

# Gfilb Cas9-CKO Strategy

**Designer: Yupeng Yang** 

Reviewer: Rui Xiong

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## **Project Overview**



**Project Name** 

Gfi1b

**Project type** 

Cas9-CKO

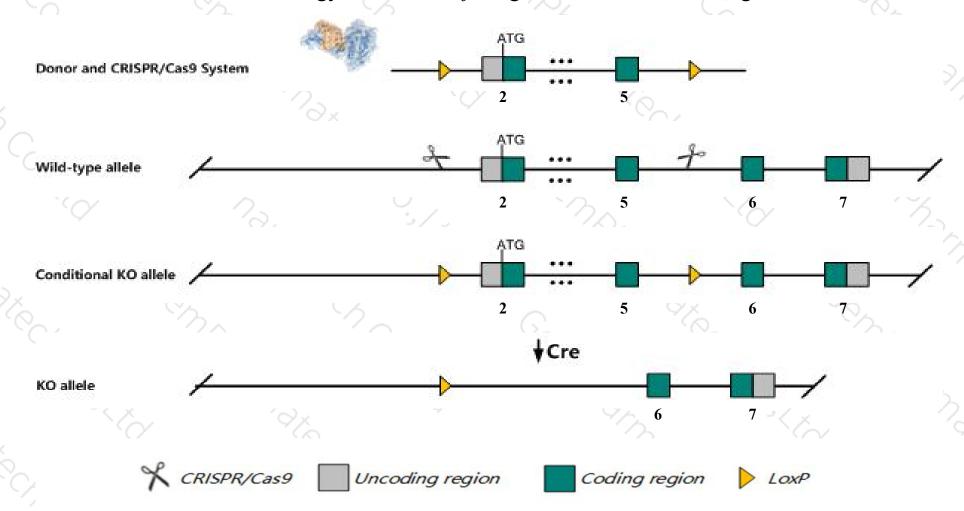
Strain background

C57BL/6JGpt

## Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Gfi1b* gene. The schematic diagram is as follows:



### Technical routes



- The *Gfi1b* gene has 4 transcripts. According to the structure of *Gfi1b* gene, exon2-exon5 of *Gfi1b*-204(ENSMUST00000164290.7) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Gfi1b* 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 disruption of this gene die as embryos by day E15. Mature adult red blood cells and megakaryocytes fail to develop.
- The *Gfi1b* gene is located on the Chr2. 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)



#### Gfi1b growth factor independent 1B [Mus musculus (house mouse)]

Gene ID: 14582, updated on 13-Mar-2020

#### Summary

☆ ?

Official Symbol Gfi1b provided by MGI

Official Full Name growth factor independent 1B provided by MGI

Primary source MGI:MGI:1276578

See related Ensembl:ENSMUSG00000026815

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 Gfi-1B

Expression Biased expression in liver E14.5 (RPKM 51.1), liver E14 (RPKM 38.6) and 2 other tissuesSee more

Orthologs <u>human</u> all

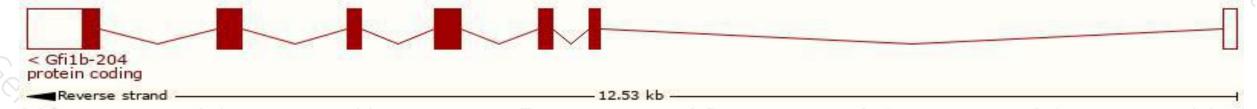
## Transcript information (Ensembl)



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

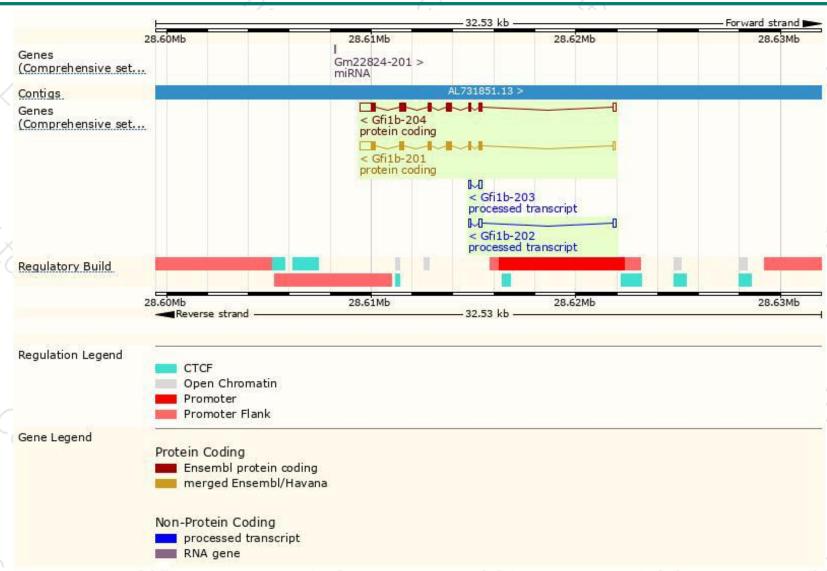
| Name      | Transcript ID        | bp   | Protein      | Biotype              | CCDS      | UniProt | Flags                           |
|-----------|----------------------|------|--------------|----------------------|-----------|---------|---------------------------------|
| Gfi1b-204 | ENSMUST00000164290.7 | 1822 | 363aa        | Protein coding       | CCDS50549 | B7ZNH2  | TSL:1 GENCODE basic APPRIS ALT2 |
| Gfi1b-201 | ENSMUST00000028156.7 | 1681 | <u>330aa</u> | Protein coding       | CCDS15843 | 070237  | TSL:1 GENCODE basic APPRIS P3   |
| Gfi1b-202 | ENSMUST00000145690.1 | 357  | No protein   | Processed transcript | 828       | 724     | TSL:2                           |
| Gfi1b-203 | ENSMUST00000155686.1 | 238  | No protein   | Processed transcript | -         | (%)     | TSL:3                           |

The strategy is based on the design of *Gfi1b-204* transcript, the transcription is shown below:



### Genomic location distribution





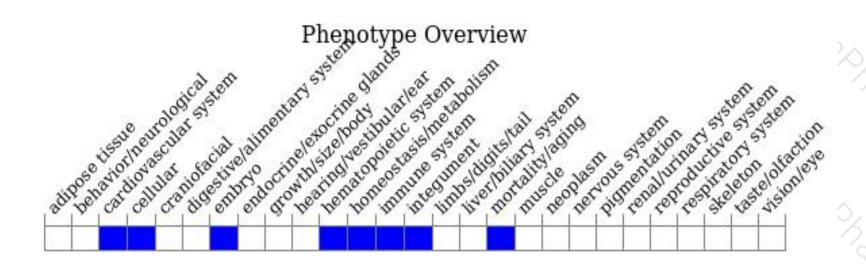
### Protein domain





## Mouse phenotype description(MGI)





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

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





