

Ccdc8 Cas9-KO Strategy

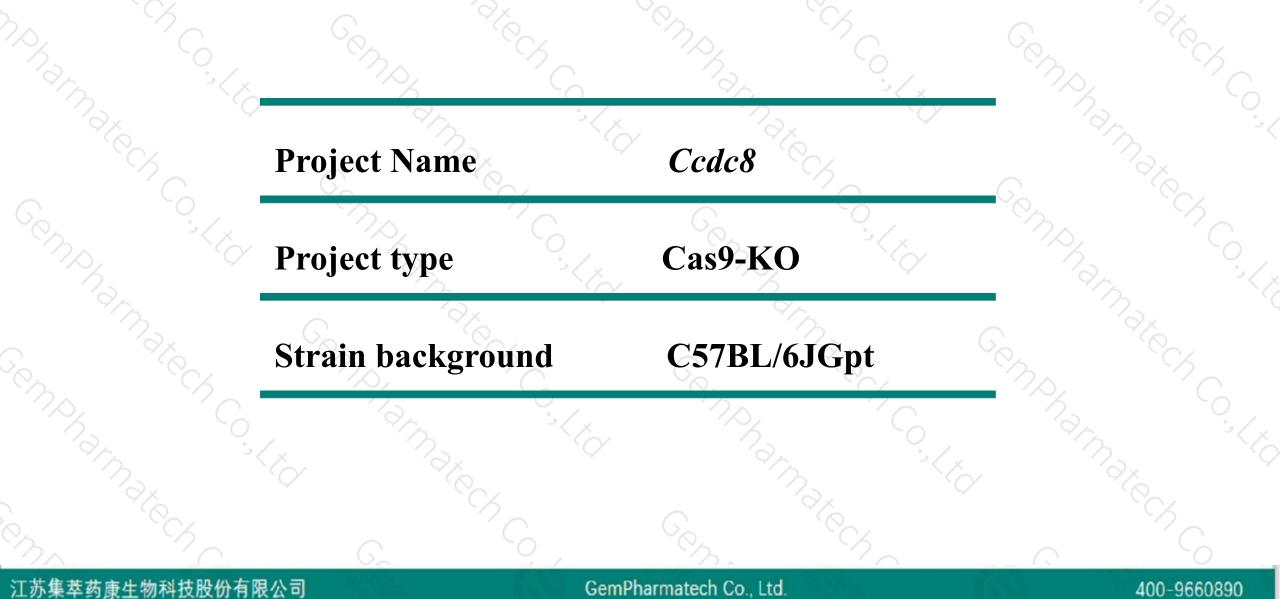
Designer: Jinling Wang

Reviewer: Miaomiao Cui

Design Date:2018-11-5

Project Overview

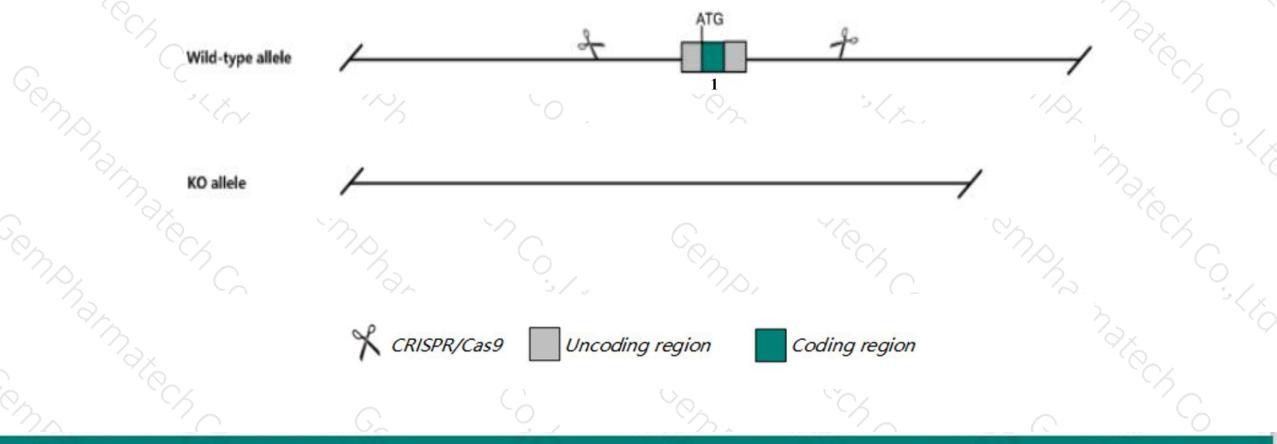




Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Ccdc8 gene. The schematic diagram is as follows:



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> The *Ccdc8* gene has 1 transcript. According to the structure of *Ccdc8* gene, exon1 of *Ccdc8*-201(ENSMUST0000094805.4) 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 *Ccdc8* 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.

- > The *Ccdc8* 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.

Notice

Gene information (NCBI)



Ccdc8 coiled-coil domain containing 8 [Mus musculus (house mouse)]

Gene ID: 434130, updated on 13-Mar-2020

Summary

Official SymbolCcdc8 provided by MGIOfficial Full Namecoiled-coil domain containing 8 provided byMGIPrimary sourceMGI:MGI:3612184See relatedEnsembl:ENSMUSG0000041117Gene typeprotein codingRefSeq statusVALIDATEDOrganismMus musculusLineageEukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Myomorpha; Muroidea; Murinae; Mus; MusOrthologshuman all

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× 1

Transcript information (Ensembl)



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The gene has 1 transcript, and the transcript is shown below:

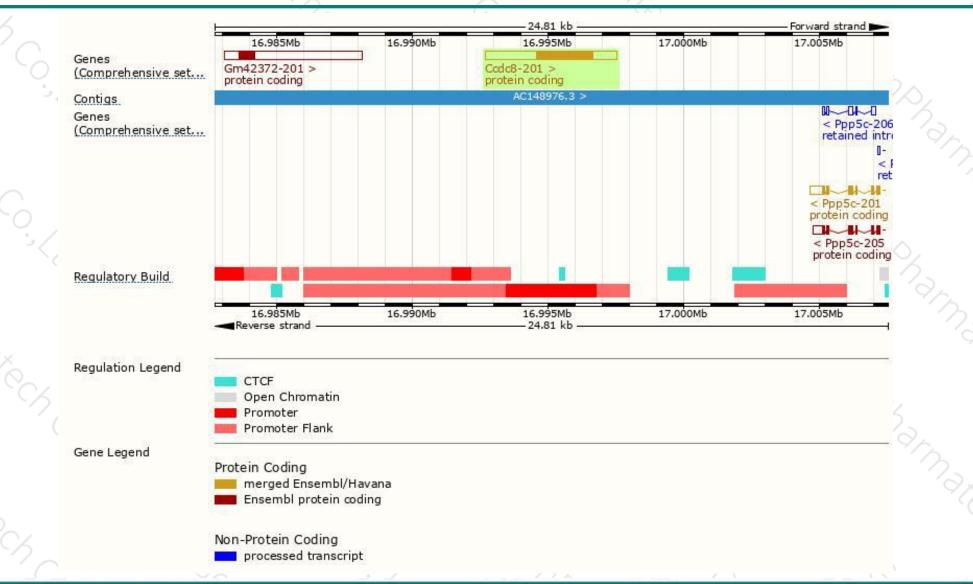
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| Name | Transcript ID | bp | Protein | Biotype | CCDS | UniProt | Flags |
|--|--|-------|--------------|--------------------|---|--------------|--------------------------------|
| cdc8-201 | ENSMUST0000094805.4 | 4809 | <u>685aa</u> | Protein coding | CCDS52044 | D3YZV8 | TSL:NA GENCODE basic APPRIS P1 |
| | | 1 | 2 | | | | |
| ~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~ | 5 | | 97. | | | 2 | |
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| | | 0 | | "3 (x | $\gamma_{\mathcal{O}}}}}}}}}}$ | | |
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| | $\mathcal{O}_{\mathcal{O}}}}}}}}}}$ | | | | | | |
| | | | | | | | |
| e strategy | is based on the design of | Ccdc8 | -201 trans | script, the transc | ription is sho | wn below: | |
| | | | | NGV. | - 0 | | |
| | | | | | | | Forward strand |
| | | | | | | | |
| c8-201 > | | | 100 | | | | |
| tein coding | | | | | | | |

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Genomic location distribution





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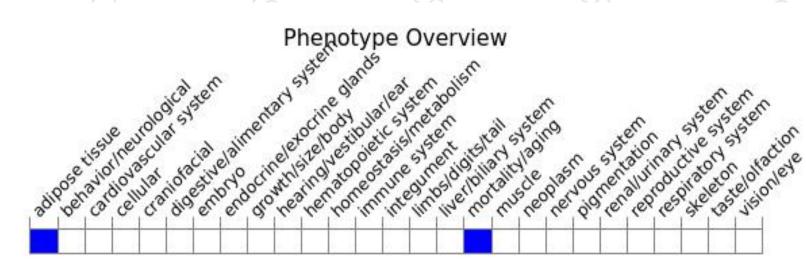
Protein domain





Mouse phenotype description(MGI)





Phenotypes affected by the gene are marked in blue.Data quoted from MGI database(http://www.informatics.jax.org/).



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



