

Cd2 Cas9-CKO Strategy

Designer: Reviewer:

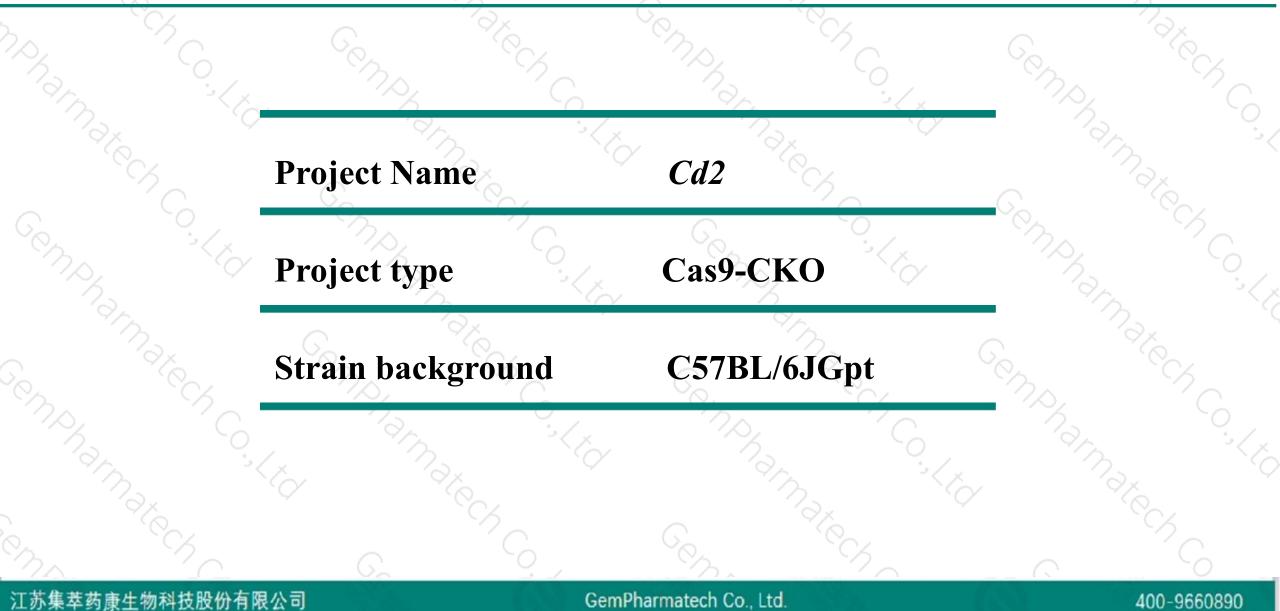
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Design Date:

Daohua Xu Huimin Su 2019-9-5

Project Overview



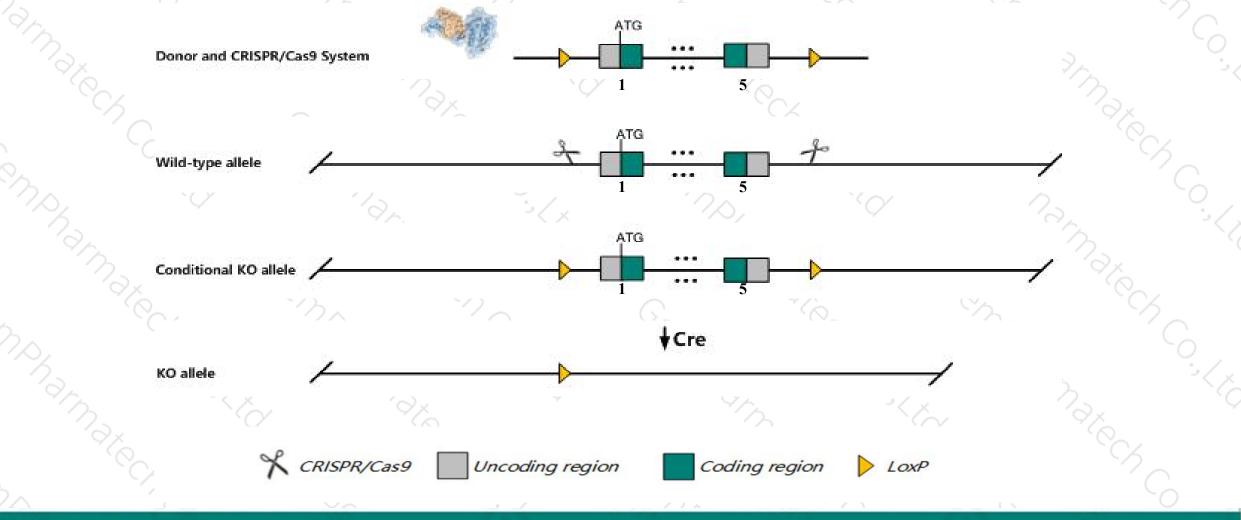


Conditional Knockout strategy



400-9660890

This model will use CRISPR/Cas9 technology to edit the Cd2 gene. The schematic diagram is as follows:



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The Cd2 gene has 3 transcripts. According to the structure of Cd2 gene, exon1-exon5 of Cd2-201 (ENSMUST00000029456.4) transcript is recommended as the knockout region. The region contains all 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 Cd2 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 mutation of this gene does not appear to result in a phenotype.
- The Cd2 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)



☆ ?

Cd2 CD2 antigen [Mus musculus (house mouse)]

Gene ID: 12481, updated on 31-Jan-2019

Summary

Official SymbolCd2 provided by MGIOfficial Full NameCD2 antigen provided byMGIPrimary sourceMGI:MGI:88320See relateEnsembl:ENSMUSG0000027863Gene typeprotein codingprotein codingPROVISIONALOrganismMus musculusLineageEukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Golires; Rodentia; Myomorpha;
Muroidea; Murinae; Mus; MusAlso knownasLFA-2, Ly-37, Ly37ExpressionBiased expression in thymus adult (RPKM 30.1), spleen adult (RPKM 18.0) and 2 other tissues
See more

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The gene has 3 transcripts, all transcripts are shown below:

| Name | Transcript ID | bp | Protein | Biotype | CCDS | UniProt | Flags | |
|---------|----------------------|------|--------------|-----------------|-----------|---------------|-------------------------------|--|
| Cd2-201 | ENSMUST00000029456.4 | 1153 | <u>344aa</u> | Protein coding | CCDS17681 | P08920 Q549Q4 | TSL:1 GENCODE basic APPRIS P1 | |
| Cd2-203 | ENSMUST00000156103.1 | 968 | No protein | Retained intron | - | ् स | TSL:1 | |
| Cd2-202 | ENSMUST00000152321.1 | 659 | No protein | Retained intron | 2 | 12 | TSL:2 | |

The strategy is based on the design of Cd2-201 transcript, The transcription is shown below

< Cd2-201 protein coding

Reverse strand ·

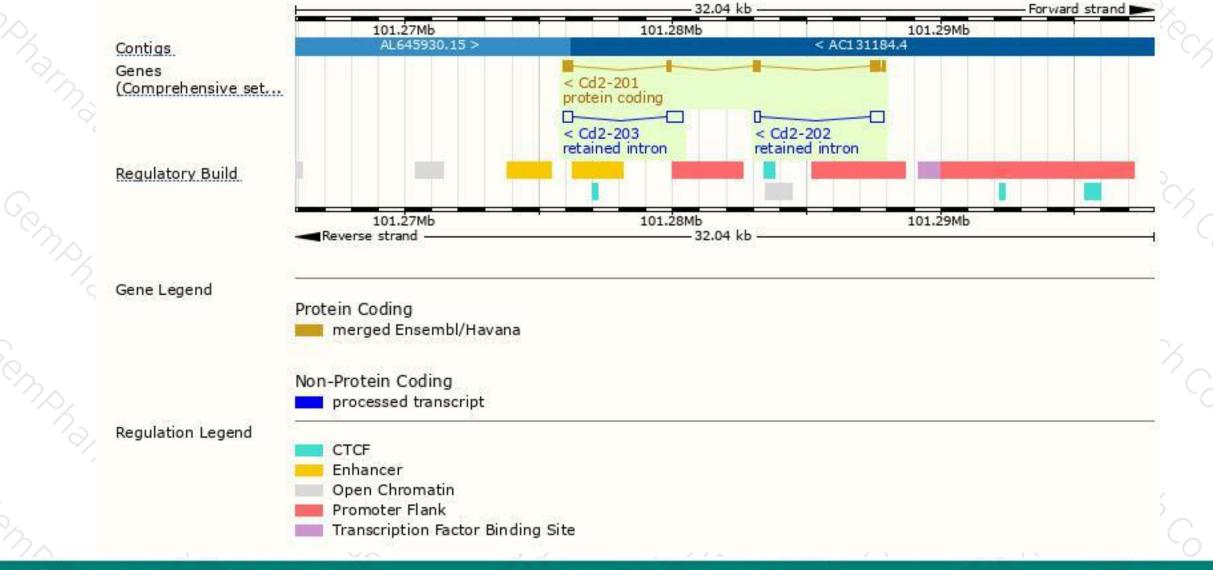
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12.04 kb

Genomic location distribution





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Protein domain



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|----|---|-------------------------|---|---------------------------------------|-----------------|-------------|-------|--------|-------|
| ₹. | ENSMUSP0000029 Transmembrane heli MobiDB lite Low complexity (Seg) Cleavage site (Sign Superfamily Prints | No. of Concession, Name | noglobulin-like do | | 5 | | | | |
| 2 | Pfam. | | l surface antigen (noglobulin V-set d | 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 | nmunoglobulin C | 2-set | | | S |
| | PANTHER | PTHR12080:SF | 14 | | | | | | ·< |
| | Gene3D | PTHR12080 | oglobulin-like fold | 145 | | | | | |
| 2 | All sequence SNPs/i | Sequence var | iants (dbSNP an | d all other source | ces) | ener ar car | 21.00 | an con | 6 |
| | Variant Legend | missense synonym | e variant Ious variant | | | | | | -< |
| | Scale bar | 6 | 40 80 | 120 | 160 | 200 | 240 | 280 | 344 |
| | °C A | | | 0 | Con . | A C | | | S Co |

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



