

Dusp6 Cas9-KO Strategy

Designer:

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Design Date:

2019-7-18

Project Overview

Project Name

Dusp6

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Dusp6* gene. The schematic diagram is as follows:



- The *Dusp6* gene has 4 transcripts. According to the structure of *Dusp6* gene, exon1-exon3 of *Dusp6-201* (ENSMUST00000020118.4) transcript is recommended as the knockout region. The region contains all of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Dusp6* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice homozygous or heterozygous for a null mutation display partial penetrance of postnatal lethality, reduced body weight, and abnormal growth plate morphology.
- The strategy affects the 5-terminal regulation of Gm48089 and affects the 3-terminal regulation of Gm34921.
- The *Dusp6* gene is located on the Chr10. 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)

Dusp6 dual specificity phosphatase 6 [Mus musculus (house mouse)]

Gene ID: 67603, updated on 5-Mar-2019

Summary



Official Symbol	Dusp6 provided by MGI
Official Full Name	dual specificity phosphatase 6 provided by MGI
Primary source	MGI:MGI:1914853
See related	Ensembl:ENSMUSG00000019960
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	1300019I03Rik, MKP-3, MKP3, PYST1
Expression	Ubiquitous expression in lung adult (RPKM 46.1), adrenal adult (RPKM 45.5) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

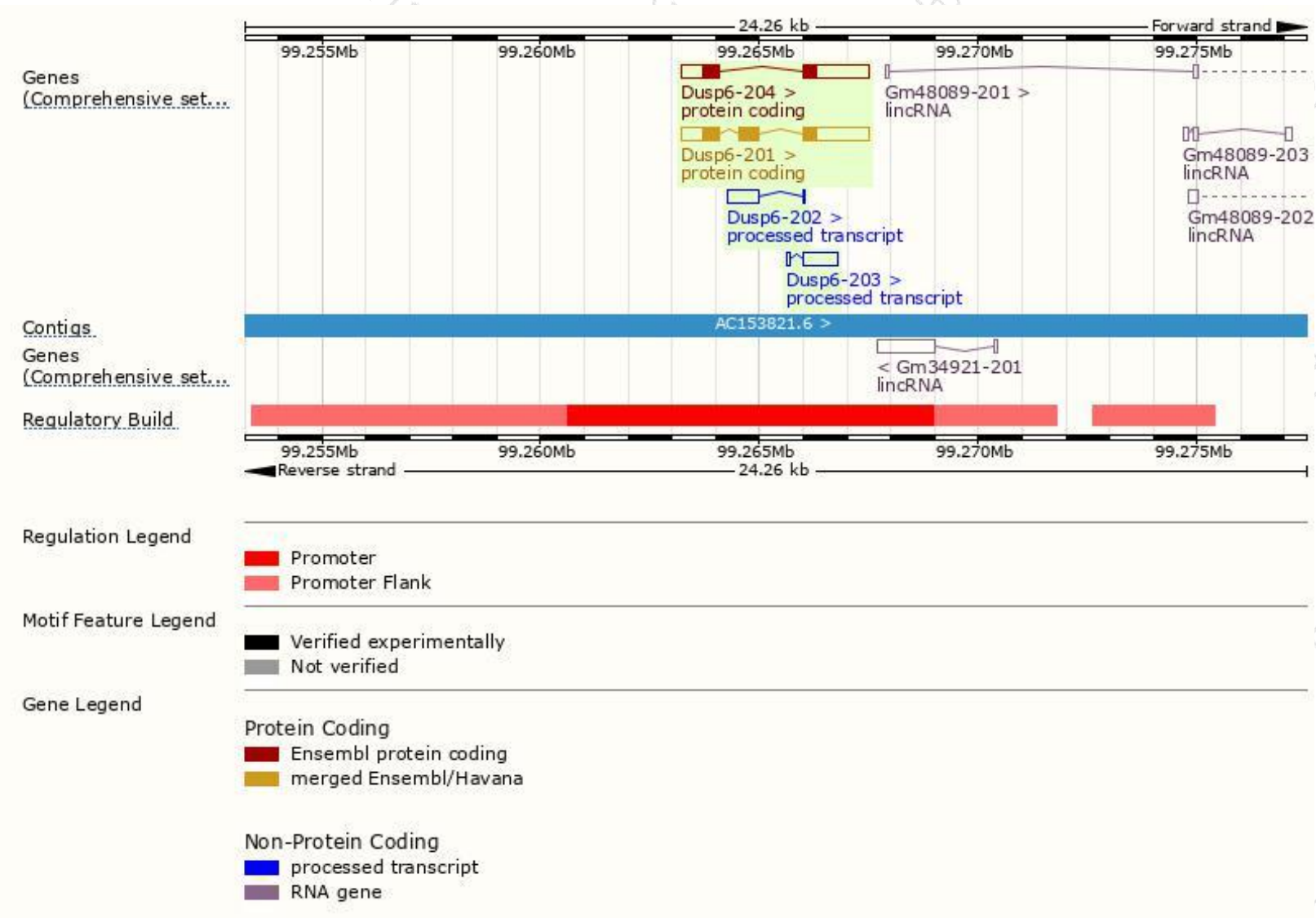
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Dusp6-201	ENSMUST00000020118.4	2796	381aa	Protein coding	CCDS24147	Q9DBB1	TSL:1 GENCODE basic APPRIS P1
Dusp6-204	ENSMUST00000220291.1	2359	235aa	Protein coding	-	A0A1W2P715	TSL:5 GENCODE basic
Dusp6-203	ENSMUST00000220218.1	876	No protein	Processed transcript	-	-	TSL:2
Dusp6-202	ENSMUST00000219988.1	758	No protein	Processed transcript	-	-	TSL:2

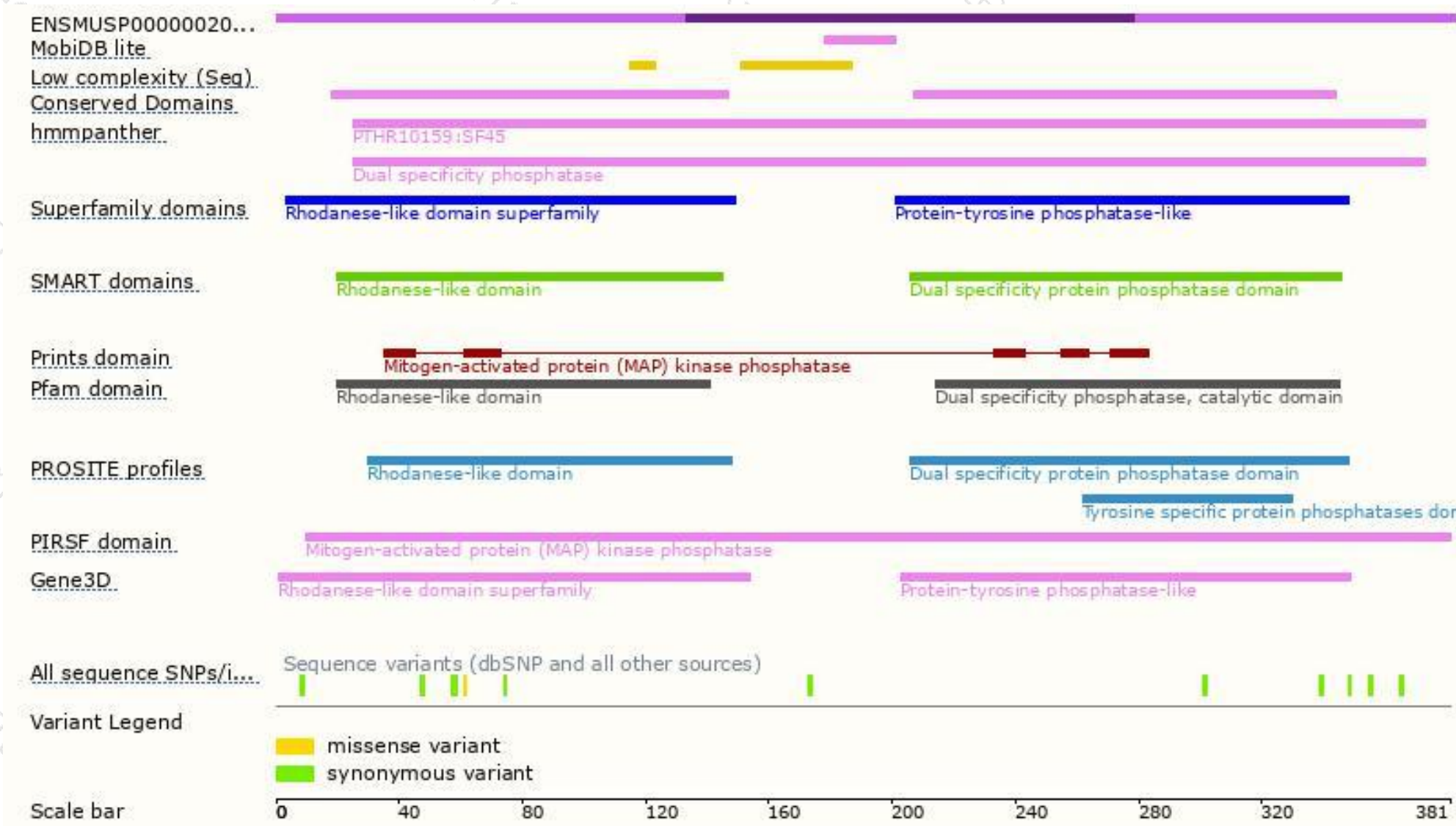
The strategy is based on the design of *Dusp6-201* transcript,The transcription is shown below



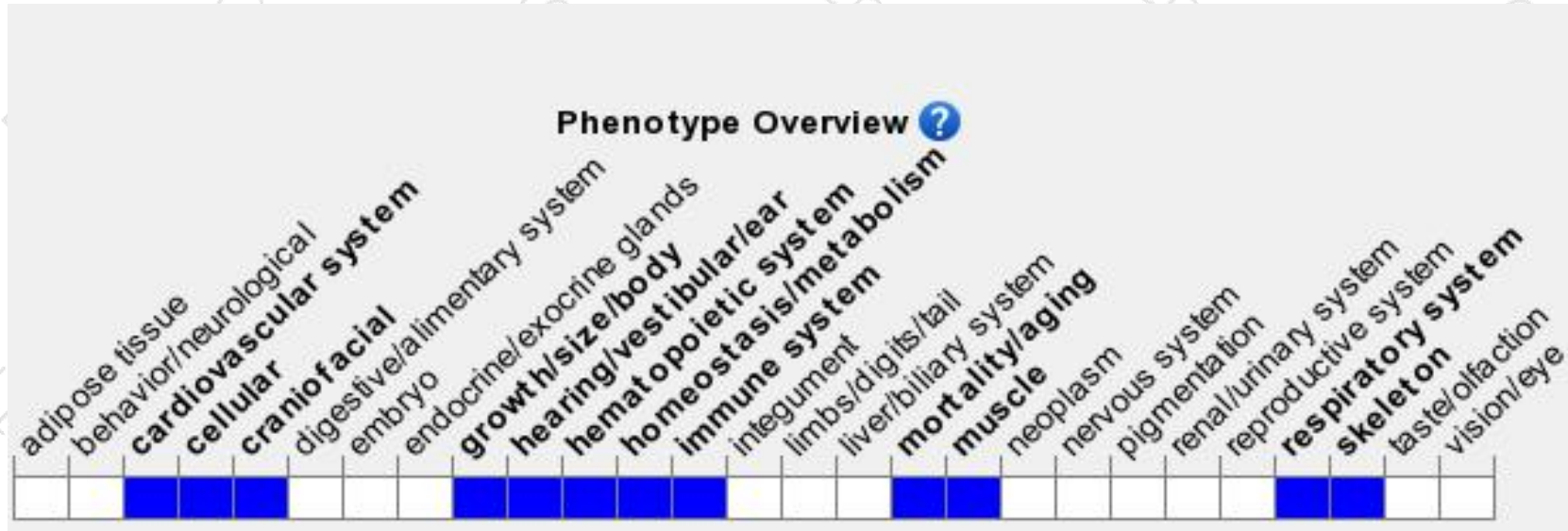
Genomic location distribution



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 or heterozygous for a null mutation display partial penetrance of postnatal lethality, reduced body weight, and abnormal growth plate morphology.

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

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