

***Clstn2* Cas9-KO Strategy**

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Project Overview

Project Name

Clstn2

Project type

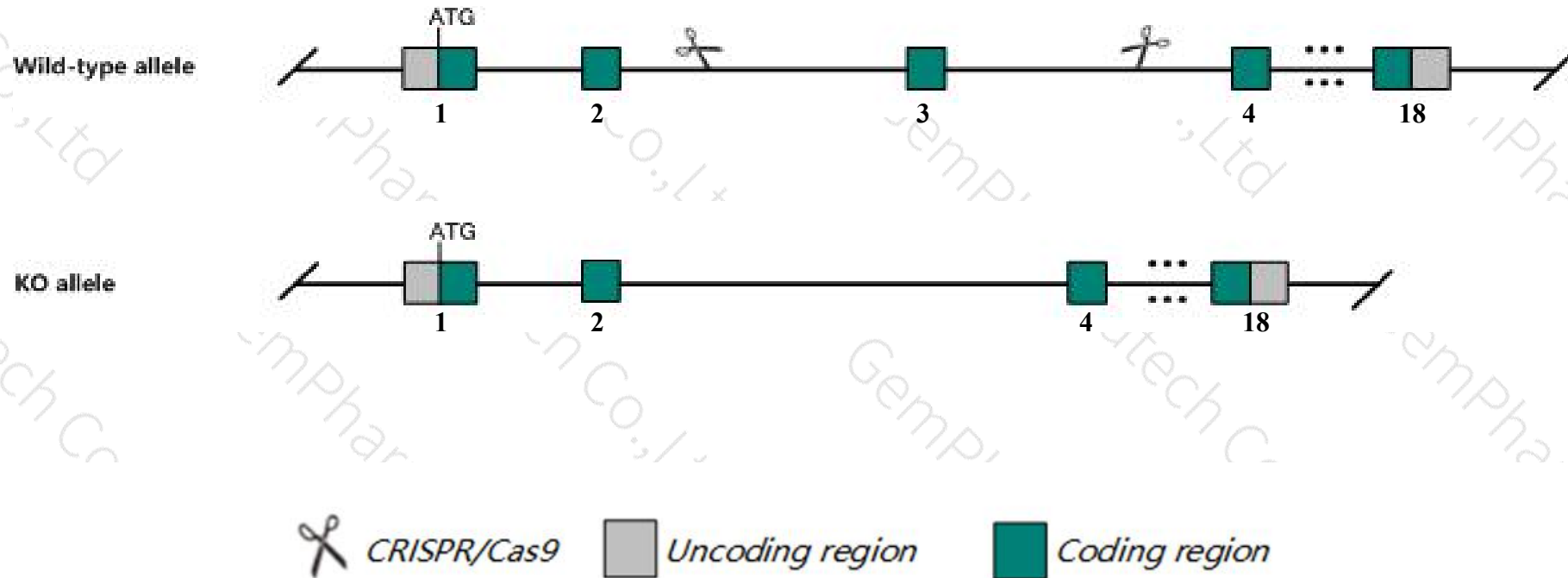
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Clstn2* gene. The schematic diagram is as follows:



- The *Clstn2* gene has 2 transcripts. According to the structure of *Clstn2* gene, exon3 of *Clstn2-201* (ENSMUST00000035027.12) transcript is recommended as the knockout region. The region contains 196bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Clstn2* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Homozygous KO mice display deficiency in spatial learning and memory in Morris water and Barnes maze tasks and increased locomotor activity in open field test.
- The *Clstn2* gene is located on the Chr9. 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)

Clstn2 calsyntenin 2 [*Mus musculus* (house mouse)]

Gene ID: 64085, updated on 11-Sep-2019

Summary

- Official Symbol** Clstn2 provided by [MGI](#)
- Official Full Name** calsyntenin 2 provided by [MGI](#)
- Primary source** [MGI:MGI:1929897](#)
- See related** [Ensembl:ENSMUSG00000032452](#)
- Gene type** protein coding
- RefSeq status** PROVISIONAL
- Organism** [Mus musculus](#)
- Lineage** Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
- Also known as** Cs2; CSTN2; Cst-2; AI448973; mKIAA4134; 2900042C18Rik
- Expression** Biased expression in cortex adult (RPKM 7.4), CNS E18 (RPKM 6.7) and 10 other tissues [See more](#)
- Orthologs** [human](#) [all](#)

Genomic context

Location: 9; 9 E3.3

See Clstn2 in [Genome Data Viewer](#)

Exon count: 17

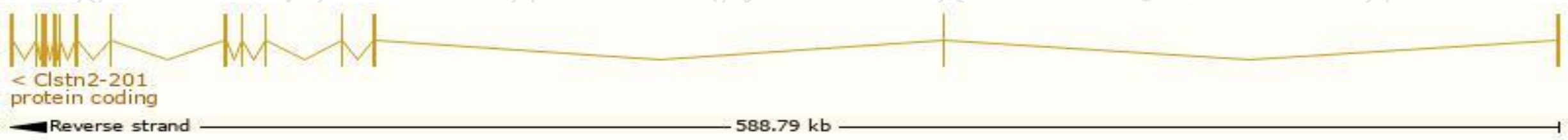
Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	9	NC_000075.6 (97444395..98033373, complement)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	9	NC_000075.5 (97344814..97933586, complement)

Transcript information (Ensembl)

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

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Clstn2-201	ENSMUST00000035027.12	4365	966aa	Protein coding	CCDS40733	Q9ER65	TSL:1 GENCODE basic APPRIS P2
Clstn2-202	ENSMUST00000162295.1	4174	960aa	Protein coding	-	Q9ER65	TSL:5 GENCODE basic APPRIS ALT2

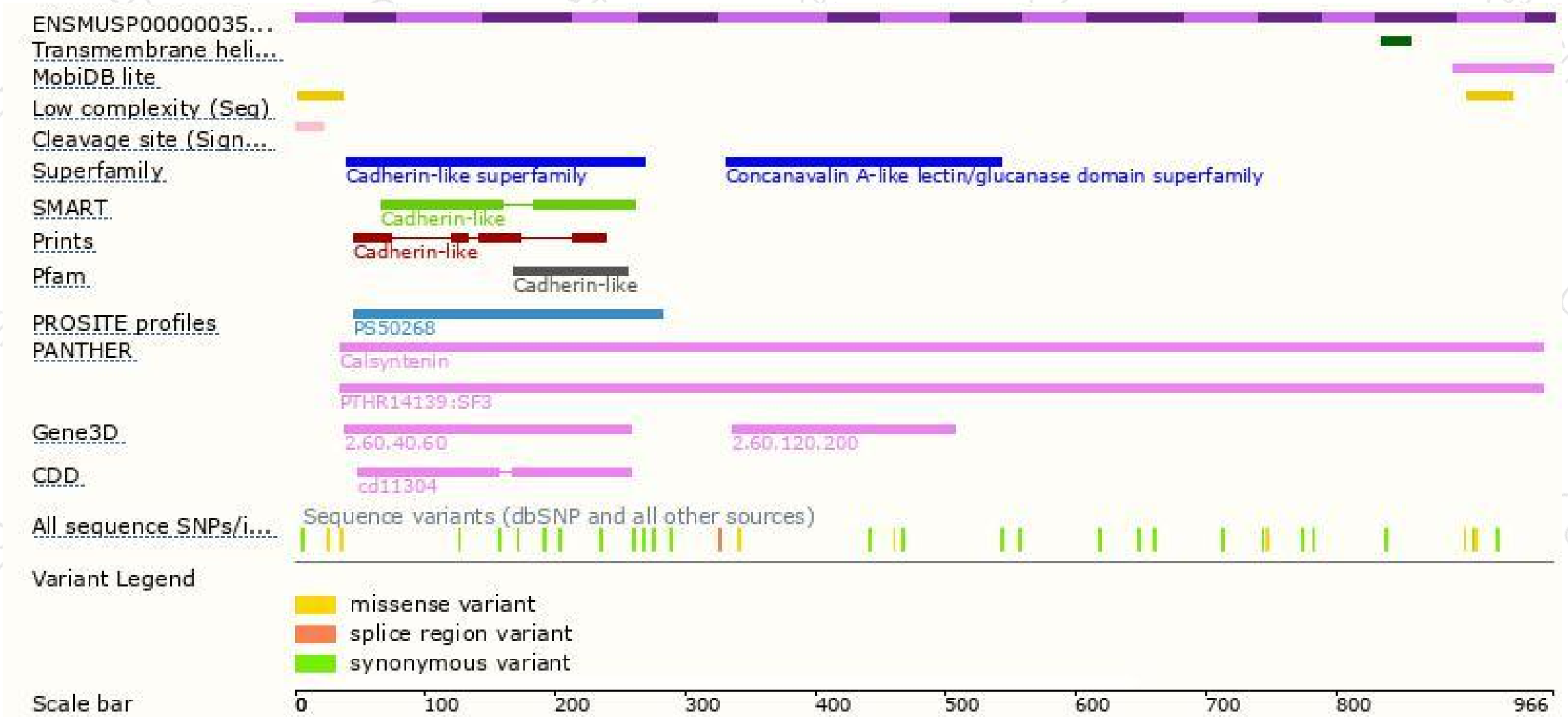
The strategy is based on the design of *Clstn2-201* transcript,The transcription is shown below



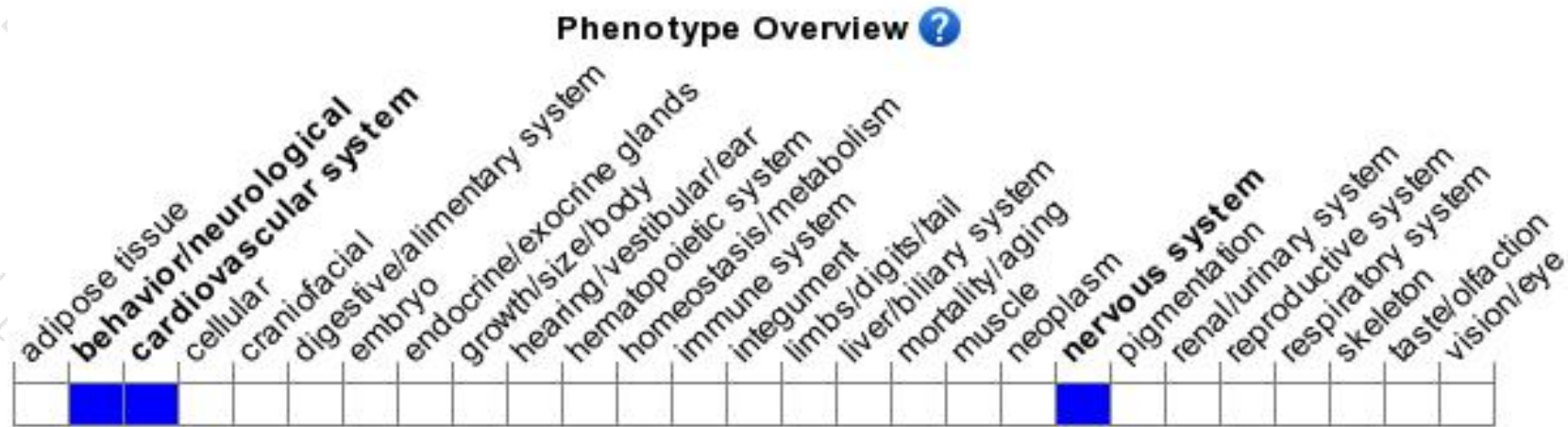
Genomic location distribution



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 KO mice display deficiency in spatial learning and memory in Morris water and Barnes maze tasks and increased locomotor activity in open field test.

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

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