

# Otud6a Cas9-CKO Strategy

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# **Project Overview**



Project Name Otud6a

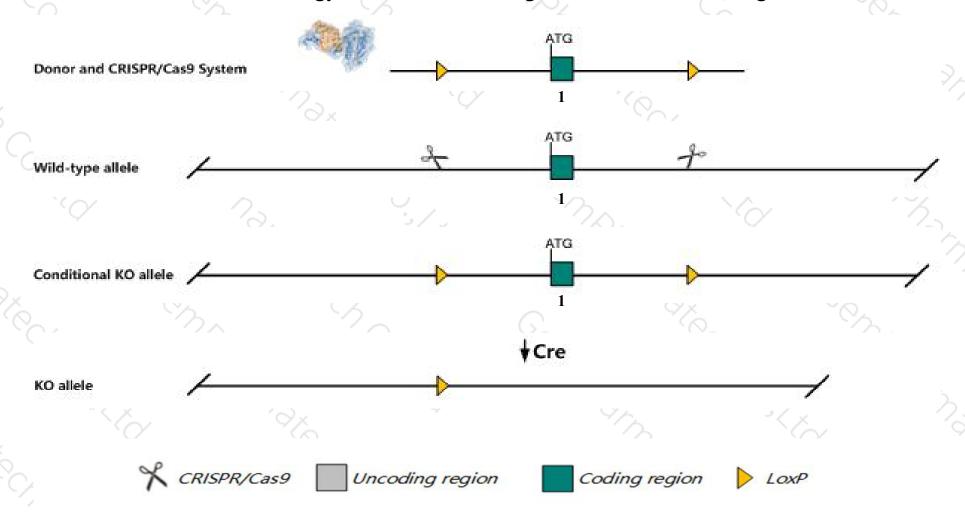
Project type Cas9-CKO

Strain background C57BL/6JGpt

## Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Otud6a* gene. The schematic diagram is as follows:



### Technical routes



- The *Otud6a* gene has 1 transcript. According to the structure of *Otud6a* gene, exon1 of *Otud6a-201* (ENSMUST00000060241.2) 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 *Otud6a* 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**



- > The *Otud6a* gene is located on the ChrX. 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)



#### Otud6a OTU domain containing 6A [Mus musculus (house mouse)]

Gene ID: 408193, updated on 13-Mar-2020

#### Summary

↑ ?

Official Symbol Otud6a provided by MGI

Official Full Name OTU domain containing 6A provided by MGI

Primary source MGI:MGI:3644685

See related Ensembl: ENSMUSG00000051582

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 EG408193, Hshin6

Orthologs <u>human</u> all

# Transcript information (Ensembl)



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

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags	
Otud6a-201	ENSMUST00000060241.2	873	290aa	Protein coding	CCDS53146	Q6IE21	TSL:NA GENCODE basic APPRIS P1	L

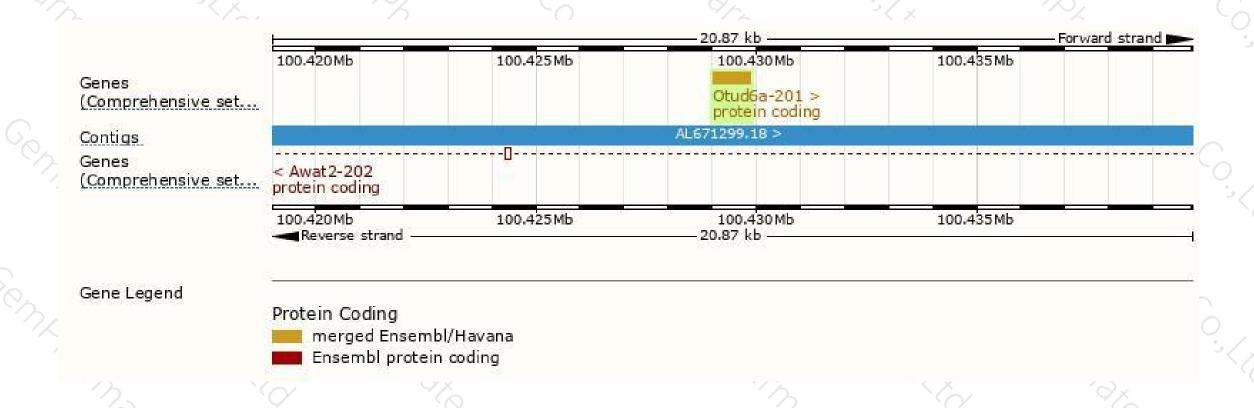
The strategy is based on the design of Otud6a-201 transcript, the transcription is shown below:

· 873 bp — Forward strand **>** 

Otud6a-201 > protein coding

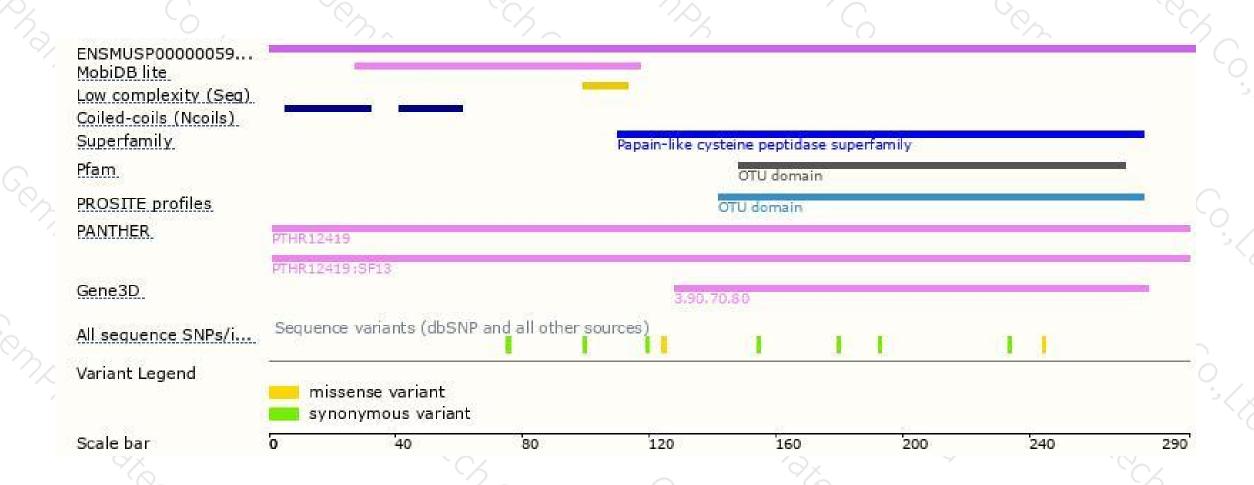
### Genomic location distribution





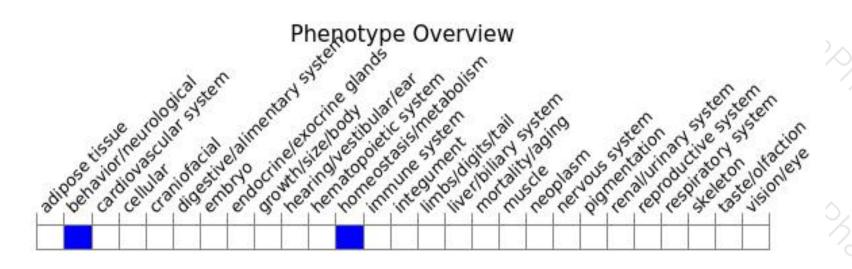
### Protein domain





# Mouse phenotype description(MGI)





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



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





