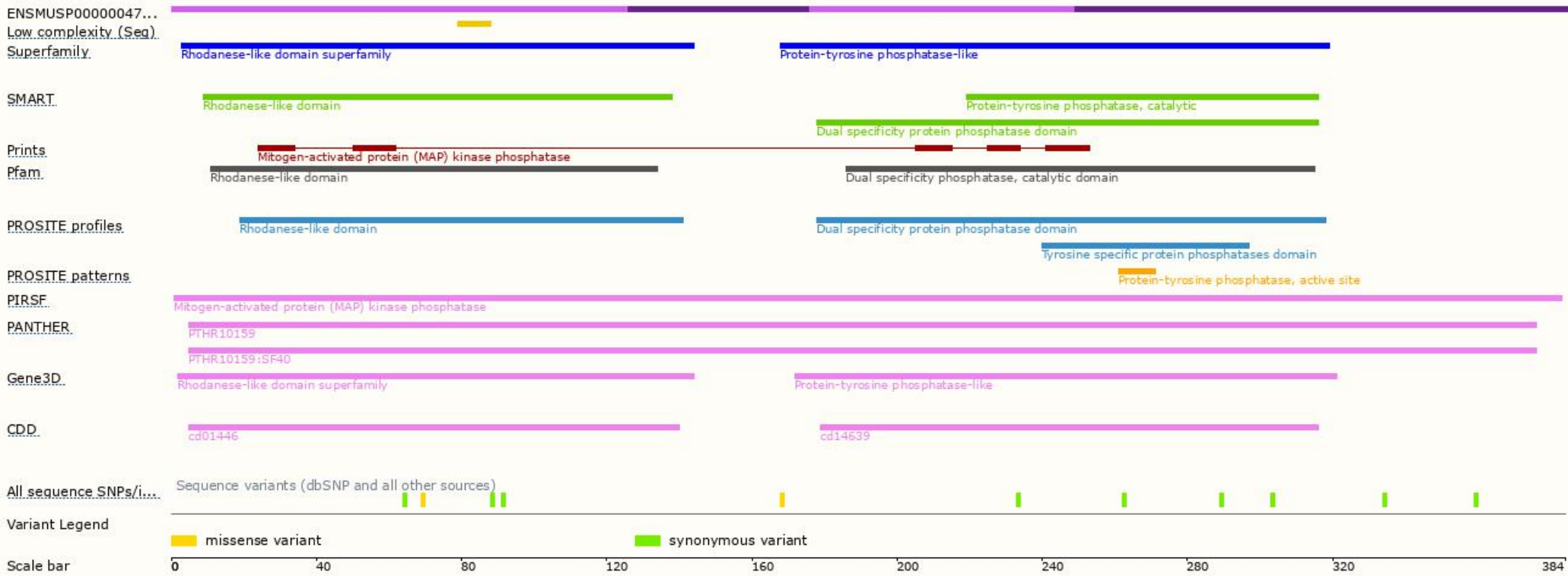
The strategy is based on the design of *Dusp5-201* transcript, The transcription is shown below



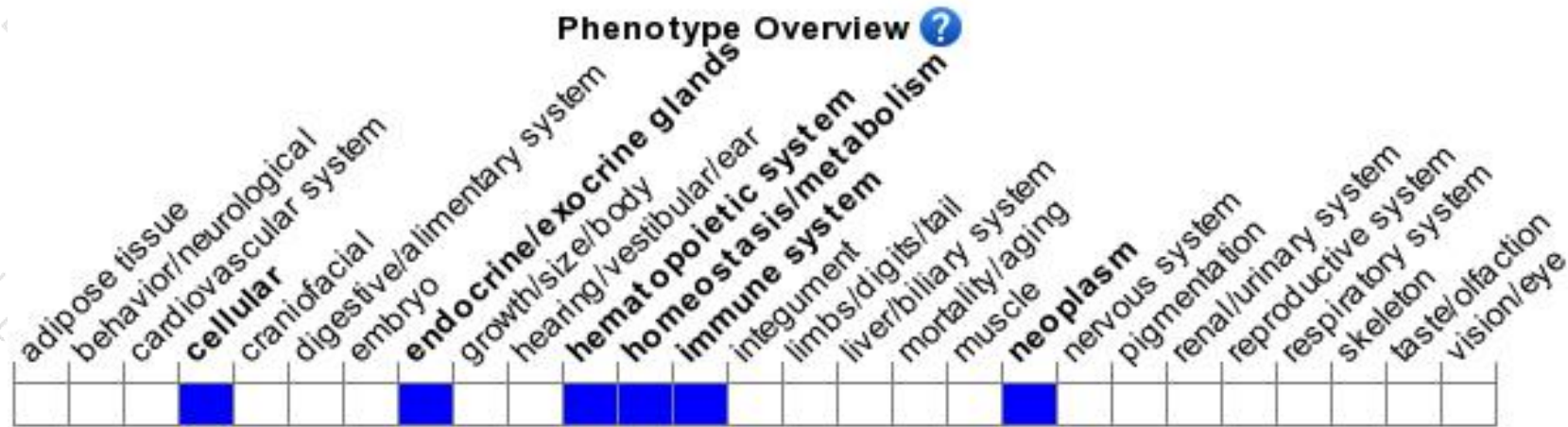
Genomic location distribution



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 increased proliferation and apoptosis and altered metabolic profiles in T cells. Mice homozygous for other null alleles exhibit altered eosinophilic response to parasitic infection and increased susceptibility to induced tumors.

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

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