

Slc5a8 Cas9-KO Strategy

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Reviewer:

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Project Overview

Project Name

Slc5a8

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Slc5a8* gene. The schematic diagram is as follows:



- The *Slc5a8* gene has 1 transcript. According to the structure of *Slc5a8* gene, exon2-exon5 of *Slc5a8-201* (ENSMUST00000020255.7) transcript is recommended as the knockout region. The region contains 341bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Slc5a8* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice homozygous for a null allele exhibit increased lactate concentrations in the saliva and urine.
- The *Slc5a8* 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)

Slc5a8 solute carrier family 5 (iodide transporter), member 8 [Mus musculus (house mouse)]

Gene ID: 216225, updated on 31-Jan-2019

Summary



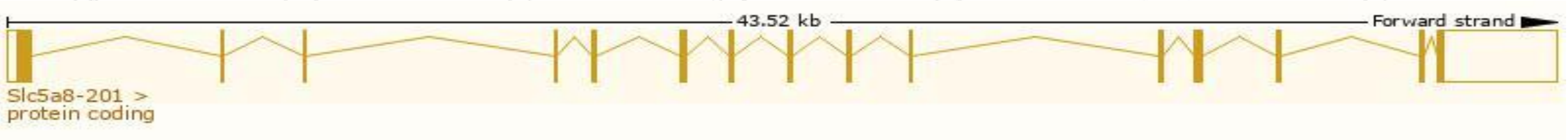
Official Symbol	Slc5a8 provided by MGI
Official Full Name	solute carrier family 5 (iodide transporter), member 8 provided by MGI
Primary source	MGI:MGI:2384916
See related	Ensembl:ENSMUSG00000020062
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Ait, SMCT
Expression	Biased expression in colon adult (RPKM 27.7), large intestine adult (RPKM 16.4) and 2 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

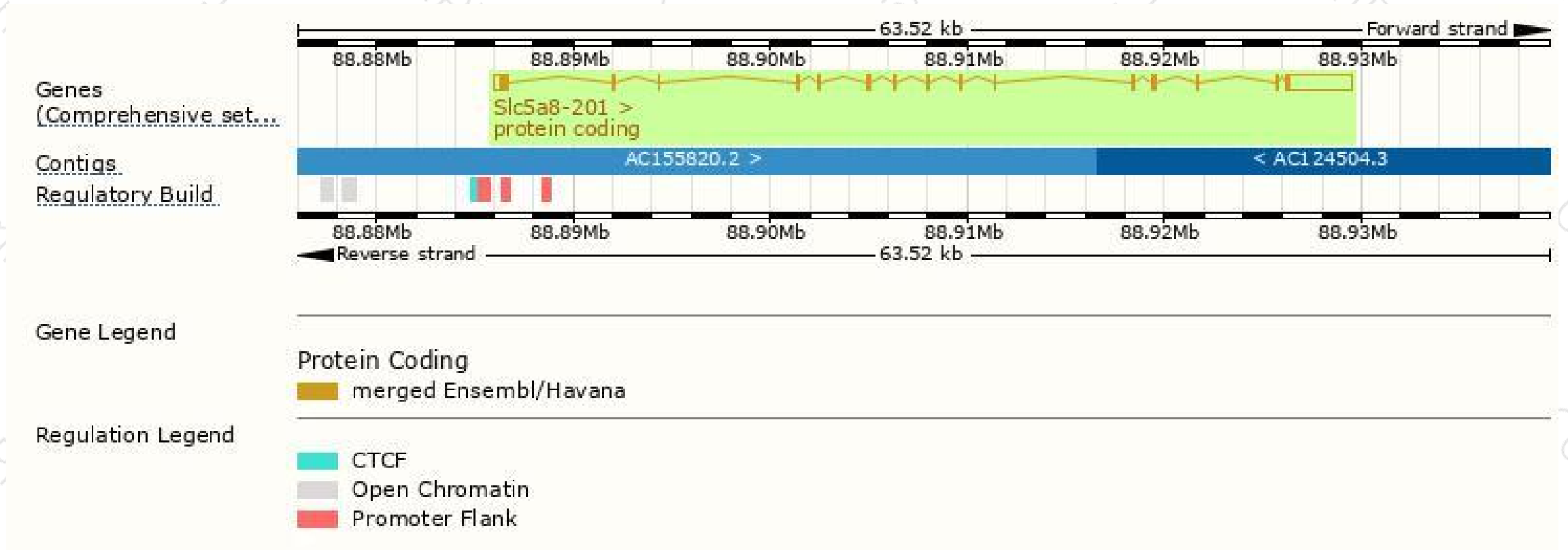
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Slc5a8-201	ENSMUST00000020255.7	5331	611aa	Protein coding	CCDS36026	Q8BYF6	TSL:1 GENCODE basic APPRIS P1

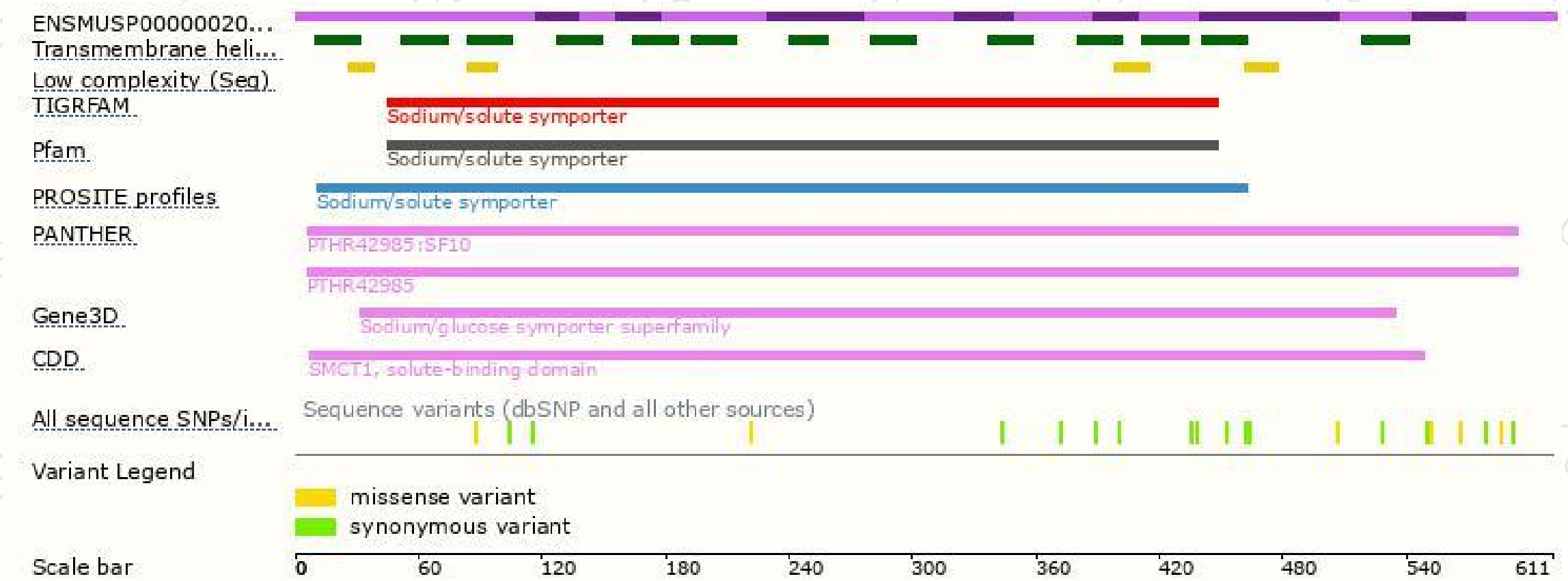
The strategy is based on the design of *Slc5a8-201* transcript, The transcription is shown below



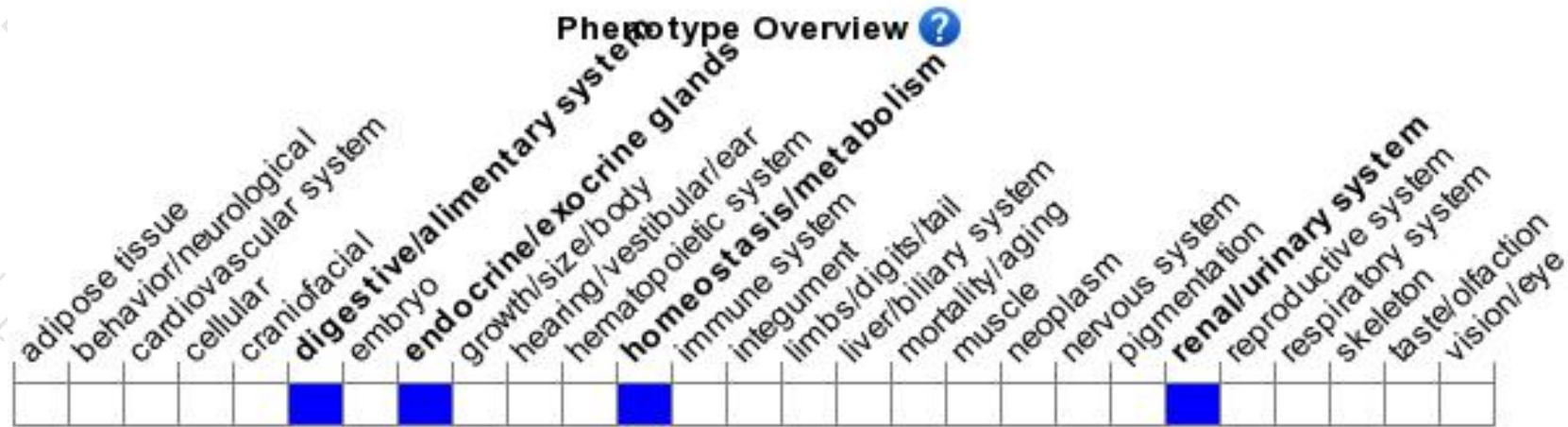
Genomic location distribution



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 null allele exhibit increased lactate concentrations in the saliva and urine.

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

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