

# Slc2a10 Cas9-KO Strategy

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# **Project Overview**



**Project Name** 

Slc2a10

**Project type** 

Cas9-KO

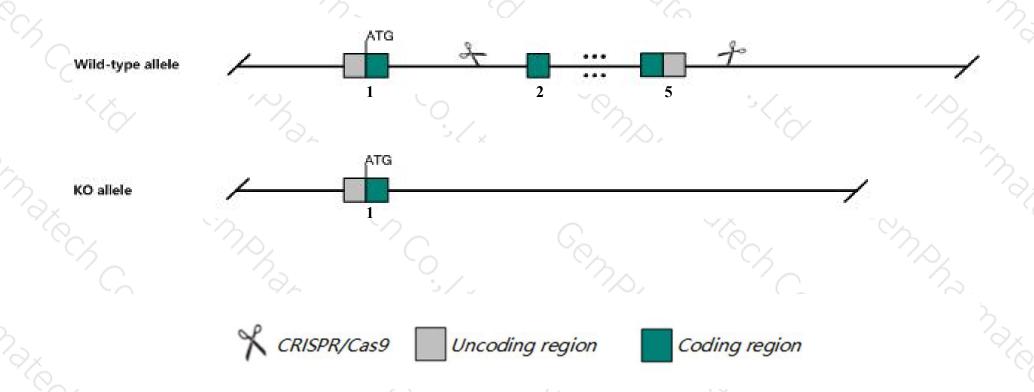
Strain background

C57BL/6JGpt

# **Knockout strategy**



This model will use CRISPR/Cas9 technology to edit the Slc2a10 gene. The schematic diagram is as follows:



### **Technical routes**



- The Slc2a10 gene has 2 transcripts. According to the structure of Slc2a10 gene, exon2-exon5 of Slc2a10-201 (ENSMUST00000029196.4) transcript is recommended as the knockout region. The region contains most of the coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify Slc2a10 gene. The brief process is as follows: CRISPR/Cas9 system w

### **Notice**



- ➤ According to the existing MGI data, Mice carrying ENU-induced mutations in this gene display thickening and aberrant vessel wall shape of large and medium size arteries, with significantly increased elastic fiber number and size. Cerebral arteries appear normal with no evidence of tortuosity, stenosis/dilatation or aneurysm.
- The *Slc2a10* gene is located on the Chr2. 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)



#### Slc2a10 solute carrier family 2 (facilitated glucose transporter), member 10 [ Mus musculus (house mouse) ]

Gene ID: 170441, updated on 21-Jan-2020

#### Summary

☆ ?

Official Symbol Slc2a10 provided by MGI

Official Full Name solute carrier family 2 (facilitated glucose transporter), member 10 provided by MGI

Primary source MGI:MGI:2156687

See related Ensembl: ENSMUSG00000027661

Gene type protein coding
RefSeq status REVIEWED
Organism Mus musculus

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;

Muroidea; Muridae; Murinae; Mus; Mus

Also known as Glut10; AA450473

Summary This gene encodes a class III facilitative glucose transporter. Mutations in the related gene in human are associated with arterial tortuosity

syndrome. [provided by RefSeq, Dec 2013]

Expression Biased expression in stomach adult (RPKM 17.5), colon adult (RPKM 9.3) and 13 other tissues See more

Orthologs human all

#### Genomic context



**Location:** 2 H3; 2 85.66 cM

See Slc2a10 in Genome Data Viewer

Exon count: 5

Annotation release	Status	Assembly	Chr	Location	
108	current	GRCm38.p6 (GCF_000001635.26) 2 NC_000068.7 (165503897165519		NC_000068.7 (165503897165519917)	
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	2	NC_000068.6 (165329478165345411)	

# Transcript information (Ensembl)



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

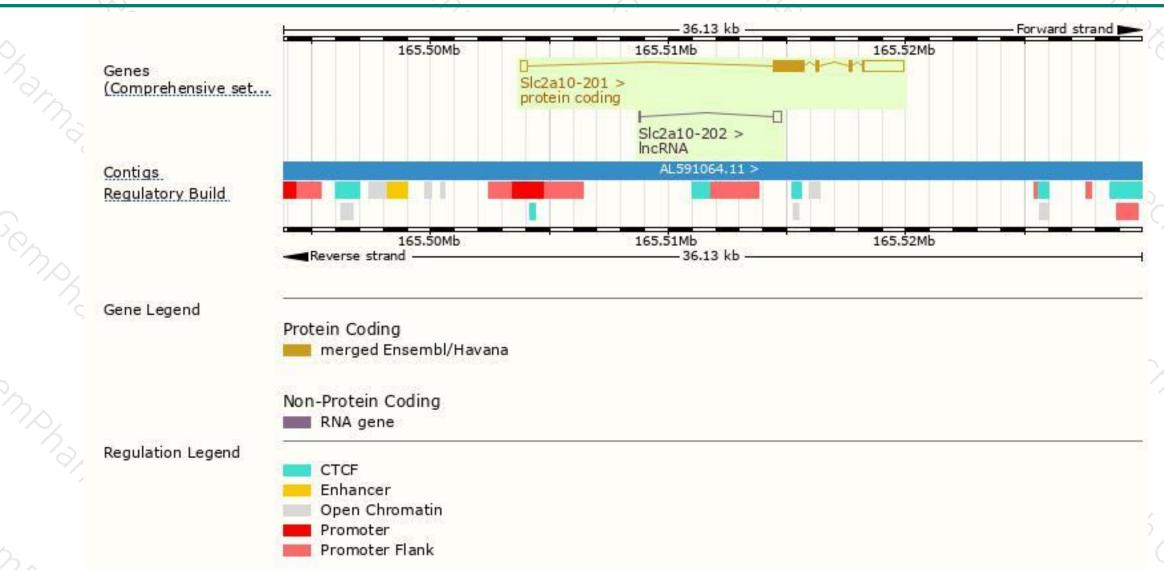
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
SIc2a10-201	ENSMUST00000029196.4	3544	536aa	Protein coding	CCDS17083	A2A4V1 Q8VHD6	TSL:1 GENCODE basic APPRIS P1
SIc2a10-202	ENSMUST00000148463.1	378	No protein	IncRNA	-	393	TSL:2

The strategy is based on the design of Slc2a10-201 transcript, The transcription is shown below



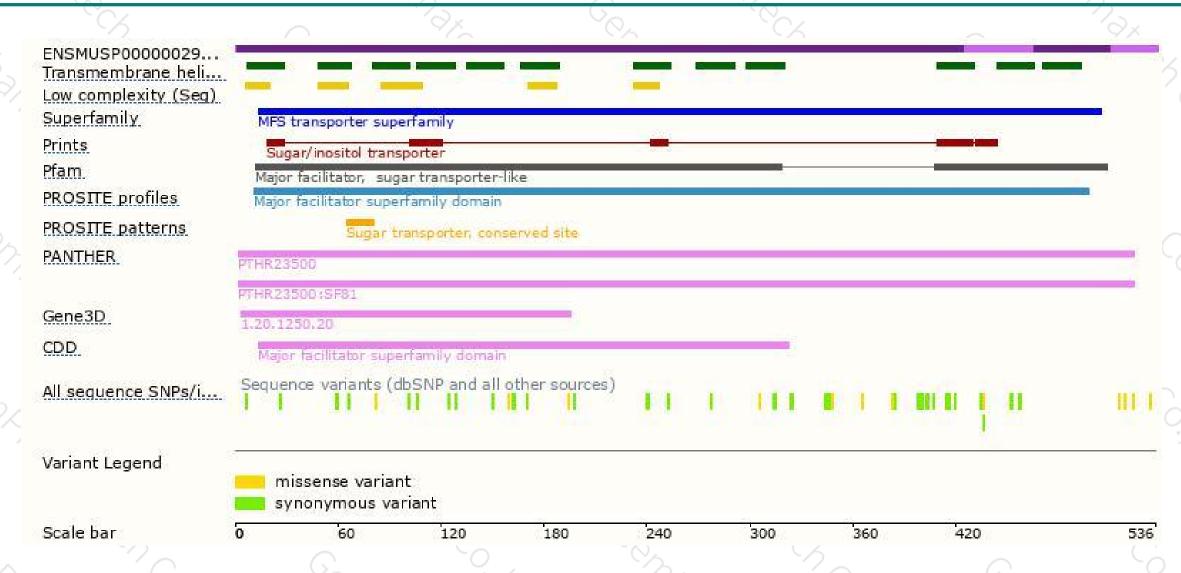
### Genomic location distribution





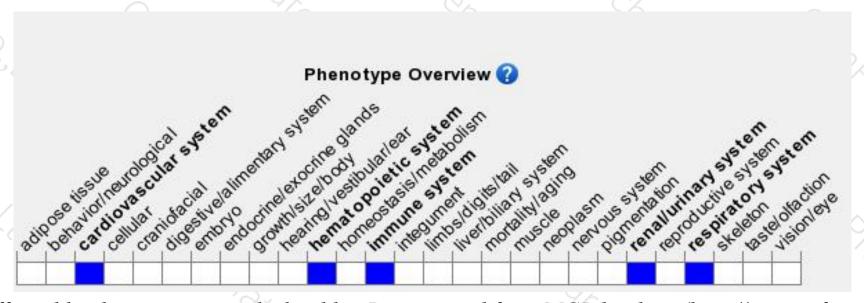
### Protein domain





# 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 carrying ENU-induced mutations in this gene display thickening and aberrant vessel wall shape of large and medium size arteries, with significantly increased elastic fiber number and size.

Cerebral arteries appear normal with no evidence of tortuosity, stenosis/dilatation or aneurysm.



If you have any questions, you are welcome to inquire. Tel: 400-9660890





