

Adgra3 Cas9-CKO Strategy

Designer:

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Design Date:

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Project Overview



Project Name

Adgra3

Project type

Cas9-CKO

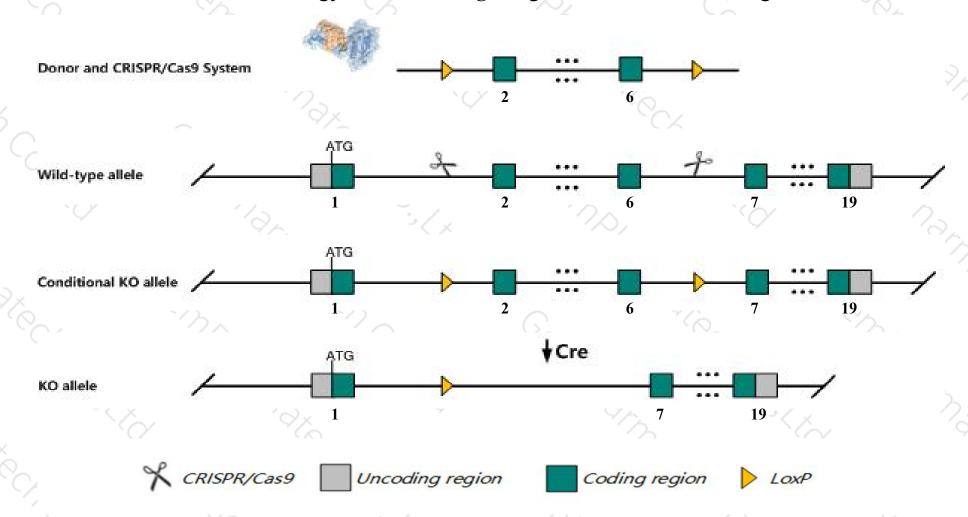
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Adgra3* gene. The schematic diagram is as follows:



Technical routes



- The *Adgra3* gene has 7 transcripts. According to the structure of *Adgra3* gene, exon2-exon6 of *Adgra3-201* (ENSMUST00000030971.6) transcript is recommended as the knockout region. The region contains 449bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Adgra3* gene. The brief process is as follows:CRISPR/Cas9 system and Donor 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.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

Notice



- > According to the existing MGI data, Homozygous mutant mice are fertile and grossly normal.
- The *Adgra3* gene is located on the Chr5. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)



Adgra3 adhesion G protein-coupled receptor A3 [Mus musculus (house mouse)]

Gene ID: 70693, updated on 31-Jan-2019

Summary

☆ ?

Official Symbol Adgra3 provided by MGI

Official Full Name adhesion G protein-coupled receptor A3 provided by MGI

Primary source MGI:MGI:1917943

See related Ensembl:ENSMUSG00000029090

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 3830613O22Rik, AU044632, Gpr125, Tem5-like

Expression Ubiquitous expression in limb E14.5 (RPKM 32.5), ovary adult (RPKM 24.3) and 27 other tissuesSee more

Orthologs human all

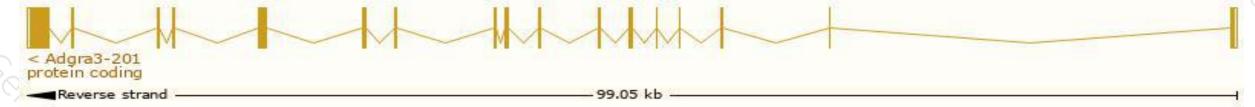
Transcript information (Ensembl)



The gene has 7 transcripts, all transcripts are shown below:

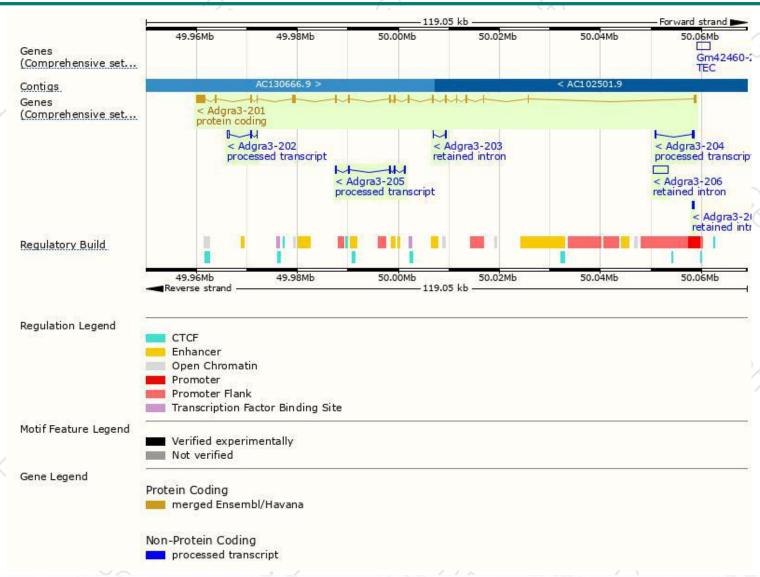
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Adgra3-201	ENSMUST00000030971.6	4480	<u>1310aa</u>	Protein coding	CCDS51500	Q7TT36	TSL:1 GENCODE basic APPRIS P1
Adgra3-205	ENSMUST00000198818.1	797	No protein	Processed transcript	÷	19 5	TSL:3
Adgra3-202	ENSMUST00000196177.1	543	No protein	Processed transcript	ě.	ÿ <u>L</u>	TSL:5
Adgra3-204	ENSMUST00000196915.4	396	No protein	Processed transcript		62	TSL:3
Adgra3-206	ENSMUST00000198868.1	2991	No protein	Retained intron	-	6.7	TSL:NA
Adgra3-207	ENSMUST00000199132.1	314	No protein	Retained intron	-	. 8 5	TSL:3
Adgra3-203	ENSMUST00000196229.1	215	No protein	Retained intron	-	102	TSL:1

The strategy is based on the design of Adgra 3-201 transcript, The transcription is shown below



Genomic location distribution





Protein domain







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





