

Chmp2a Cas9-KO Strategy

Designer: Wenjing Li

Reviewer: Jiayuan Yao

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Project Overview



Project Name

Chmp2a

Project type

Cas9-KO

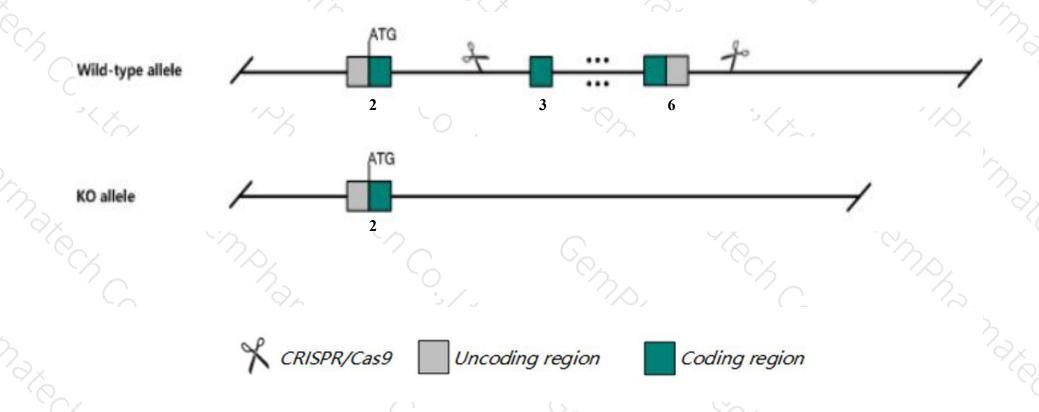
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Chmp2a* gene. The schematic diagram is as follows:



Technical routes



- The *Chmp2a* gene has 6 transcripts. According to the structure of *Chmp2a* gene, exon3-exon6 of *Chmp2a-201*(ENSMUST00000005711.5) 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 *Chmp2a* gene. The brief process is as follows: CRISPR/Cas9 system were microinjected into the fertilized eggs of C57BL/6JGpt mice. Fertilized eggs were transplanted to obtain positive F0 mice which were confirmed by PCR and sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6JGpt mice.

Notice



- > The N-terminal of Chmp2a gene will remain several amino acids, it may remain the partial function of Chmp2a gene.
- The floxed region is near to the C-terminal of *Trim28* gene, this strategy may influence the regulatory function of the C-terminal of *Trim28* gene.
- The *Chmp2a* gene is located on the Chr7. 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)



Chmp2a charged multivesicular body protein 2A [Mus musculus (house mouse)]

Gene ID: 68953, updated on 13-Mar-2020





Official Symbol Chmp2a provided by MGI

Official Full Name charged multivesicular body protein 2A provided by MGI

Primary source MGI:MGI:1916203

See related Ensembl: ENSMUSG000000033916

Gene type protein coding

RefSeq status VALIDATED

Organism <u>Mus musculus</u>

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

Muroidea; Muridae; Murinae; Mus; Mus

Also known as 1500016L11Rik, mVps2

Expression Ubiquitous expression in placenta adult (RPKM 67.0), bladder adult (RPKM 65.4) and 28 other tissuesSee more

Orthologs <u>human</u> <u>all</u>

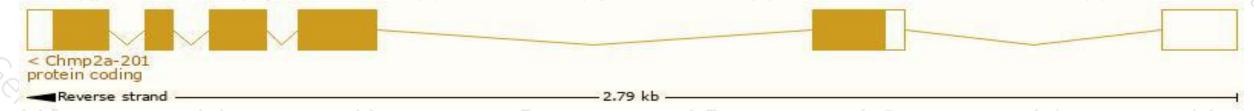
Transcript information (Ensembl)



The gene has 6 transcripts, all transcripts are shown below:

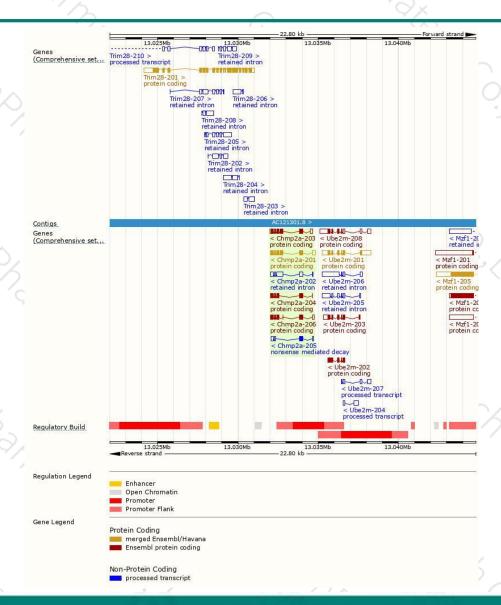
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags	
Chmp2a-201	ENSMUST00000005711.5	949	222aa	Protein coding	CCDS20824	Q9DB34	TSL:1 GENCODE basic APPRIS P1	
Chmp2a-203	ENSMUST00000210587.1	930	222aa	Protein coding	CCDS20824	Q9DB34	TSL:3 GENCODE basic APPRIS P1	
Chmp2a-206	ENSMUST00000211626.1	726	<u>172aa</u>	Protein coding	120	A0A1B0GR63	TSL:2 GENCODE basic	
Chmp2a-204	ENSMUST00000211344.1	606	162aa	Protein coding		A0A1B0GT75	TSL:5 GENCODE basic	
Chmp2a-205	ENSMUST00000211369.1	521	<u>56aa</u>	Nonsense mediated decay	848	A0A1B0GSH4	TSL:3	
Chmp2a-202	ENSMUST00000209487.1	888	No protein	Retained intron	853	40	TSL:5	

The strategy is based on the design of *Chmp2a-201* transcript, the transcription is shown below:



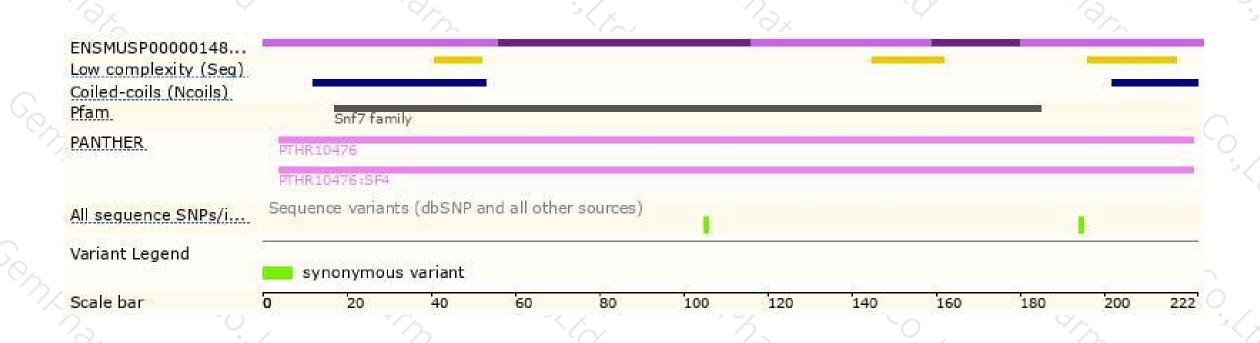
Genomic location distribution





Protein domain







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





