

Slc30a8 Cas9-KO Strategy

Designer: Xueting Zhang

Reviewer: Yanhua Shen

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Project Overview

Project Name

Slc30a8

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Slc30a8* gene. The schematic diagram is as follows:



- The *Slc30a8* gene has 1 transcript. According to the structure of *Slc30a8* gene, exon2 of *Slc30a8-201* (ENSMUST00000037240.2) transcript is recommended as the knockout region. The region contains 197bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Slc30a8* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice homozygous for a knock-out allele exhibit reduced islet zinc levels, circulating insulin levels, and glucose-stimulated insulin secretion.
- The *Slc30a8* gene is located on the Chr15. 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)

Slc30a8 solute carrier family 30 (zinc transporter), member 8 [*Mus musculus* (house mouse)]

Gene ID: 239436, updated on 31-Dec-2019

Summary

- Official Symbol** Slc30a8 provided by MGI
- Official Full Name** solute carrier family 30 (zinc transporter), member 8 provided by MGI
- Primary source** MGI:MGI:2442682
- See related** Ensembl:ENSMUSG00000022315
- 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** ZnT8; ZnT-8; C820002P14Rik
- Expression** Low expression observed in reference dataset [See more](#)
- Orthologs** [human](#) [all](#)

Genomic context

Location: 15; 15 C See Slc30a8 in [Genome Data Viewer](#)

Exon count: 8

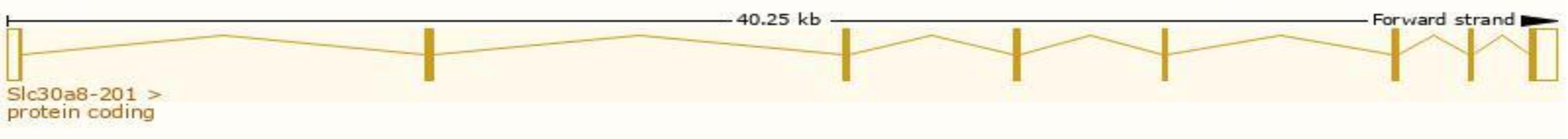
Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	15	NC_000081.6 (52295553..52335798)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	15	NC_000081.5 (52127108..52167288)

Transcript information (Ensembl)

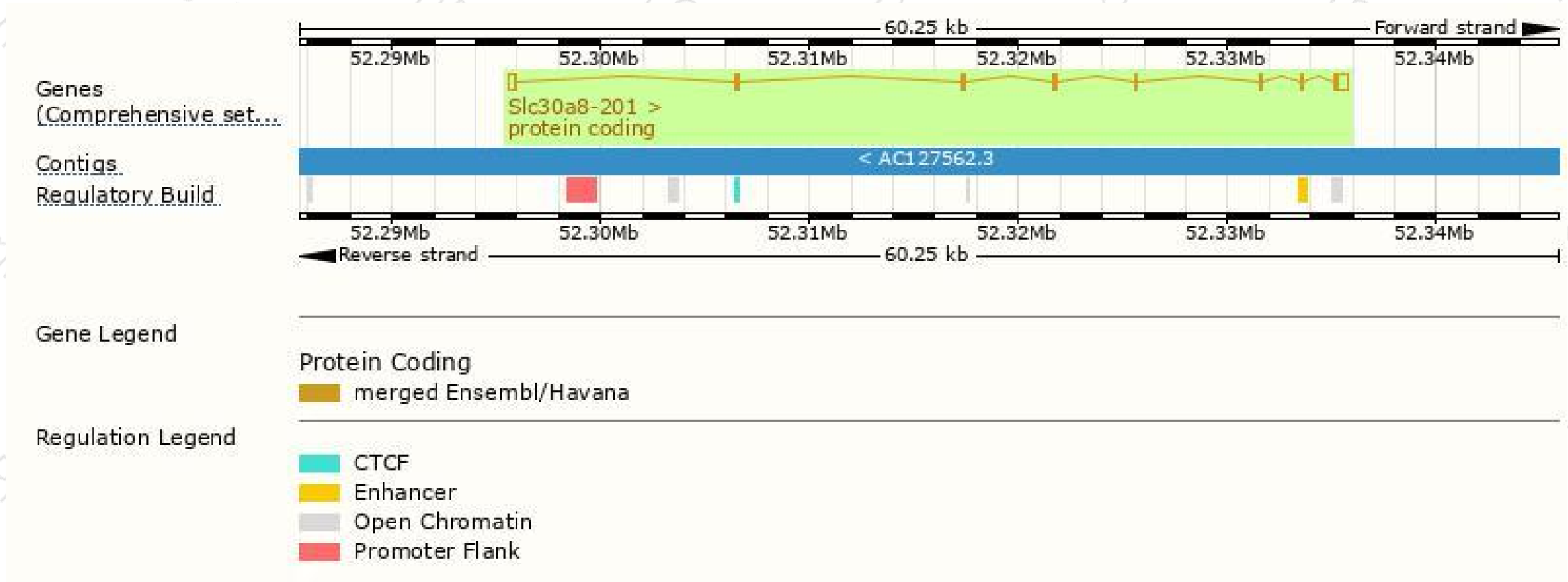
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Slc30a8-201	ENSMUST00000037240.2	1971	367aa	Protein coding	CCDS27464	Q8BGG0	TSL:1 GENCODE basic APPRIS P1

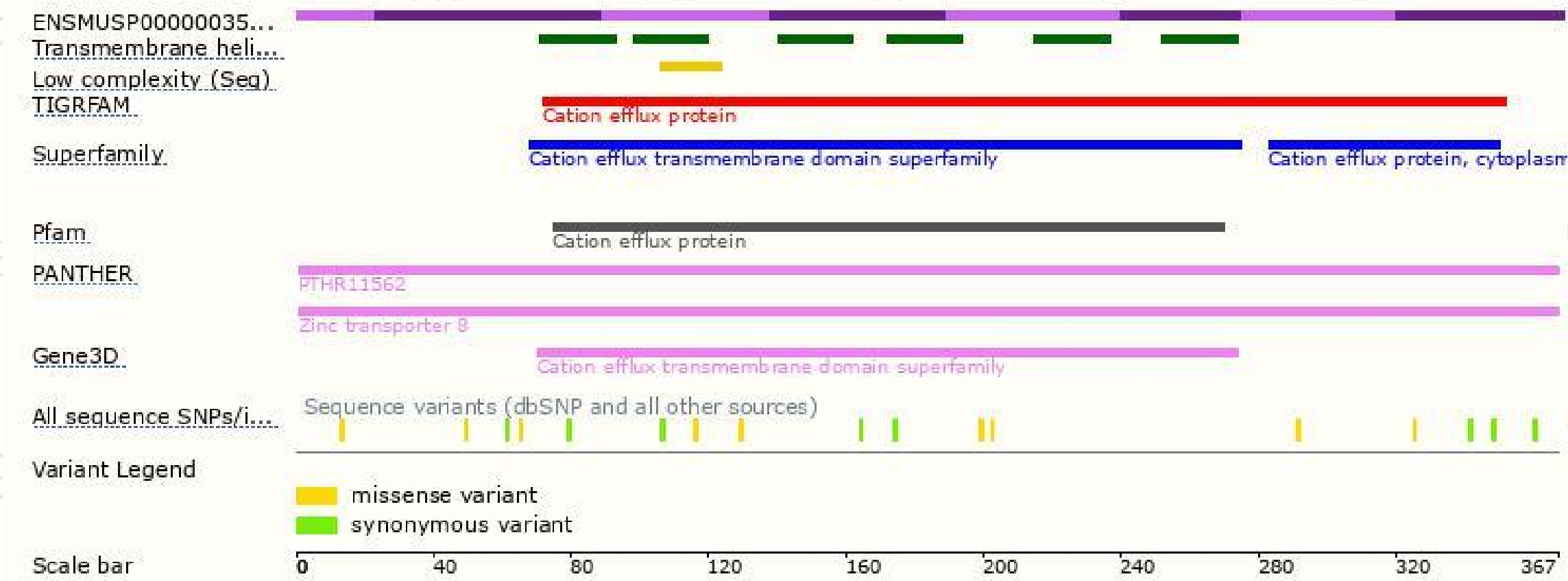
The strategy is based on the design of *Slc30a8-201* transcript, The transcription is shown below



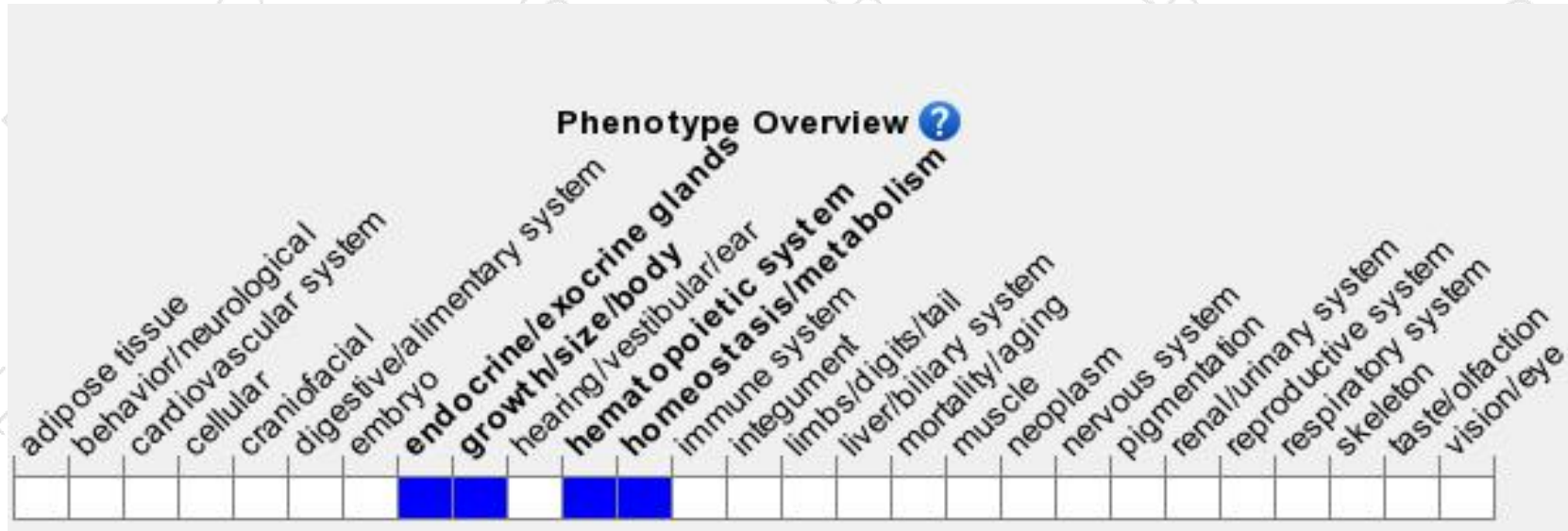
Genomic location distribution



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 reduced islet zinc levels, circulating insulin levels, and glucose-stimulated insulin secretion.

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

Tel: 400-9660890

