

Slc9a5 Cas9-KO Strategy

Designer:

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Project Overview

Project Name

Slc9a5

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Slc9a5* gene. The schematic diagram is as follows:



- The *Slc9a5* gene has 2 transcripts. According to the structure of *Slc9a5* gene, exon2-exon12 of *Slc9a5-201* (ENSMUST00000073149.6) transcript is recommended as the knockout region. The region contains 1655bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Slc9a5* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Slc9a5* gene is located on the Chr8. 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)

Slc9a5 solute carrier family 9 (sodium/hydrogen exchanger), member 5 [Mus musculus (house mouse)]

Gene ID: 277973, updated on 31-Jan-2019

Summary



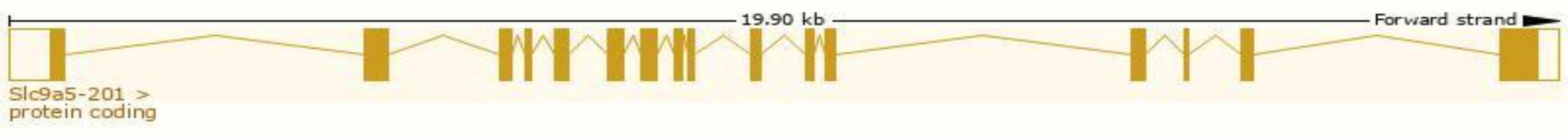
Official Symbol	Slc9a5 provided by MGI
Official Full Name	solute carrier family 9 (sodium/hydrogen exchanger), member 5 provided by MGI
Primary source	MGI:MGI:2685542
See related	Ensembl:ENSMUSG00000014786
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Gm696
Expression	Broad expression in CNS E18 (RPKM 13.3), limb E14.5 (RPKM 12.5) and 21 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

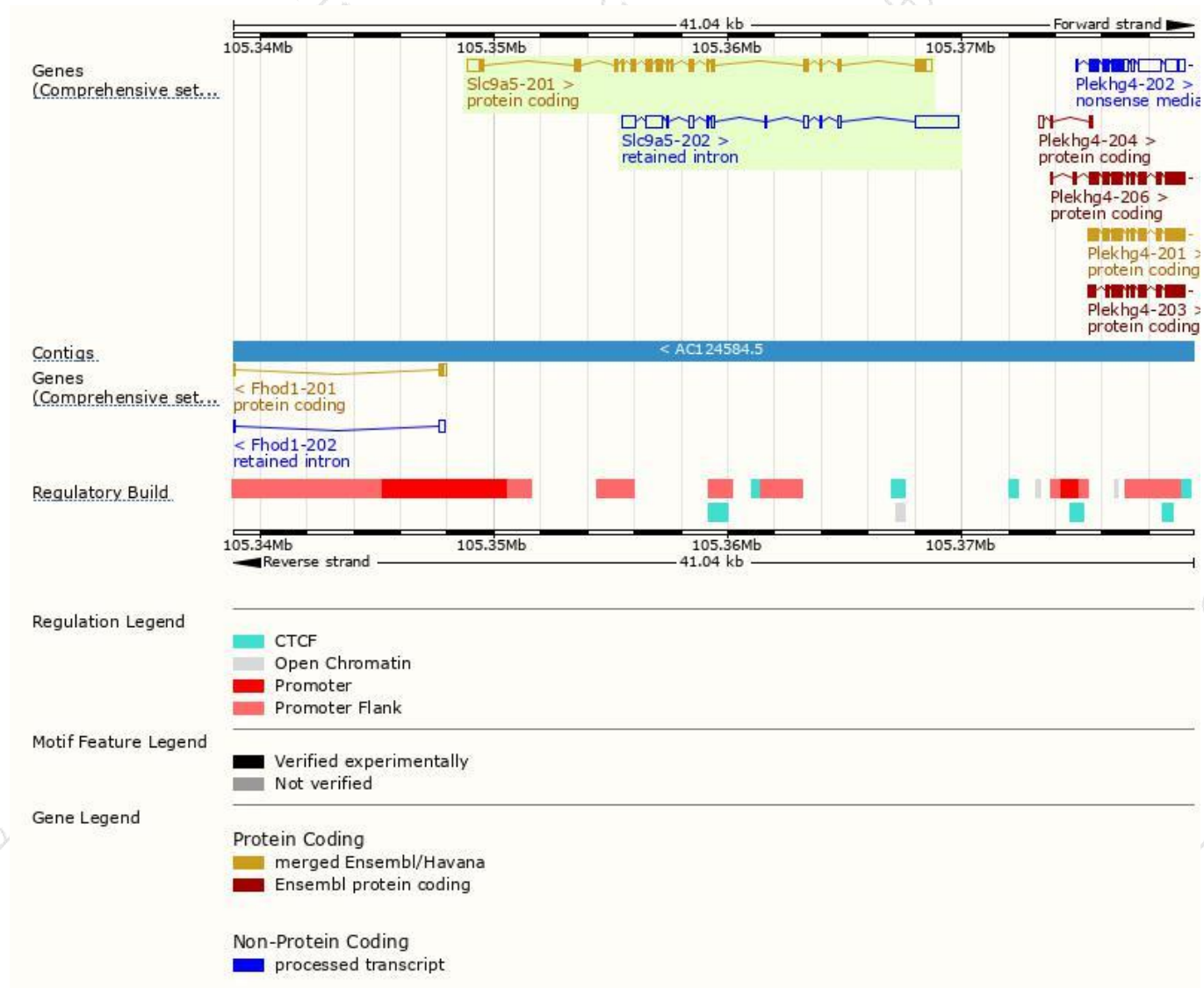
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Slc9a5-201	ENSMUST00000073149.6	3499	898aa	Protein coding	CCDS40457	B2RXE2	TSL:1 GENCODE basic APPRIS P1
Slc9a5-202	ENSMUST00000212772.1	3972	No protein	Retained intron	-	-	TSL:1

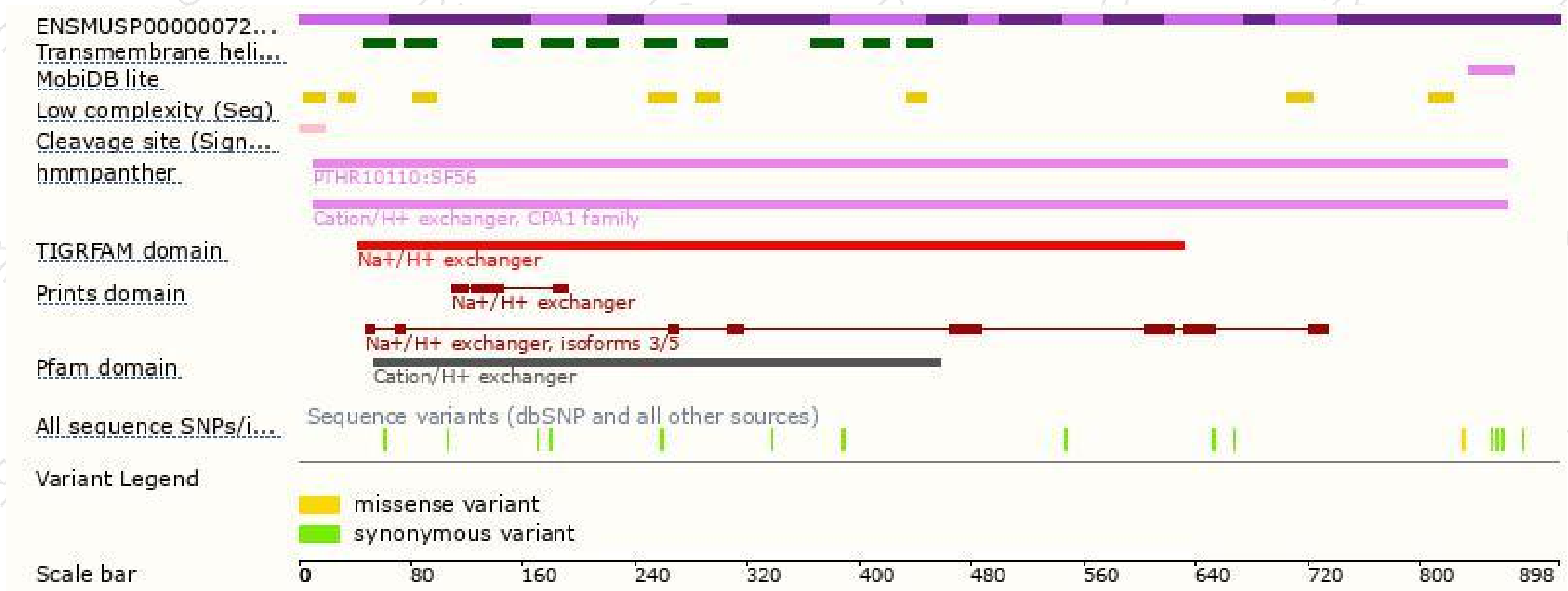
The strategy is based on the design of *Slc9a5-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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