

Cib3 Cas9-CKO Strategy

Designer: Yanhua Shen

Project Overview

Project Name

Cib3

Project type

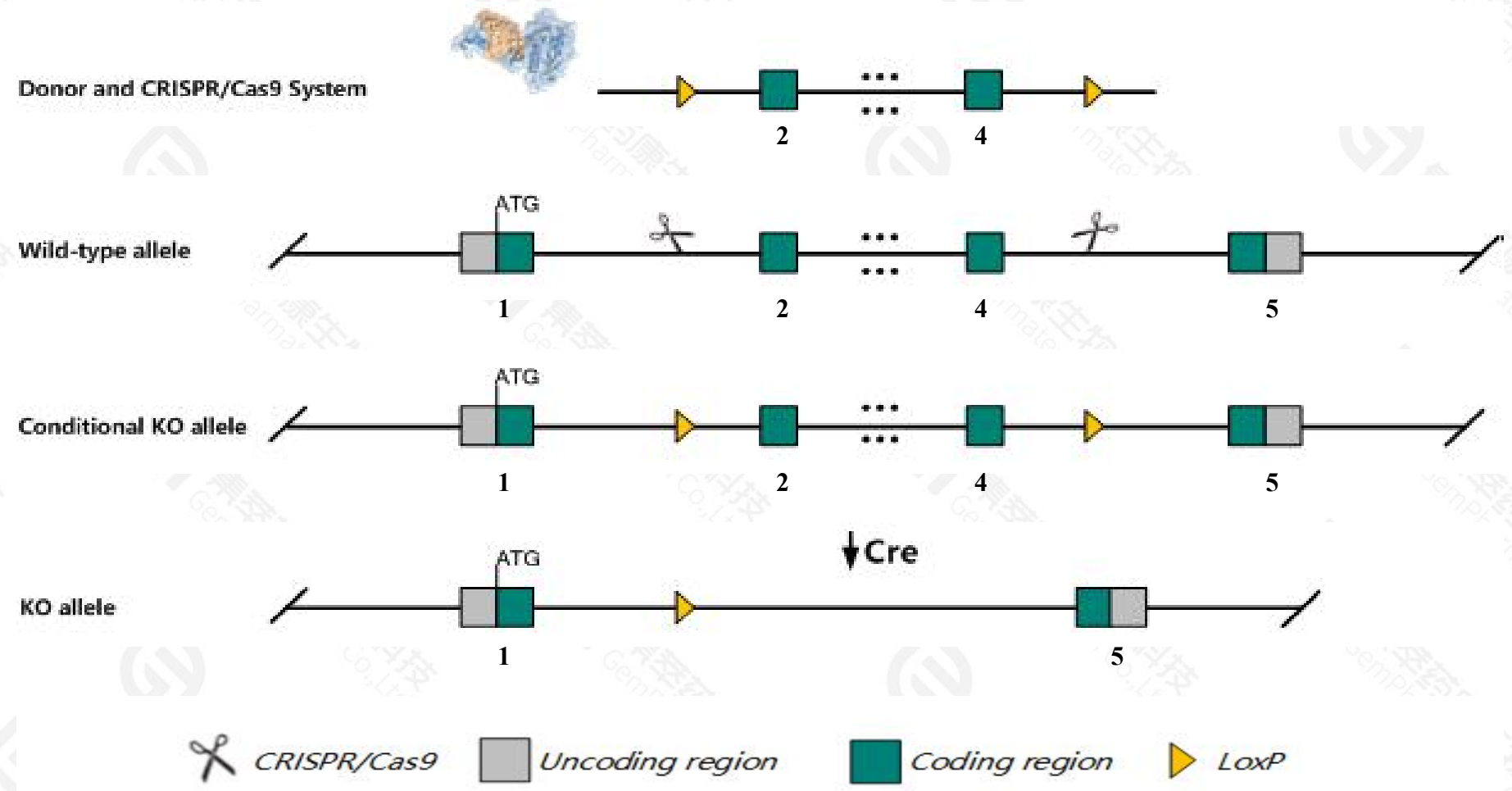
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Cib3* gene. The schematic diagram is as follows:



Technical routes

- The *Cib3* gene has 3 transcripts. According to the structure of *Cib3* gene, exon2-exon4 of *Cib3*-201(ENSMUST00000098630.5) transcript is recommended as the knockout region. The region contains most 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 *Cib3* 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.

- The KO region deletes most of the coding sequence, but does not result in frameshift.
- The *Cib3* gene is located on the Chr8. 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)

Cib3 calcium and integrin binding family member 3 [Mus musculus (house mouse)]

Gene ID: 234421, updated on 26-Sep-2020

Summary



Official Symbol Cib3 provided by [MGI](#)

Official Full Name calcium and integrin binding family member 3 provided by [MGI](#)

Primary source [MGI:MGI:2685953](#)

See related [Ensembl:ENSMUSG00000074240](#)

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 C730014M21Rik, Gm1107, KI, KIP3

Expression Low expression observed in reference dataset [See more](#)

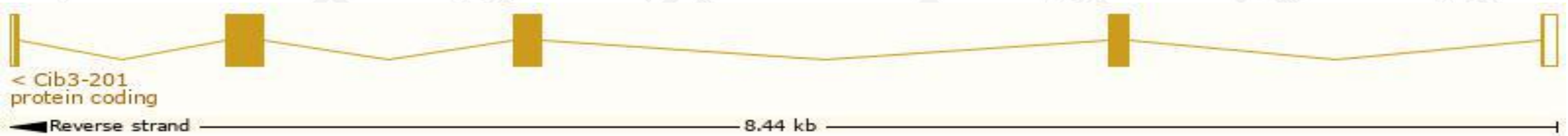
Orthologs [human](#) [all](#)

Transcript information (Ensembl)

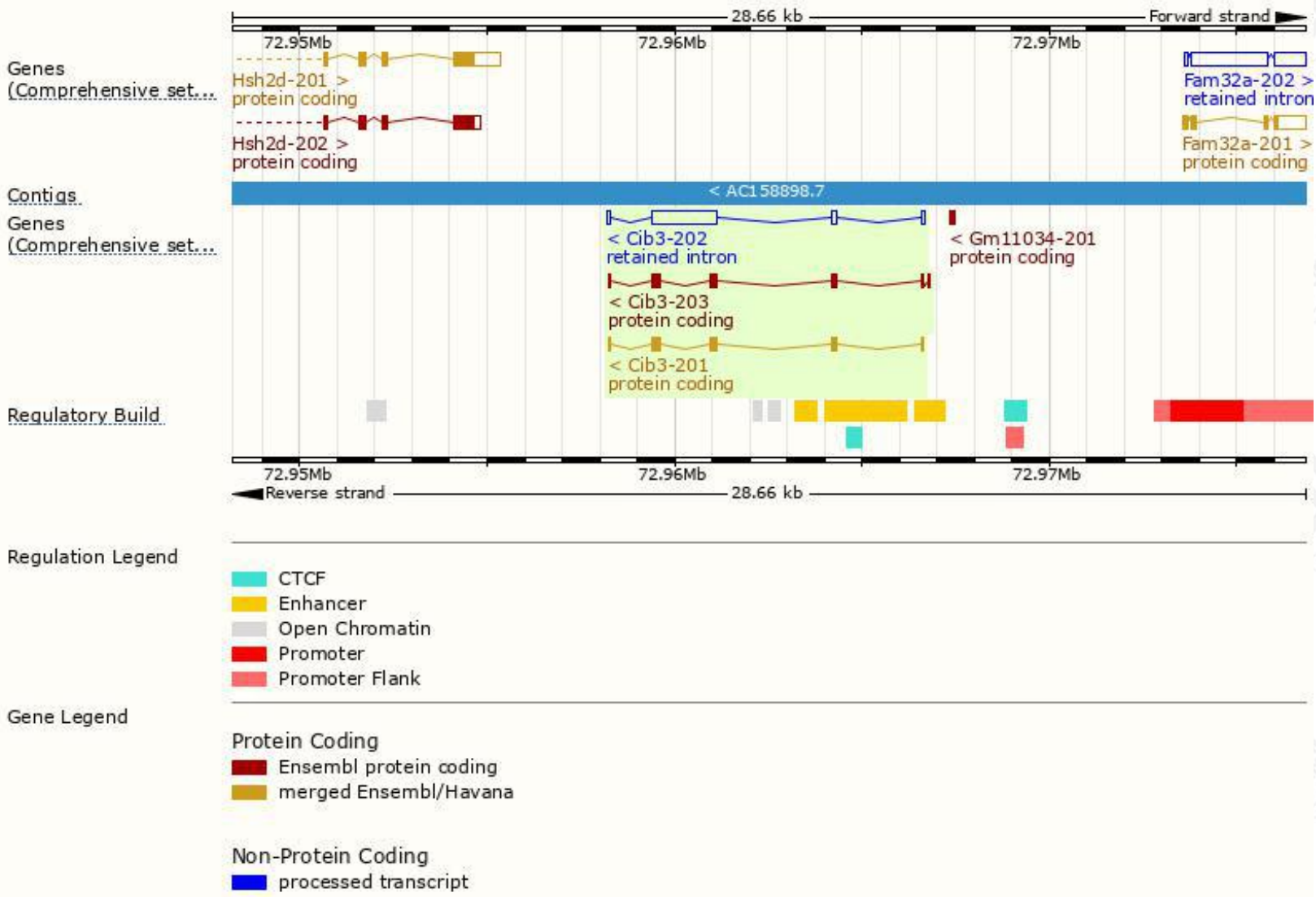
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Cib3-201	ENSMUST00000098630.5	585	160aa	Protein coding	CCDS52594		TSL:1 , GENCODE basic ,
Cib3-203	ENSMUST00000238973.2	617	187aa	Protein coding	-		GENCODE basic , APPRIS P1 ,
Cib3-202	ENSMUST00000211946.2	2014	No protein	Retained intron	-		TSL:1 ,

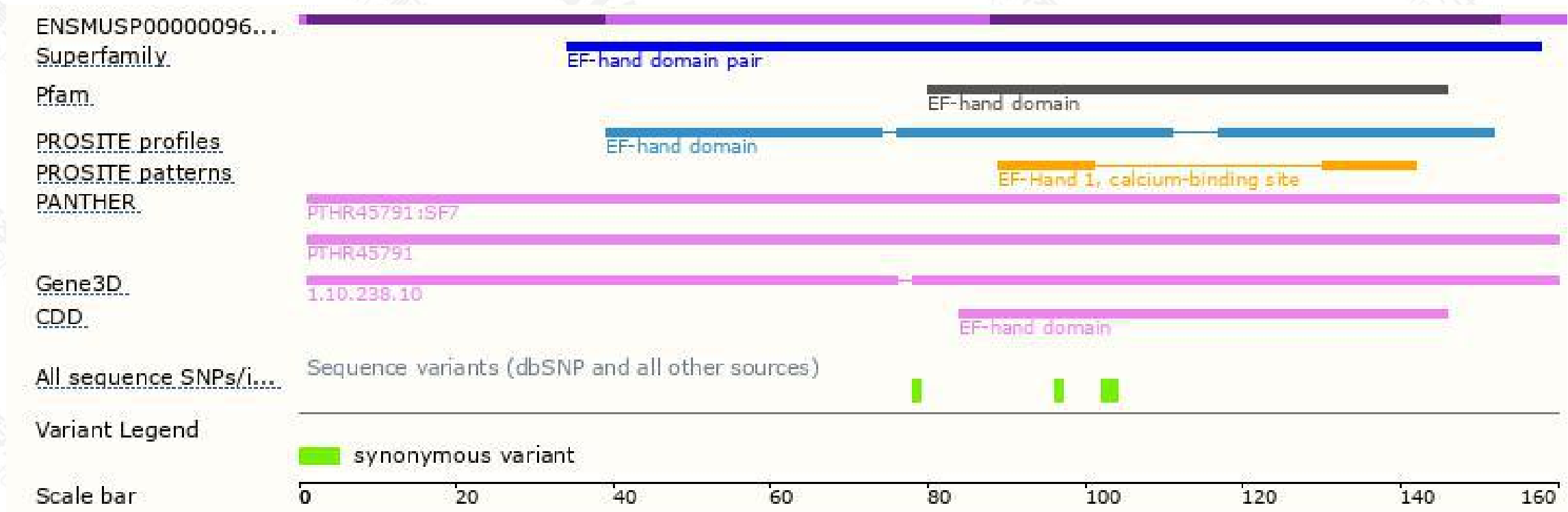
The strategy is based on the design of *Cib3-201* transcript,the transcription is shown below:



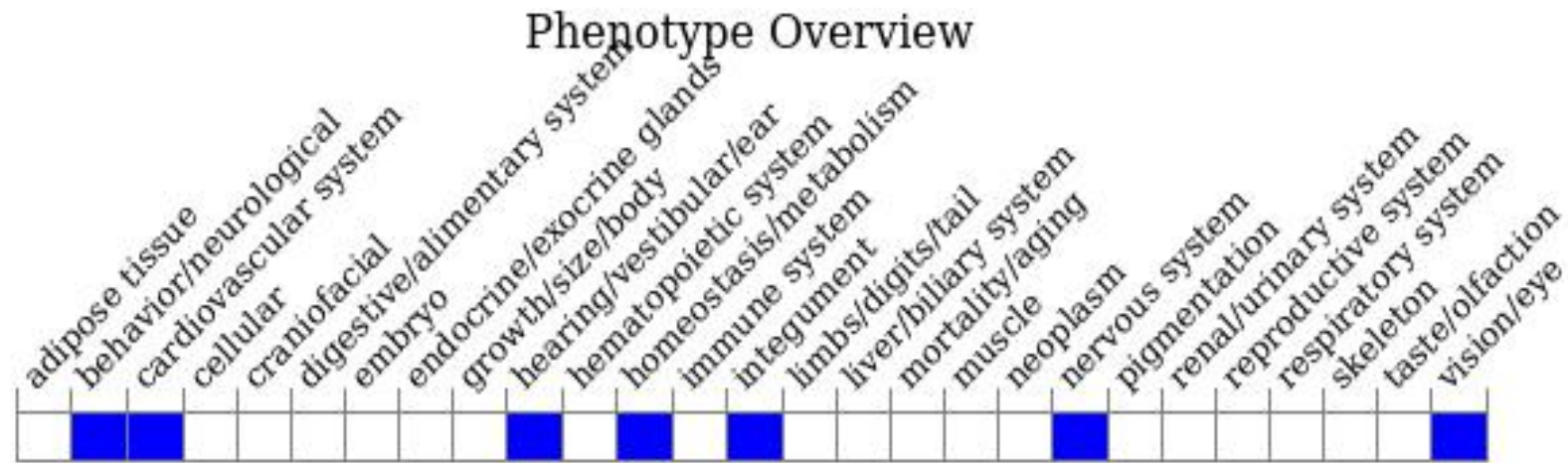
Genomic location distribution



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/>).

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

