

Sod1 Cas9-KO Strategy

Designer:

Daohua Xu

Reviewer:

Huimin Su

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Project Overview

Project Name

Sod1

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Sod1* gene. The schematic diagram is as follows:



- The *Sod1* gene has 2 transcripts. According to the structure of *Sod1* gene, exon2 of *Sod1-201* (ENSMUST00000023707.10) transcript is recommended as the knockout region. The region contains 97bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Sod1* gene. The brief process is as follows: CRISPR/Cas9 system v

- According to the existing MGI data, Homozygous mutants exhibit increased motor neuron loss after axonal injury and enhanced susceptibility to ischemic reperfusion injury. Homozygous females have irregular and small litters, and for some alleles exhibit immature ovarian follicles with few corpora lutea.
- The *Sod1* gene is located on the Chr16. 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)

Sod1 superoxide dismutase 1, soluble [Mus musculus (house mouse)]

Gene ID: 20655, updated on 31-Jan-2019

Summary



Official Symbol	Sod1 provided by MGI
Official Full Name	superoxide dismutase 1, soluble provided by MGI
Primary source	MGI:MGI:98351
See related	Ensembl:ENSMUSG00000022982
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	B430204E11Rik, Cu/Zn-SOD, CuZnSOD, Ipo-1, Ipo1, SODC, Sod-1
Expression	Ubiquitous expression in liver adult (RPKM 521.8), liver E14 (RPKM 341.6) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

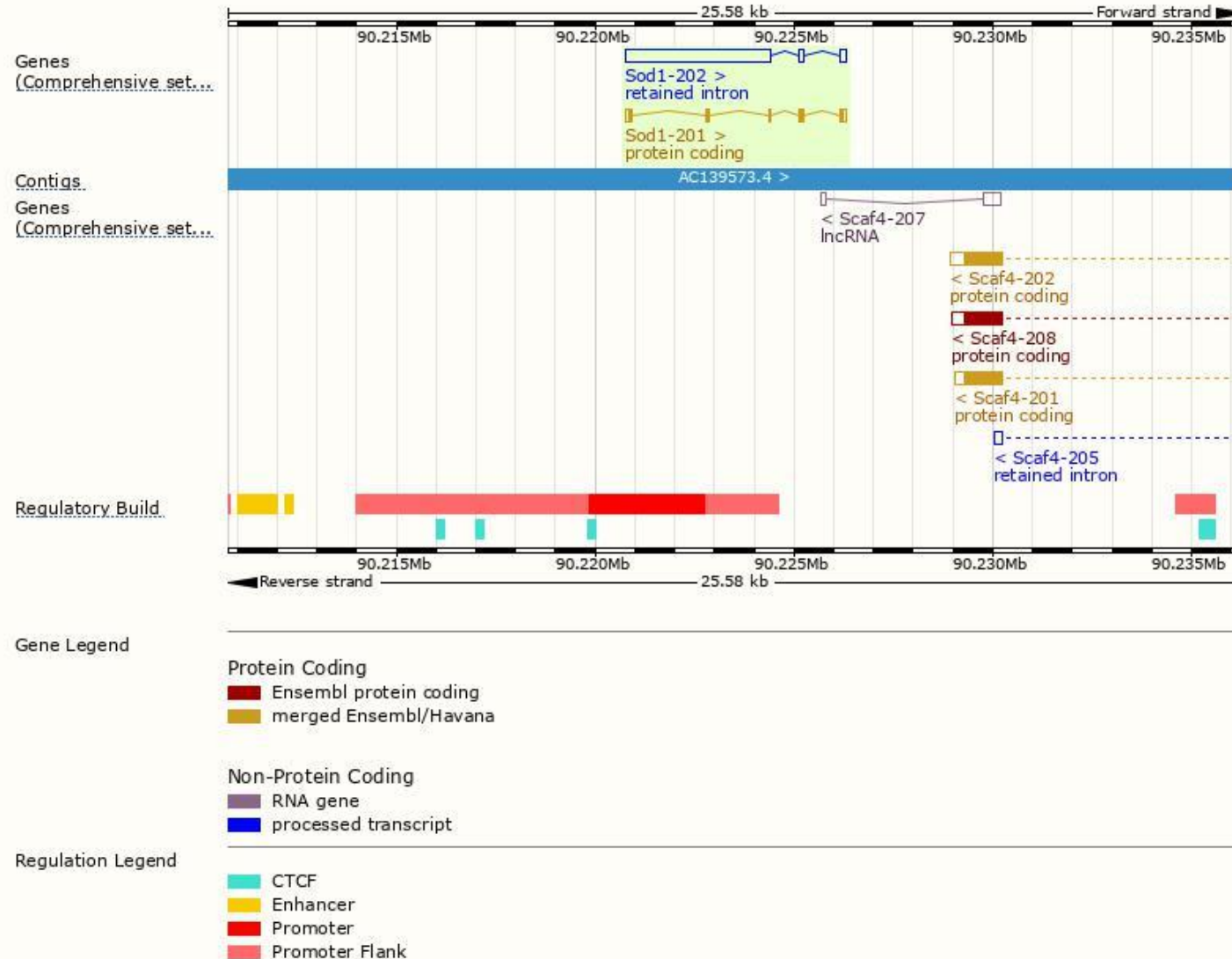
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Sod1-201	ENSMUST00000023707.10	641	154aa	Protein coding	CCDS37395	P08228	TSL:1 Gencode basic APPRIS P1
Sod1-202	ENSMUST000000232505.1	3970	No protein	Retained intron	-	-	-

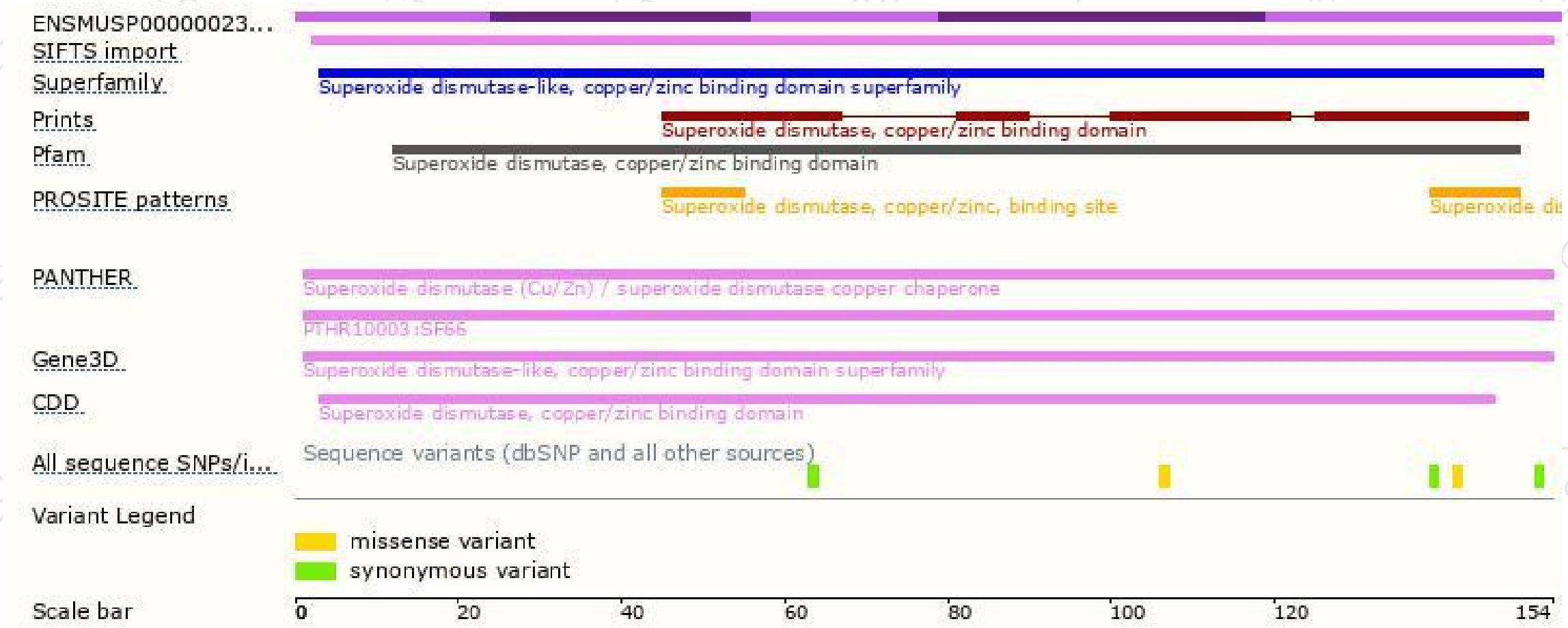
The strategy is based on the design of *Sod1-201* transcript,The transcription is shown below



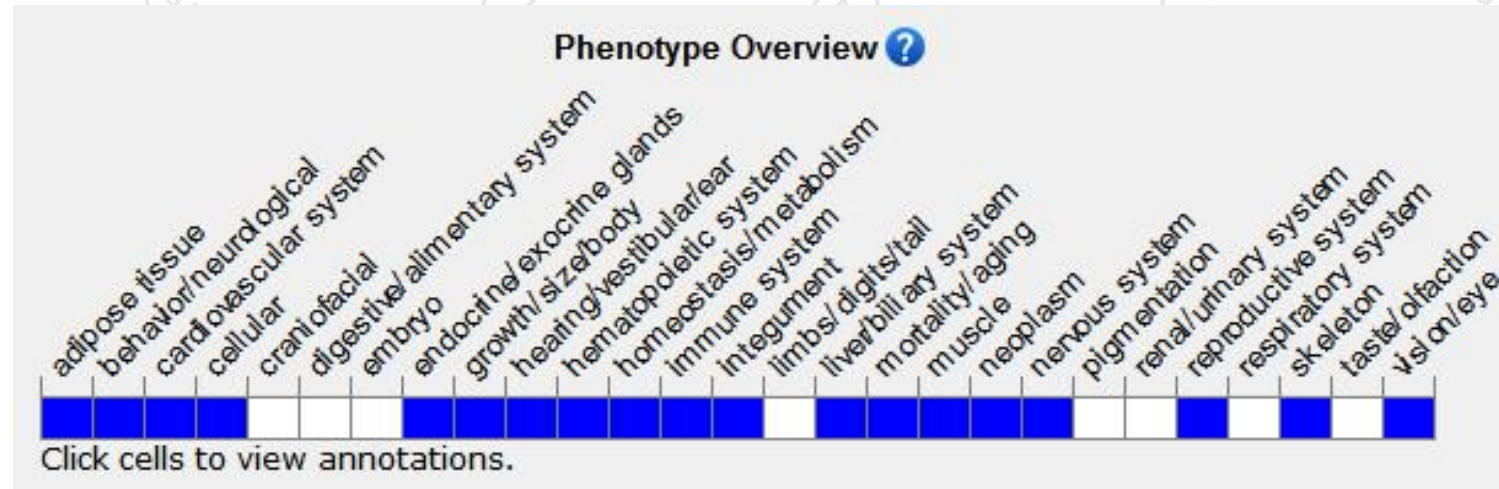
Genomic location distribution



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, Homozygous mutants exhibit increased motor neuron loss after axonal injury and enhanced susceptibility to ischemic reperfusion injury. Homozygous females have irregular and small litters, and for some alleles exhibit immature ovarian follicles with few corpora lutea.

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

Tel: 400-9660890

