

# Slc6a6 Cas9-KO Strategy

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**Reviewer:** 

**Design Date:** 2019-11-19

## **Project Overview**



**Project Name** 

Slc6a6

**Project type** 

Cas9-KO

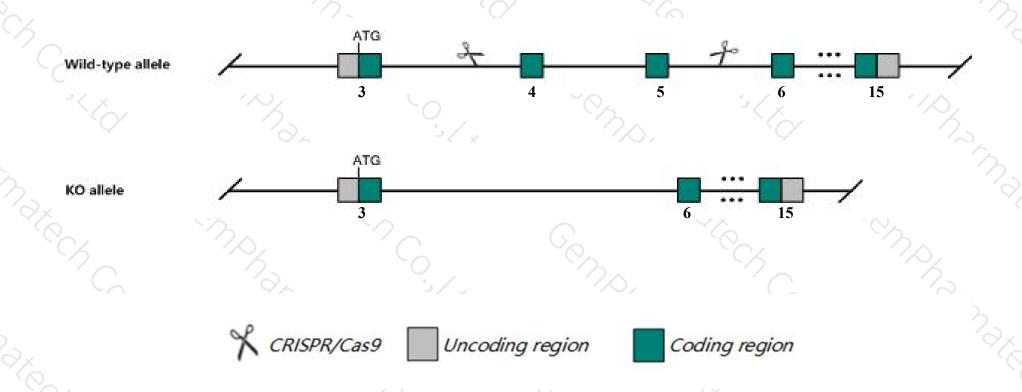
Strain background

C57BL/6JGpt

# **Knockout strategy**



This model will use CRISPR/Cas9 technology to edit the *Slc6a6* gene. The schematic diagram is as follows:



### **Technical routes**



- ➤ The *Slc6a6* gene has 11 transcripts. According to the structure of *Slc6a6* gene, exon4-exon5 of *Slc6a6-201*(ENSMUST00000032185.8) transcript is recommended as the knockout region. The region contains 370bp coding sequence.

  Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify Slc6a6 gene. The brief process is as follows: CRISPR/Cas9 system

### **Notice**



- ➤ According to the existing MGI data, Homozygous mutant mice have impaired vision associated with retinal degeneration. In addition to the visual defects, mutant mice exhibit reduced female fertility and decreased levels of taurine in a variety of tissues.
- > The *Slc6a6* gene is located on the Chr6. 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)



#### SIc6a6 solute carrier family 6 (neurotransmitter transporter, taurine), member 6 [ Mus musculus (house mouse) ]

Gene ID: 21366, updated on 10-Oct-2019



☆?

Official Symbol Slc6a6 provided by MGI

Official Full Name solute carrier family 6 (neurotransmitter transporter, taurine), member 6 provided by MGI

Primary source MGI:MGI:98488

See related Ensembl: ENSMUSG00000030096

Gene type protein coding
RefSeq status VALIDATED
Organism Mus musculus

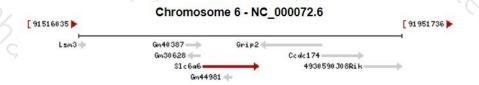
Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;

Muroidea; Muridae; Murinae; Mus; Mus

Also known as Taut; C80501; AA589629

Expression Ubiquitous expression in adrenal adult (RPKM 68.2), kidney adult (RPKM 51.4) and 28 other tissues See more

Orthologs human all



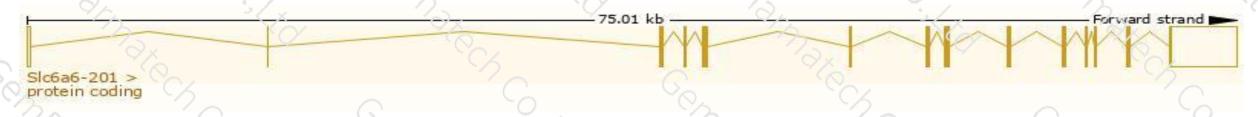
# Transcript information (Ensembl)



The gene has 11 transcripts, all transcripts are shown below:

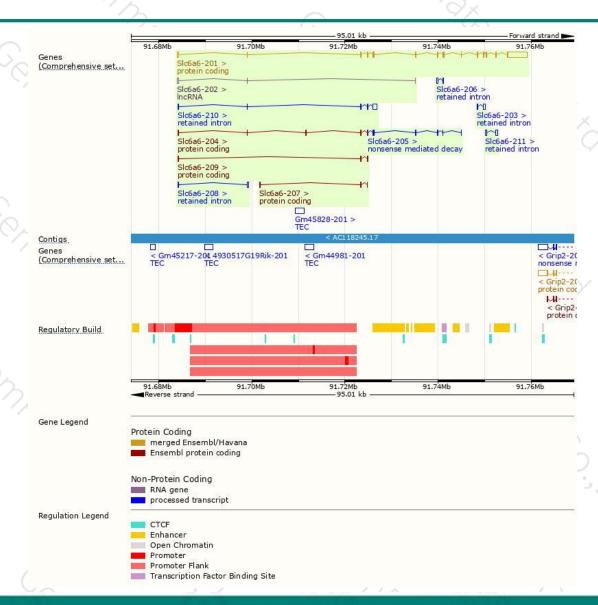
Name 🌲	Transcript ID	bp 🌲	Protein	Translation ID	Biotype	CCDS +	UniProt	Flags	
SIC686-201	ENSMUST00000032185.8	6186	<u>621aa</u>	ENSMUSP00000032185.7	Protein coding	CCDS39570₽	035316@ Q3UPI8@	TSL:1 GENCODE basic	APPRIS P1
Ic6a6-204	ENSMUST00000205480.1	691	110aa	ENSMUSP00000145794.1	Protein coding	2 3	A0A0U1RP20₽	CDS 3' incomplete	TSL:5
SIC6a6-209	ENSMUST00000206545.1	557	<u>121aa</u>	ENSMUSP00000146306.1	Protein coding	973	A0A0U1RQA0⊌	CDS 3' incomplete	TSL:3
SIc6a6-207	ENSMUST00000205828.1	431	83aa	ENSMUSP00000146312.1	Protein coding	- CX	A0A0U1RQA5©	CDS 3' incompleta	TSL:5
Ic6a6-205	ENSMUST00000205663.1	738	49aa	ENSMUSP00000145986.1	Nonsense mediated decay	1 1	A0A0U1RPH7@	CDS 5' incomplete	ISL:5
SIC6a6-210	ENSMUST00000206835.1	1553	No protein		Retained intron	- 4	0	TSL:1	75
Sic6a6-211	ENSMUST00000206988.1	682	No protein	2 30	Retained intron	-	3/30	TSL:3	
SIC6a6-203	ENSMUST00000205477.1	559	No protein	2	Retained intron		``C <del>'</del> 9	/ TSL:3	
SIC6a6-208	ENSMUST00000205764.1	510	No protein	7/2 (8)	Retained intron	75 -	7-	TSL:2	
Ic6a6-208	ENSMUST00000206451.1	481	No protein	72.5	Retained intron	77		TSL:3	
Ic6a6-202	ENSMUST00000205443.1	374	No protein	17/20	IncRNA	2)5	184	TSL:3	b.

The strategy is based on the design of Slc6a6-201 transcript, The transcription is shown below



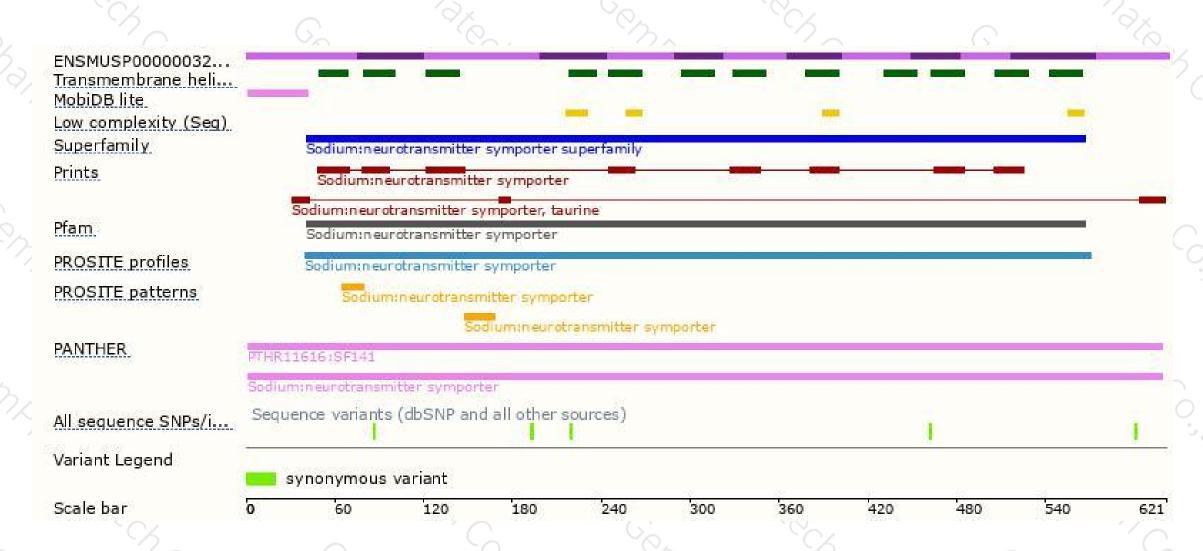
### Genomic location distribution





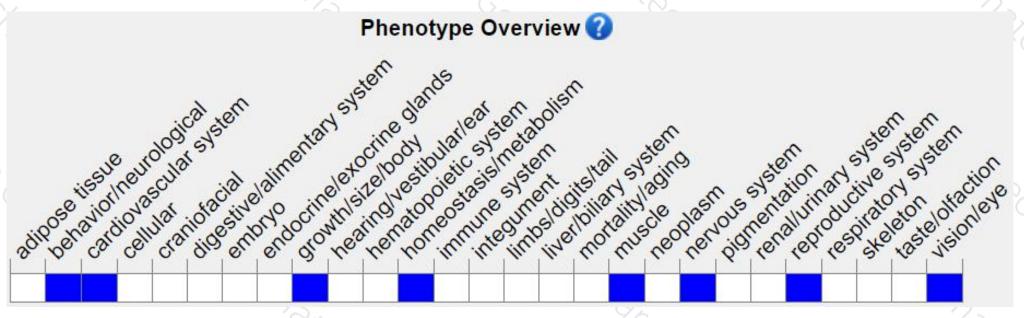
### Protein domain





# 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, Homozygous mutant mice have impaired vision associated with retinal degeneration. In addition to the visual defects, mutant mice exhibit reduced female fertility and decreased levels of taurine in a variety of tissues.



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





