

Slc2a13 Cas9-KO Strategy

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Project Overview

Project Name

Slc2a13

Project type

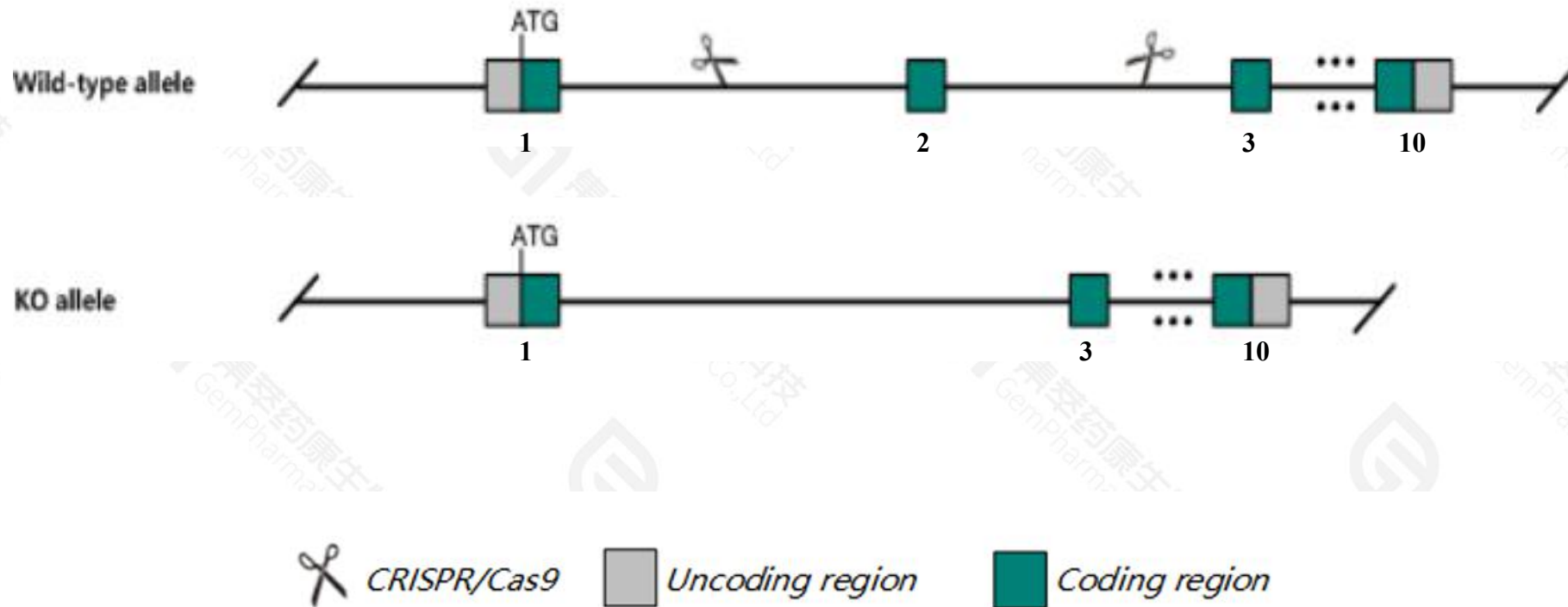
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Slc2a13* gene. The schematic diagram is as follows:



- The *Slc2a13* gene has 3 transcripts. According to the structure of *Slc2a13* gene, exon2 of *Slc2a13*-202(ENSMUST00000109283.2) transcript is recommended as the knockout region. The region contains 160bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Slc2a13* gene. The brief process is as follows: CRISPR/Cas9 system were microinjected into the fertilized eggs of C57BL/6JGpt mice. Fertilized eggs were transplanted to obtain positive F0 mice which were confirmed by PCR and sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6JGpt mice.

- The *Slc2a13* 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.

Slc2a13 solute carrier family 2 (facilitated glucose transporter), member 13 [Mus musculus (house mouse)]

Gene ID: 239606, updated on 25-Sep-2020

Summary



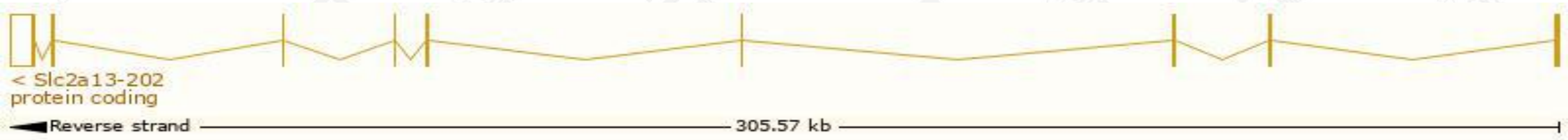
Official Symbol	Slc2a13 provided by MGI
Official Full Name	solute carrier family 2 (facilitated glucose transporter), member 13 provided by MGI
Primary source	MGI:MGI:2146030
See related	Ensembl:ENSMUSG00000036298
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	6530403A04, A630029G22Rik, AI505012, Gm308
Expression	Broad expression in cortex adult (RPKM 10.3), frontal lobe adult (RPKM 7.6) and 15 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

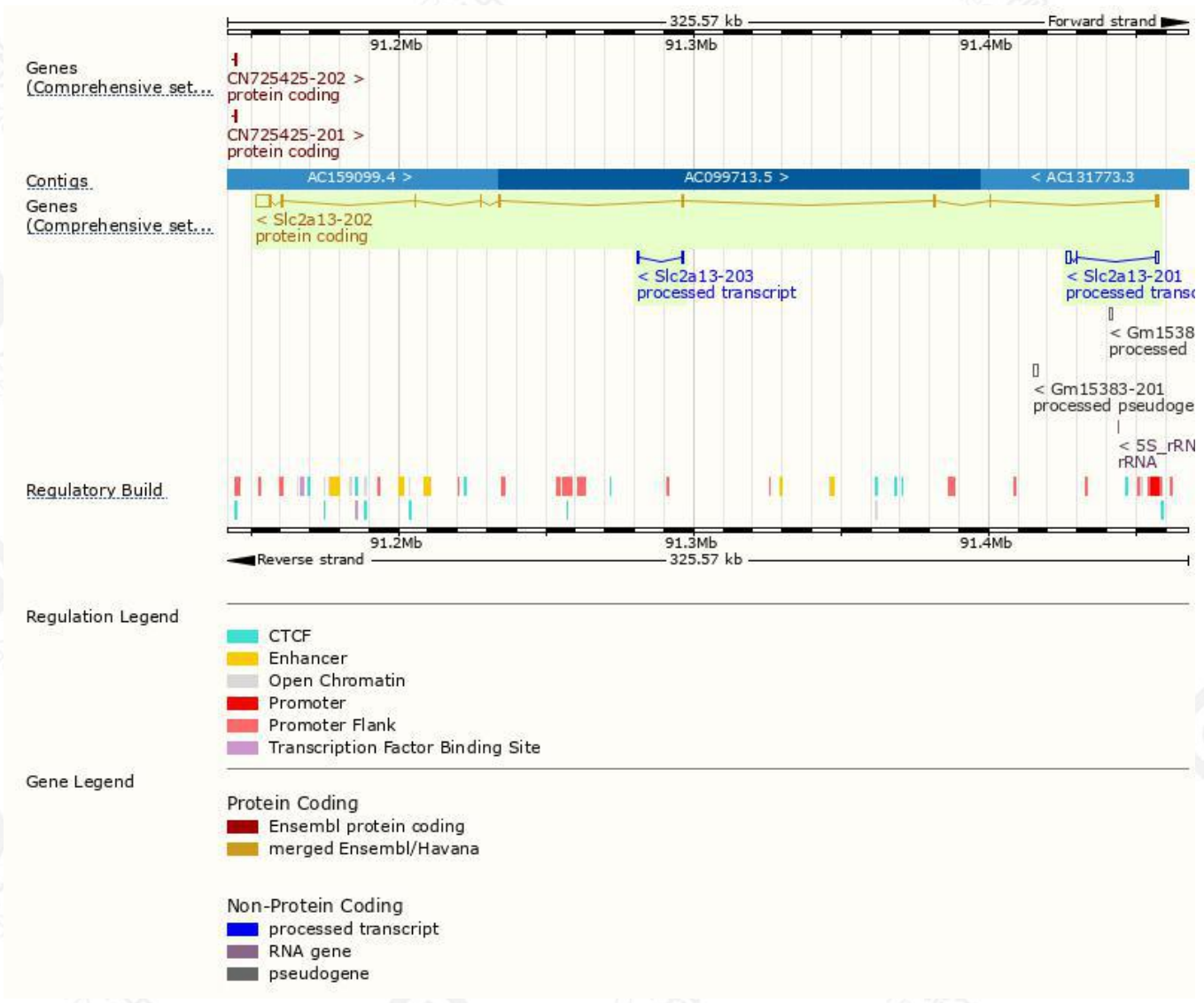
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Slc2a13-202	ENSMUST00000109283.2	6536	637aa	Protein coding	CCDS27761		TSL:1 , GENCODE basic , APPRIS P1 ,
Slc2a13-201	ENSMUST00000109282.2	2081	No protein	Processed transcript	-		TSL:1 ,
Slc2a13-203	ENSMUST00000156971.2	479	No protein	Processed transcript	-		TSL:5 ,

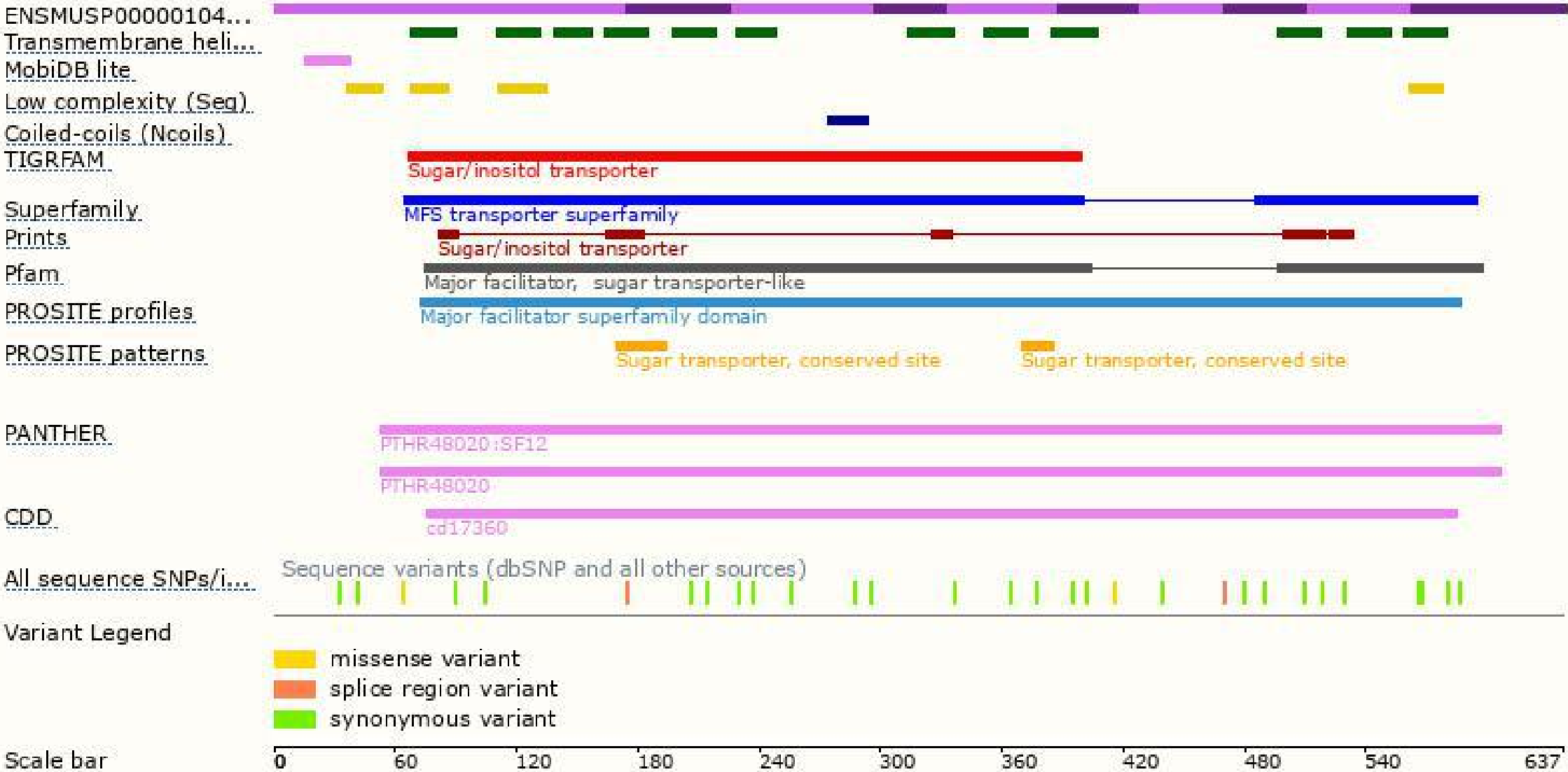
The strategy is based on the design of *Slc2a13-202* transcript,the transcription is shown below:



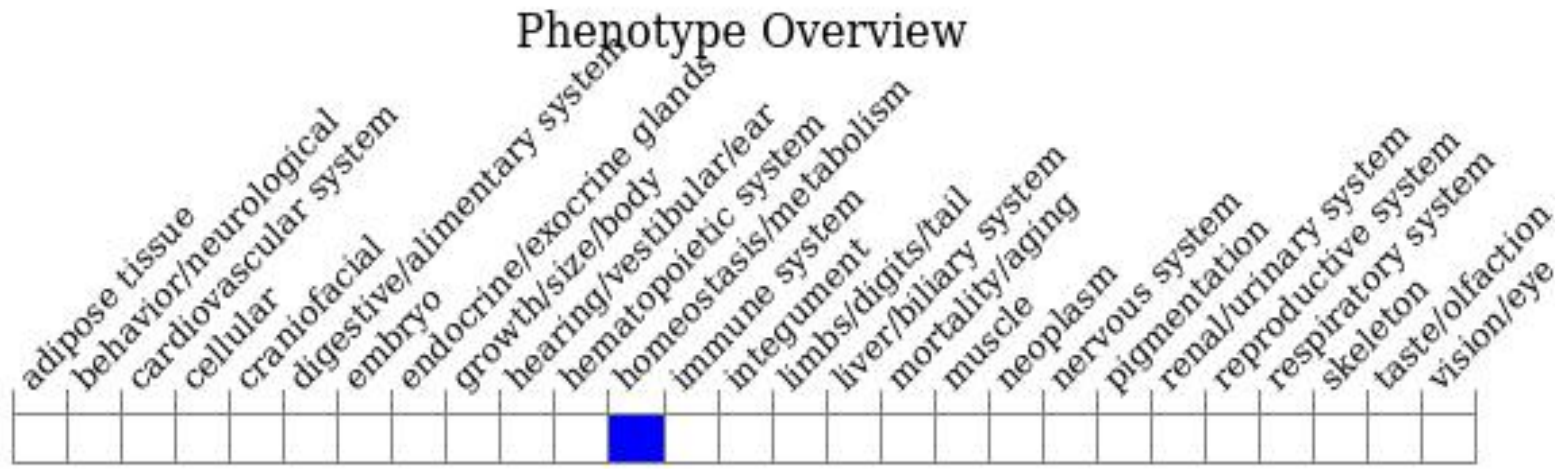
Genomic location distribution



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/>).

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