

# Dcdc2a Cas9-CKO Strategy

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Reviewer: Xiaojing Li

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# **Project Overview**



**Project Name** 

Dcdc2a

**Project type** 

Cas9-CKO

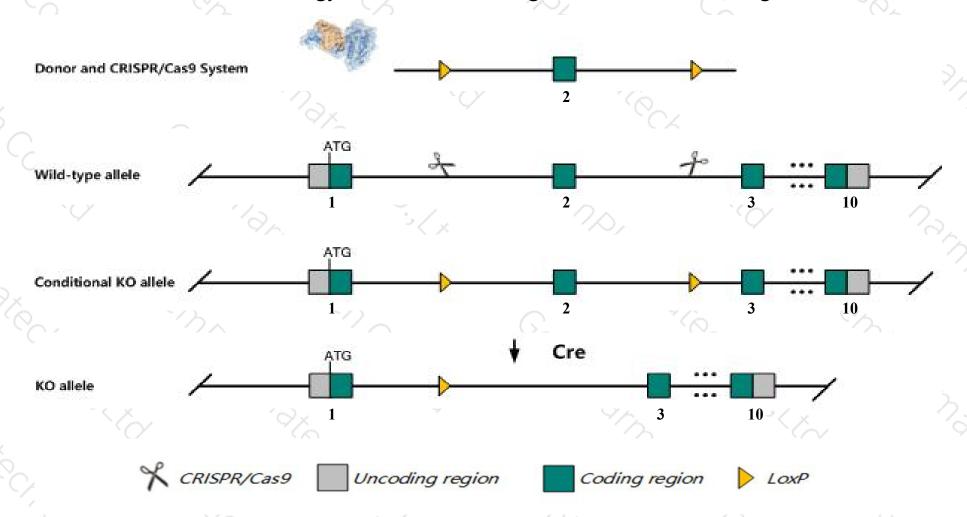
Strain background

C57BL/6JGpt

# Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Dcdc2a* gene. The schematic diagram is as follows:



## Technical routes



- ➤ The *Dcdc2a* gene has 4 transcripts. According to the structure of *Dcdc2a* gene, exon2 of *Dcdc2a-202*(ENSMUST00000069614.6) transcript is recommended as the knockout region. The region contains 55bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Dcdc2a* 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, Mice homozygous for a knock-out allele exhibit impaired short term object recognition, impaired visuo-spatial learning and memory and increased anxiety-related response.
- The *Dcdc2a* gene is located on the Chr13. 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)



#### Dcdc2a doublecortin domain containing 2a [Mus musculus (house mouse)]

Gene ID: 195208, updated on 31-Jan-2019

#### Summary

↑ ?

Official Symbol Dcdc2a provided by MGI

Official Full Name doublecortin domain containing 2a provided by MGI

Primary source MGI:MGI:2652818

See related Ensembl:ENSMUSG00000035910

Gene type protein coding
RefSeq status REVIEWED
Organism Mus musculus

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

Muroidea; Muridae; Murinae; Mus; Mus

Also known as AW492955, Dcdc2, RU2

Summary This gene encodes a member of the doublecortin family. The protein encoded by this gene contains two doublecortin domains. The

doublecortin domain has been demonstrated to bind tubulin and enhance microtubule polymerization. Alternatively spliced transcript variants

encoding distinct isoforms have been found for this gene. [provided by RefSeq, Sep 2010]

Expression Low expression observed in reference datasetSee more

Orthologs <u>human all</u>

# Transcript information (Ensembl)



The gene has 4 transcripts, all transcripts are shown below:

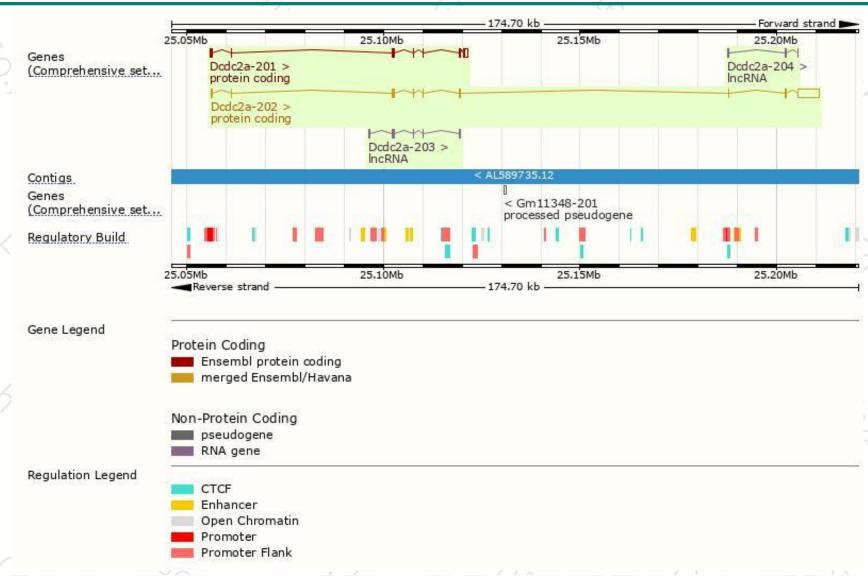
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Dcdc2a-202	ENSMUST00000069614.6	6634	<u>475aa</u>	Protein coding	CCDS26386	R4GML1	TSL:1 GENCODE basic APPRIS P1
Dcdc2a-201	ENSMUST00000036932.14	2345	<u>368aa</u>	Protein coding	CCDS56873	Q5DU00	TSL:1 GENCODE basic
Dcdc2a-203	ENSMUST00000159129.1	678	No protein	IncRNA	2	84	TSL:3
Dcdc2a-204	ENSMUST00000160363.1	661	No protein	IncRNA		12	TSL:3

The strategy is based on the design of Dcdc2a-202 transcript, The transcription is shown below



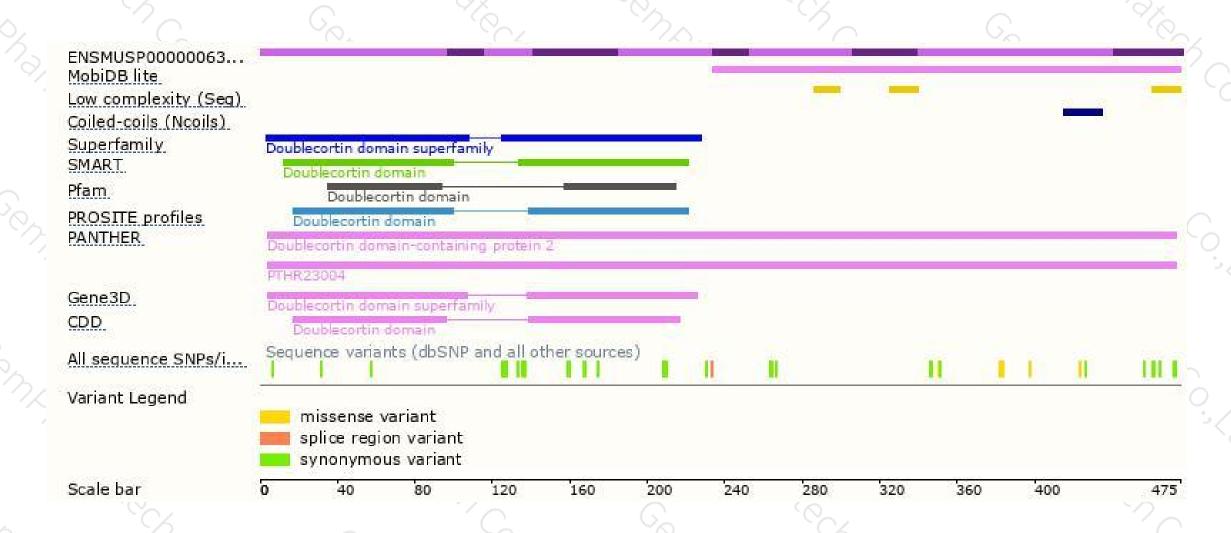
### Genomic location distribution





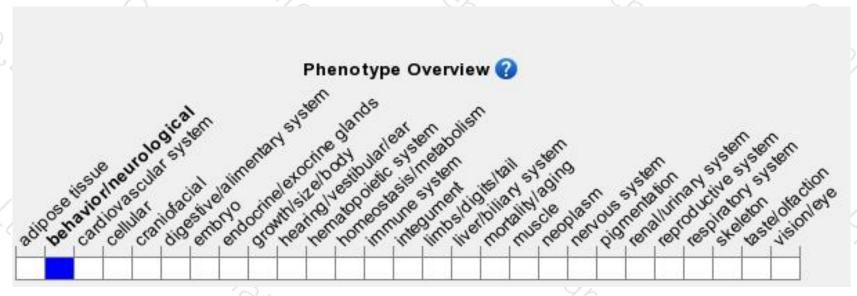
### Protein domain





# Mouse phenotype description(MGI)





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

According to the existing MGI data, Mice homozygous for a knock-out allele exhibit impaired short term object recognition, impaired visuo-spatial learning and memory and increased anxiety-related response.



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





