

Glt8d1 Cas9-KO Strategy

Designer: Yanhua Shen

Reviewer: Xueting Zhang

Design Date: 2020-2-15

Project Overview



Project Name

Glt8d1

Project type

Cas9-KO

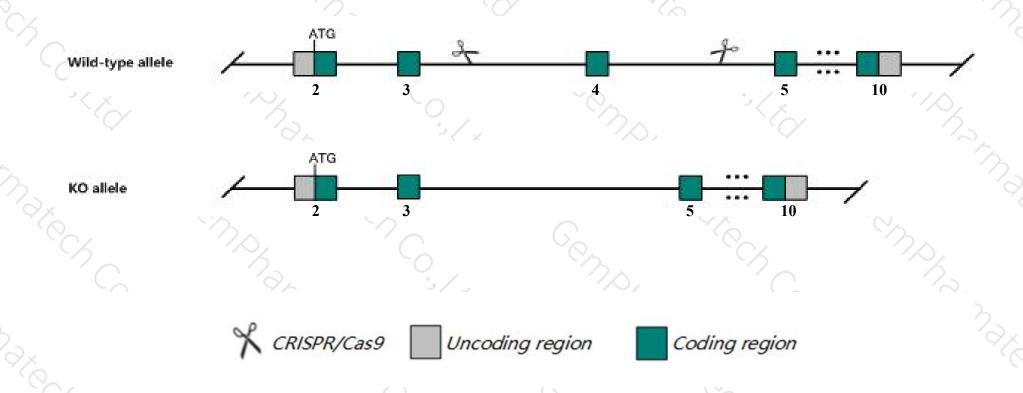
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Glt8d1* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Glt8d1* gene has 10 transcripts. According to the structure of *Glt8d1* gene, exon4 of *Glt8d1-201*(ENSMUST00000022476.8) transcript is recommended as the knockout region. The region contains 214bp coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Glt8d1* gene. The brief process is as follows: CRISPR/Cas9 system

Notice



- > According to the existing MGI data, Mice homozygous for a disruption in this gene display exhibited impaired sensorimotor gating/attention during prepulse inhibition testing.
- ➤ Some amino acids will remain at the N-terminus and some functions may be retained.
- Transcripts 206,207 may not be affected. The effect of transcript 205 is unknown.
- > The *Glt8d1* gene is located on the Chr14. 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)



Glt8d1 glycosyltransferase 8 domain containing 1 [Mus musculus (house mouse)]

Gene ID: 76485, updated on 14-Aug-2019

Summary

↑ ?

Official Symbol Glt8d1 provided by MGI

Official Full Name glycosyltransferase 8 domain containing 1 provided by MGI

Primary source MGI:MGI:1923735

See related Ensembl: ENSMUSG00000021916

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 Al450005; 2410004H05Rik; 5430414N14Rik

Summary This gene encodes a member of the glycosyltransferase family. The encoded protein is a type II transmembrane protein containing a glycosyltransferase 8 domain in the

lumenal (C-terminal) portion. The specific function of this protein has not been determined. Two alternatively spliced variants encoding the same isoform have been

described. [provided by RefSeq, Sep 2009]

Expression Broad expression in testis adult (RPKM 75.2), limb E14.5 (RPKM 20.1) and 21 other tissues See more

Orthologs human all

Genomic context

Location: 14; 14 B See Glt8d1 in Genome Data Viewer

Exon count: 12

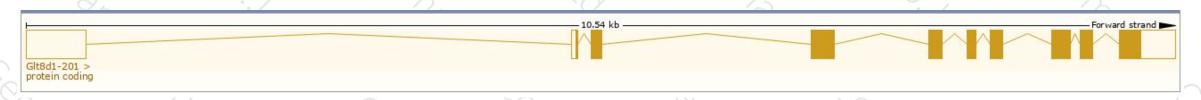
Transcript information (Ensembl)



The gene has 10 transcripts, all transcripts are shown below:

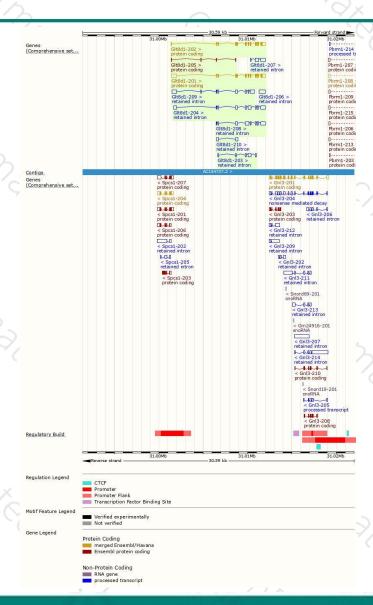
Glt8d1-202 ENSMUST00000168584.8 1498 371aa Protein coding CCDS26904 Q6NSU3 TSL:1 GENCODE basic APPRIS Glt8d1-205 ENSMUST00000226378.1 213 18aa Protein coding - A0A2I3BRI4 CDS 3' incomplete Glt8d1-204 ENSMUST00000226348.1 1923 No protein Retained intron - - Glt8d1-208 ENSMUST00000228755.1 1692 No protein Retained intron - - Glt8d1-209 ENSMUST00000228718.1 1681 No protein Retained intron - - Glt8d1-207 ENSMUST0000022627783.1 955 No protein Retained intron - - Glt8d1-206 ENSMUST00000226971.1 772 No protein Retained intron - - Glt8d1-206 ENSMUST00000226959.1 684 No protein Retained intron - -	Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Glt8d1-205 ENSMUST00000226378.1 213 18aa Protein coding - A0A2I3BRI4 CDS 3' incomplete Glt8d1-204 ENSMUST00000226348.1 1923 No protein Retained intron - - Glt8d1-208 ENSMUST00000228255.1 1692 No protein Retained intron - - Glt8d1-209 ENSMUST00000228718.1 1681 No protein Retained intron - - Glt8d1-207 ENSMUST00000227783.1 955 No protein Retained intron - - Glt8d1-208 ENSMUST00000226271.1 772 No protein Retained intron - - Glt8d1-206 ENSMUST00000226959.1 684 No protein Retained intron - -	Glt8d1-201	ENSMUST00000022476.8	2018	<u>371aa</u>	Protein coding	CCDS26904	Q6NSU3	TSL:1 GENCODE basic APPRIS P1
Git8d1-204 ENSMUST00000226348.1 1923 No protein Retained intron - -	Glt8d1-202	ENSMUST00000168584.8	1498	<u>371aa</u>	Protein coding	CCDS26904	Q6NSU3	TSL:1 GENCODE basic APPRIS P1
Glt8d1-208 ENSMUST00000228255.1 1692 No protein Retained intron - - Glt8d1-209 ENSMUST00000228718.1 1681 No protein Retained intron - - Glt8d1-207 ENSMUST00000227783.1 955 No protein Retained intron - - Glt8d1-203 ENSMUST00000226271.1 772 No protein Retained intron - - Glt8d1-206 ENSMUST00000226959.1 684 No protein Retained intron - -	Glt8d1-205	ENSMUST00000226378.1	213	<u>18aa</u>	Protein coding	120	A0A2I3BRI4	CDS 3' incomplete
Glt8d1-209 ENSMUST00000228718.1 1681 No protein Retained intron - - Glt8d1-207 ENSMUST00000227783.1 955 No protein Retained intron - - Glt8d1-203 ENSMUST00000226271.1 772 No protein Retained intron - - Glt8d1-206 ENSMUST00000226959.1 684 No protein Retained intron - -	Glt8d1-204	ENSMUST00000226348.1	1923	No protein	Retained intron	12/	-	
Glt8d1-207 ENSMUST00000227783.1 955 No protein Retained intron - - Glt8d1-203 ENSMUST00000226271.1 772 No protein Retained intron - - Glt8d1-206 ENSMUST00000226959.1 684 No protein Retained intron - -	GIt8d1-208	ENSMUST00000228255.1	1692	No protein	Retained intron	151	1. st	
Glt8d1-203 <u>ENSMUST00000226271.1</u> 772 No protein <u>Retained intron</u>	Glt8d1-209	ENSMUST00000228718.1	1681	No protein	Retained intron	-	. s .	
GIt8d1-206 ENSMUST00000226959.1 684 No protein Retained intron	GIt8d1-207	ENSMUST00000227783.1	955	No protein	Retained intron	120	11-	
	GIt8d1-203	ENSMUST00000226271.1	772	No protein	Retained intron	12/	-	
Glt8d1-210 ENSMUST00000228914.1 487 No protein Betained intron	Glt8d1-206	ENSMUST00000226959.1	684	No protein	Retained intron	181		
	Glt8d1-210	ENSMUST00000228914.1	487	No protein	Retained intron	-		

The strategy is based on the design of Glt8d1-201 transcript, The transcription is shown below



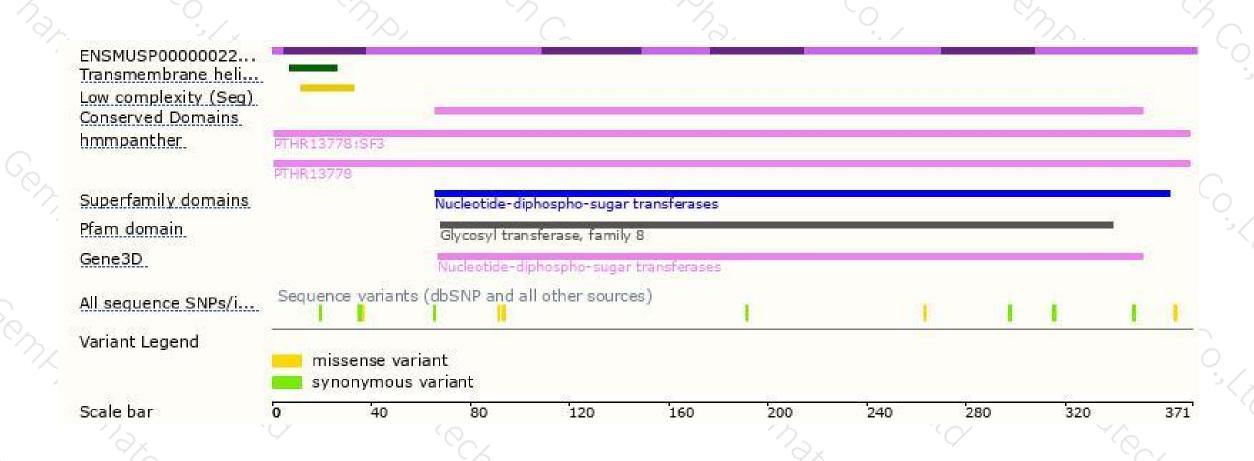
Genomic location distribution





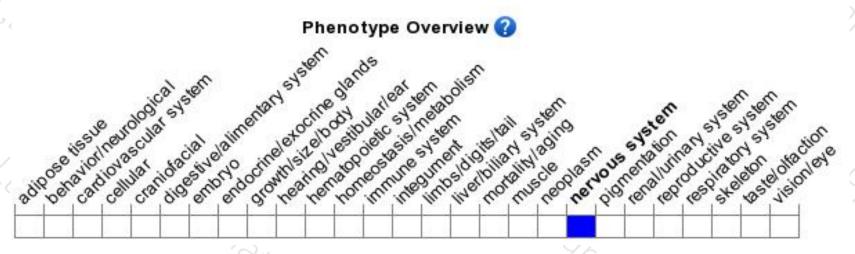
Protein domain





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 disruption in this gene display exhibited impaired sensorimotor gating/attention during prepulse inhibition testing.



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





