

Gtdc1 Cas9-KO Strategy

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Reviewer: Xueting Zhang

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Project Overview



Project Name Gtdc1

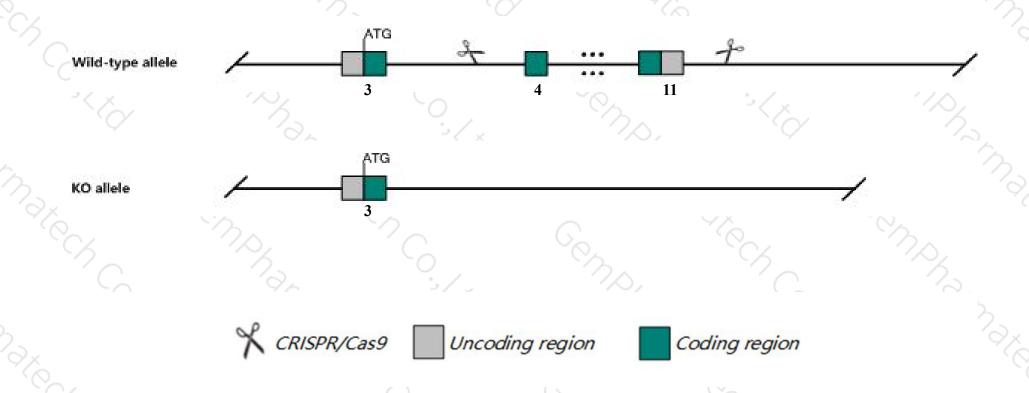
Project type Cas9-KO

Strain background C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Gtdc1* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Gtdc1* gene has 13 transcripts. According to the structure of *Gtdc1* gene, exon4-exon11 of *Gtdc1-203*(ENSMUST00000112810.7) transcript is recommended as the knockout region. The region contains most of coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Gtdc1* gene. The brief process is as follows: CRISPR/Cas9 system v

Notice



- ➤ Transcripts 202,206,209 may not be affected. The effect of transcripts 205,210,211,213 is unknown.
- The *Gtdc1* 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)



Gtdc1 glycosyltransferase-like domain containing 1 [Mus musculus (house mouse)]

Gene ID: 227835, updated on 24-Oct-2019

Summary

☆ ?

Official Symbol Gtdc1 provided by MGI

Official Full Name glycosyltransferase-like domain containing 1 provided by MGI

Primary source MGI:MGI:2444269

See related Ensembl: ENSMUSG00000036890

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 E330008O22Rik

Expression Broad expression in cortex adult (RPKM 8.8), CNS E18 (RPKM 7.0) and 23 other tissues See more

Orthologs human all

Genomic context

Location: 2; 2 B

Exon count: 18

See Gtdc1 in Genome Data Viewer

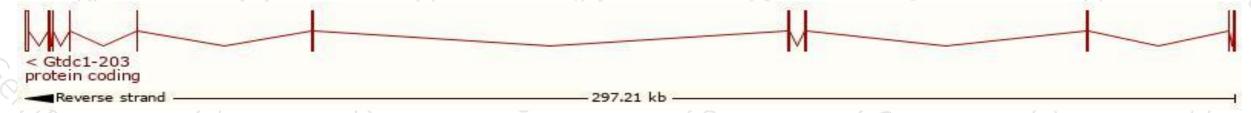
Transcript information (Ensembl)



The gene has 13 transcripts, all transcripts are shown below:

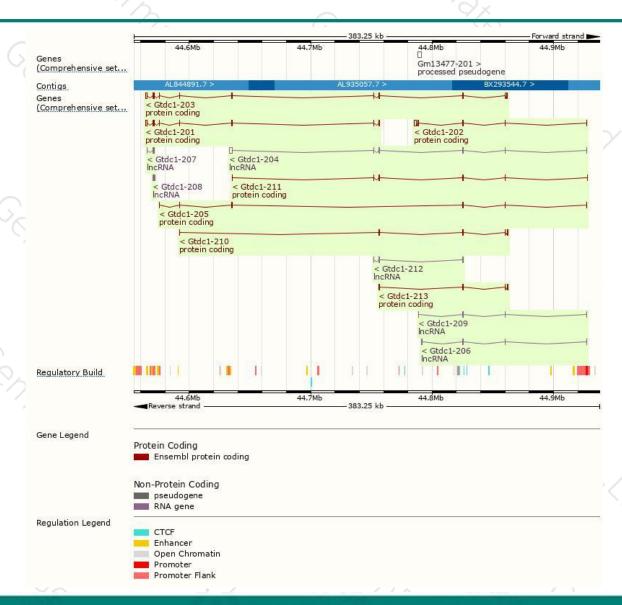
| | | ′ /) . | | | | |
|-----------------------|--|--|--|---|---|--|
| Transcript ID | bp | Protein | Biotype | CCDS | UniProt | Flags |
| ENSMUST00000112810.7 | 2564 | 445aa | Protein coding | CCDS71050 | Q8BW56 | TSL:1 GENCODE basic APPRIS P1 |
| ENSMUST00000100127.8 | 2422 | 106aa | Protein coding | - | Q8BW56 | TSL:1 GENCODE basic |
| ENSMUST00000049051.12 | 2197 | 386aa | Protein coding | 17 E | G8JL42 | TSL:1 GENCODE basic |
| ENSMUST00000130991.7 | 784 | 238aa | Protein coding | 62 | B1AY87 | CDS 3' incomplete TSL:5 |
| ENSMUST00000148279.7 | 677 | <u>191aa</u> | Protein coding | 1.5 m | A0A0A0MQK2 | CDS 3' incomplete TSL:5 |
| ENSMUST00000146694.7 | 652 | <u>142aa</u> | Protein coding | - | B1AY88 | CDS 3' incomplete TSL:3 |
| ENSMUST00000154744.7 | 571 | <u>115aa</u> | Protein coding | () | A0A0A0MQI8 | CDS 3' incomplete TSL:2 |
| ENSMUST00000129240.7 | 2882 | No protein | IncRNA | 62 | 2 | TSL:1 |
| ENSMUST00000133442.1 | 646 | No protein | IncRNA | 1.5 | - | TSL:5 |
| ENSMUST00000143333.1 | 643 | No protein | IncRNA | | - | TSL:2 |
| ENSMUST00000143766.7 | 638 | No protein | IncRNA | 1/2 1/2 | 2 | TSL:3 |
| ENSMUST00000134813.1 | 629 | No protein | IncRNA | 12 | 2 | TSL:1 |
| ENSMUST00000148786.1 | 547 | No protein | IncRNA | 15 | 5. | TSL:2 |
| | ENSMUST00000112810.7 ENSMUST00000100127.8 ENSMUST00000049051.12 ENSMUST00000130991.7 ENSMUST00000148279.7 ENSMUST00000146694.7 ENSMUST00000154744.7 ENSMUST00000154744.7 ENSMUST00000133442.1 ENSMUST00000143333.1 ENSMUST00000143766.7 ENSMUST00000134813.1 | ENSMUST00000112810.7 2564 ENSMUST00000100127.8 2422 ENSMUST00000049051.12 2197 ENSMUST00000130991.7 784 ENSMUST00000148279.7 677 ENSMUST00000146694.7 652 ENSMUST00000154744.7 571 ENSMUST00000129240.7 2882 ENSMUST00000133442.1 646 ENSMUST00000143333.1 643 ENSMUST00000143766.7 638 ENSMUST00000134813.1 629 | ENSMUST00000112810.7 2564 445aa ENSMUST00000100127.8 2422 106aa ENSMUST00000049051.12 2197 386aa ENSMUST00000130991.7 784 238aa ENSMUST00000148279.7 677 191aa ENSMUST00000146694.7 652 142aa ENSMUST00000154744.7 571 115aa ENSMUST00000129240.7 2882 No protein ENSMUST00000133442.1 646 No protein ENSMUST00000143333.1 643 No protein ENSMUST00000143766.7 638 No protein ENSMUST00000134813.1 629 No protein | ENSMUST00000112810.7 2564 445aa Protein coding ENSMUST00000100127.8 2422 106aa Protein coding ENSMUST00000049051.12 2197 386aa Protein coding ENSMUST00000130991.7 784 238aa Protein coding ENSMUST00000148279.7 677 191aa Protein coding ENSMUST00000146694.7 652 142aa Protein coding ENSMUST00000154744.7 571 115aa Protein coding ENSMUST00000129240.7 2882 No protein IncRNA ENSMUST00000133442.1 646 No protein IncRNA ENSMUST00000143333.1 643 No protein IncRNA ENSMUST00000143766.7 638 No protein IncRNA ENSMUST00000134813.1 629 No protein IncRNA | ENSMUST00000112810.7 2564 445aa Protein coding CCDS71050 ENSMUST00000100127.8 2422 106aa Protein coding - ENSMUST00000049051.12 2197 386aa Protein coding - ENSMUST00000130991.7 784 238aa Protein coding - ENSMUST00000148279.7 677 191aa Protein coding - ENSMUST00000146694.7 652 142aa Protein coding - ENSMUST00000154744.7 571 115aa Protein coding - ENSMUST00000129240.7 2882 No protein IncRNA - ENSMUST00000133442.1 646 No protein IncRNA - ENSMUST00000143333.1 643 No protein IncRNA - ENSMUST00000143766.7 638 No protein IncRNA - ENSMUST00000134813.1 629 No protein IncRNA - | ENSMUST00000112810.7 2564 445aa Protein coding CCDS71050 Q8BW56 ENSMUST00000100127.8 2422 106aa Protein coding - Q8BW56 ENSMUST00000049051.12 2197 386aa Protein coding - G8JL42 ENSMUST00000130991.7 784 238aa Protein coding - B1AY87 ENSMUST00000148279.7 677 191aa Protein coding - A0A0A0MQK2 ENSMUST00000146694.7 652 142aa Protein coding - B1AY88 ENSMUST00000154744.7 571 115aa Protein coding - A0A0A0MQI8 ENSMUST00000129240.7 2882 No protein IncRNA - - ENSMUST00000133442.1 646 No protein IncRNA - - ENSMUST000001433333.1 643 No protein IncRNA - - ENSMUST00000134813.1 629 No protein IncRNA - - |

The strategy is based on the design of Gtdc1-203 transcript, The transcription is shown below



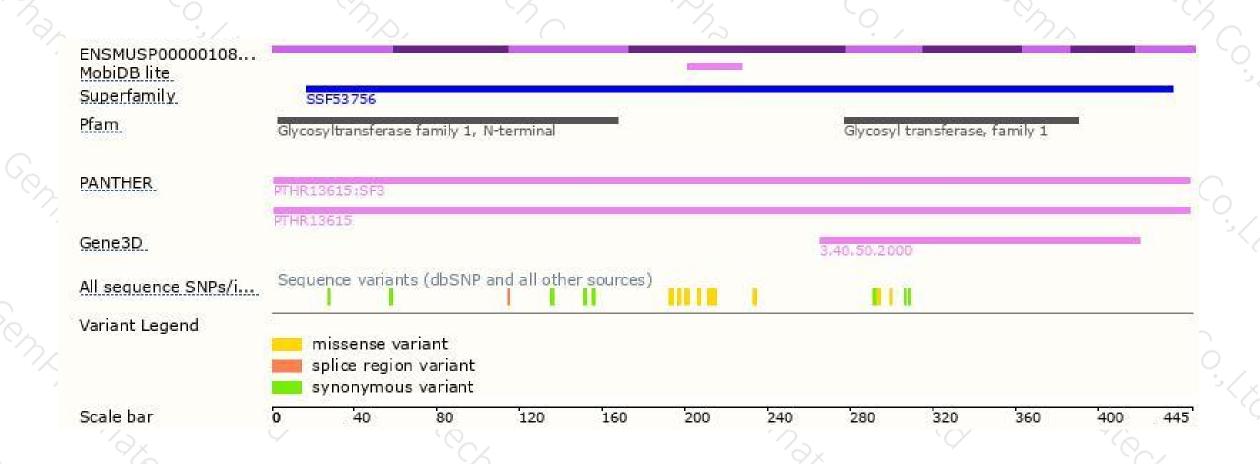
Genomic location distribution





Protein domain







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





