

# Zw10 Cas9-KO Strategy

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# **Project Overview**



Project Name Zw10

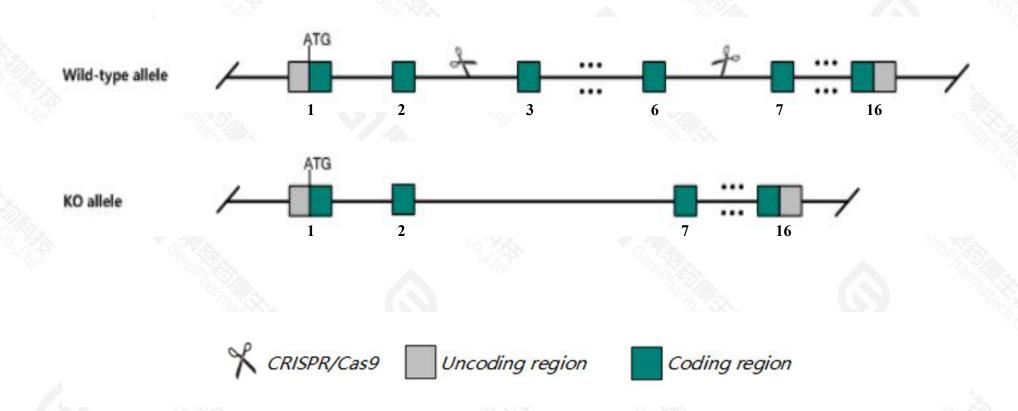
Project type Cas9-KO

Strain background C57BL/6JGpt

# **Knockout strategy**



This model will use CRISPR/Cas9 technology to edit the Zw10 gene. The schematic diagram is as follows:



### **Technical routes**



- The Zw10 gene has 3 transcripts. According to the structure of Zw10 gene, exon3-exon6 of Zw10201(ENSMUST00000034803.10) transcript is recommended as the knockout region. The region contains 493bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Zw10* 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.

### **Notice**



- > The Zw10 gene is located on the Chr9. 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.
- > The KO region nears the 1700042D02Rik gene. Knockout the region may affect the function of 1700042D02Rik gene.

## Gene information (NCBI)



#### Zw10 zw10 kinetochore protein [Mus musculus (house mouse)]

Gene ID: 26951, updated on 17-Dec-2020

#### Summary

☆ ?

Official Symbol Zw10 provided by MGI

Official Full Name zw10 kinetochore protein provided byMGI

Primary source MGI:MGI:1349478

See related Ensembl: ENSMUSG00000032264

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 6330566F14Rik, MmZw10

Expression Ubiquitous expression in CNS E11.5 (RPKM 15.1), limb E14.5 (RPKM 13.7) and 28 other tissuesSee more

Orthologs <u>human all</u>

# Transcript information (Ensembl)



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

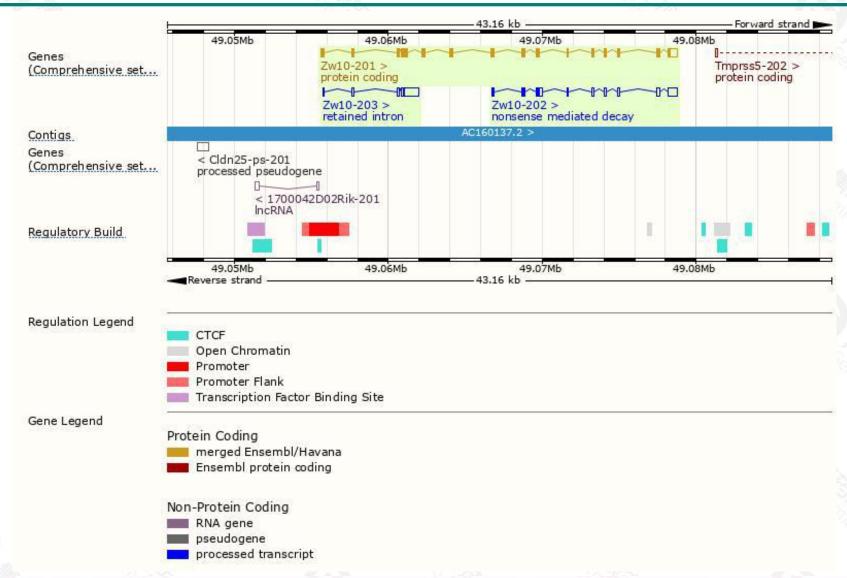
| Name     | Transcript ID         | bp   | Protein      | Biotype                 | CCDS      | UniProt | Flags                            |
|----------|-----------------------|------|--------------|-------------------------|-----------|---------|----------------------------------|
| Zw10-201 | ENSMUST00000034803.10 | 2863 | <u>779aa</u> | Protein coding          | CCDS23161 |         | TSL:1, GENCODE basic, APPRIS P1, |
| Zw10-202 | ENSMUST00000213787.2  | 2013 | <u>187aa</u> | Nonsense mediated decay | -         |         | CDS 5' incomplete , TSL:1 ,      |
| Zw10-203 | ENSMUST00000215706.2  | 1272 | No protein   | Retained intron         | 12        |         | TSL:2,                           |

The strategy is based on the design of Zw10-201 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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