

Adgrl4 Cas9-CKO Strategy

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Project Overview



Project Name Adgrl4

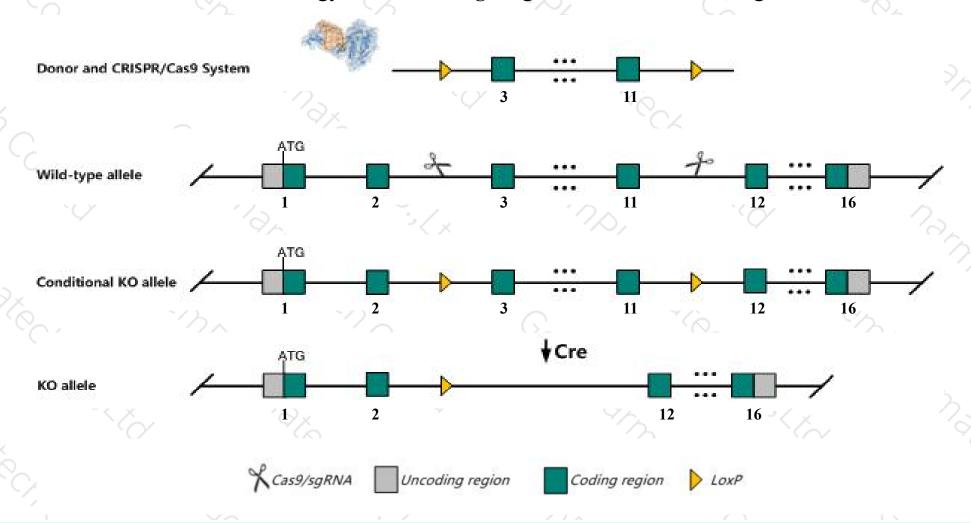
Project type Cas9-CKO

Strain background C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Adgrl4 gene. The schematic diagram is as follows:



Technical routes



- The *Adgrl4* gene has 6 transcripts. According to the structure of *Adgrl4* gene, exon3-exon11 of *Adgrl4-201* (ENSMUST00000046977.11) transcript is recommended as the knockout region. The region contains 1436bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Adgrl4* gene. The brief process is as follows:sgRNA was transcribed in vitro, donor vector was constructed.Cas9, sgRNA 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 was 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, For a targeted mutation, no significant differences were detected between homozygous mice and controls in a high-throughput screen.
- > The *Adgrl4* gene is located on the Chr3. 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)



Adgrl4 adhesion G protein-coupled receptor L4 [Mus musculus (house mouse)]

Gene ID: 170757, updated on 31-Jan-2019

Summary

↑ ?

Official Symbol Adgrl4 provided by MGI

Official Full Name adhesion G protein-coupled receptor L4 provided by MGI

Primary source MGI:MGI:2655562

See related Ensembl:ENSMUSG00000039167

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 1110033N21Rik, ETL1, Eltd1, Etl

Expression Broad expression in lung adult (RPKM 19.1), subcutaneous fat pad adult (RPKM 9.6) and 19 other tissuesSee more

Orthologs <u>human</u> all

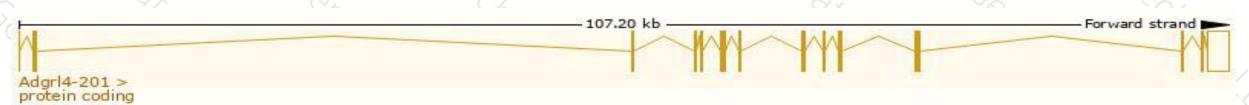
Transcript information (Ensembl)



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

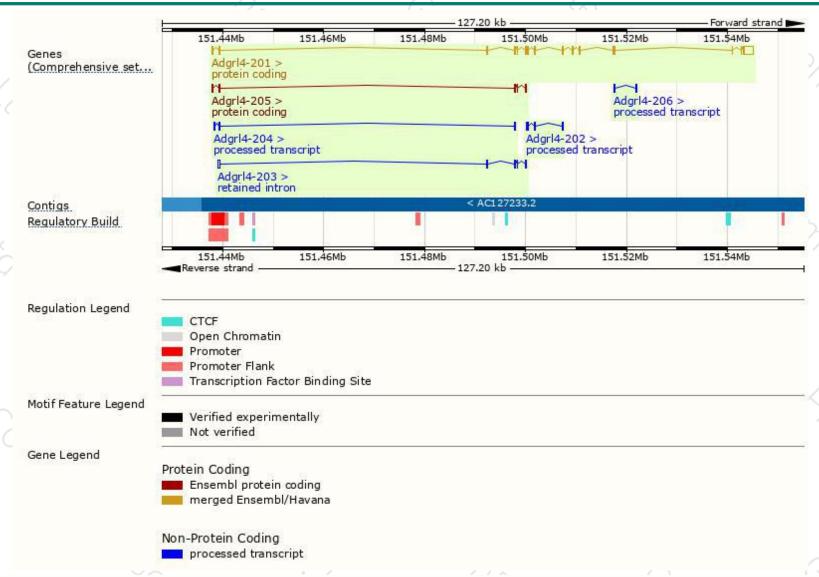
| | | | | <u> </u> | | l kun |
|-----------------------|---|--|---|--|--|--|
| Transcript ID | bp | Protein | Biotype | CCDS | UniProt | Flags |
| ENSMUST00000046977.11 | 4106 | 739aa | Protein coding | CCDS17910 | Q923X1 | TSL:1 GENCODE basic APPRIS P1 |
| ENSMUST00000196970.1 | 598 | <u>181aa</u> | Protein coding | * | A0A0G2JGY1 | CDS 3' incomplete TSL:3 |
| ENSMUST00000129283.1 | 625 | No protein | Processed transcript | - | 10 | TSL:3 |
| ENSMUST00000198893.1 | 498 | No protein | Processed transcript | 24 | 20 | TSL:3 |
| ENSMUST00000155652.1 | 345 | No protein | Processed transcript | - | | TSL:3 |
| ENSMUST00000141038.1 | 783 | No protein | Retained intron | -8 | . . 8 | TSL:2 |
| | ENSMUST00000196977.11 ENSMUST00000196970.1 ENSMUST00000129283.1 ENSMUST00000198893.1 ENSMUST00000155652.1 | ENSMUST000000196970.1 4106 ENSMUST00000196970.1 598 ENSMUST00000129283.1 625 ENSMUST00000198893.1 498 ENSMUST00000155652.1 345 | ENSMUST000000196970.1 4106 739aa ENSMUST00000196970.1 598 181aa ENSMUST00000129283.1 625 No protein ENSMUST00000198893.1 498 No protein ENSMUST00000155652.1 345 No protein | ENSMUST00000046977.11 4106 739aa Protein coding ENSMUST00000196970.1 598 181aa Protein coding ENSMUST00000129283.1 625 No protein Processed transcript ENSMUST00000198893.1 498 No protein Processed transcript ENSMUST00000155652.1 345 No protein Processed transcript | ENSMUST00000046977.11 4106 739aa Protein coding CCDS17910 ENSMUST00000196970.1 598 181aa Protein coding - ENSMUST00000129283.1 625 No protein Processed transcript - ENSMUST00000198893.1 498 No protein Processed transcript - ENSMUST00000155652.1 345 No protein Processed transcript - | ENSMUST00000046977.11 4106 739aa Protein coding CCDS17910 Q923X1 ENSMUST00000196970.1 598 181aa Protein coding - A0A0G2JGY1 ENSMUST00000129283.1 625 No protein Processed transcript - - ENSMUST00000198893.1 498 No protein Processed transcript - - ENSMUST00000155652.1 345 No protein Processed transcript - - |

The strategy is based on the design of Adgrl4-201 transcript, The transcription is shown below



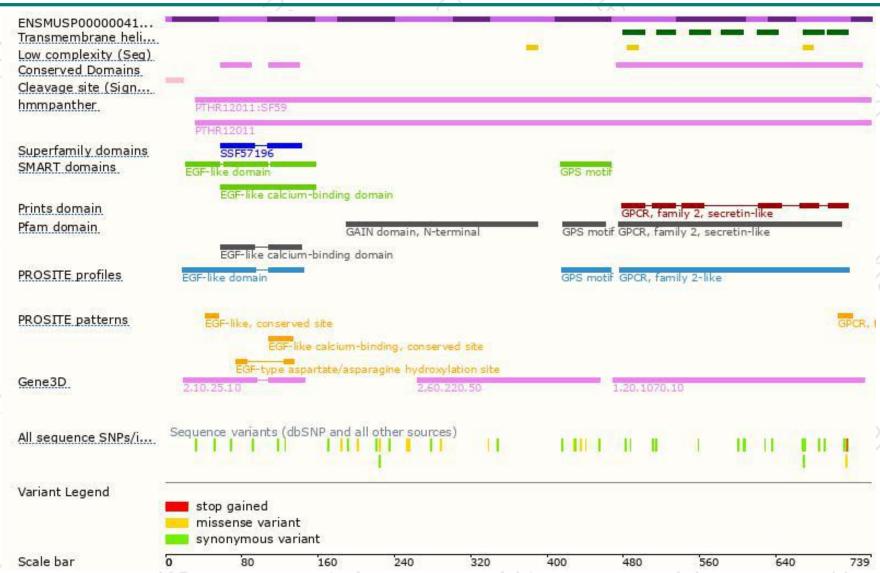
Genomic location distribution





Protein domain







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

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