

# Asb13 Cas9-CKO Strategy

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**Reviewer:** 

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# **Project Overview**



**Project Name** 

Asb13

**Project type** 

Cas9-CKO

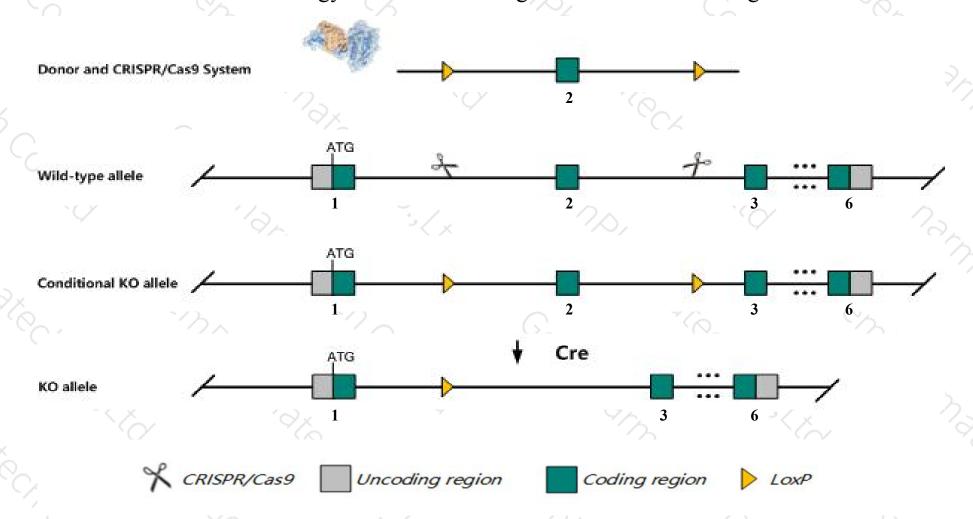
Strain background

C57BL/6JGpt

## Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Asb13 gene. The schematic diagram is as follows:



### Technical routes



- ➤ The *Asb13* gene has 4 transcripts. According to the structure of *Asb13* gene, exon2 of *Asb13-201*(ENSMUST00000042288.7) transcript is recommended as the knockout region. The region contains 188bp coding sequence.

  Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Asb13* 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 *Asb13* 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)



#### Asb13 ankyrin repeat and SOCS box-containing 13 [Mus musculus (house mouse)]

Gene ID: 142688, updated on 31-Jan-2019

#### Summary

☆ ?

Official Symbol Asb13 provided by MGI

Official Full Name ankyrin repeat and SOCS box-containing 13 provided by MGI

Primary source MGI:MGI:2145525

See related Ensembl: ENSMUSG00000033781

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 2210015B19Rik, 6430573K02Rik, Asb-13, C85285

Expression Ubiquitous expression in kidney adult (RPKM 25.8), liver adult (RPKM 18.0) and 27 other tissuesSee more

Orthologs human all

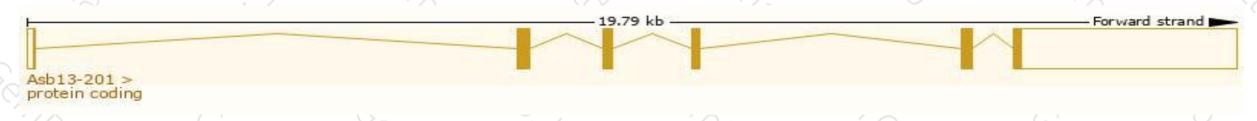
# Transcript information (Ensembl)



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

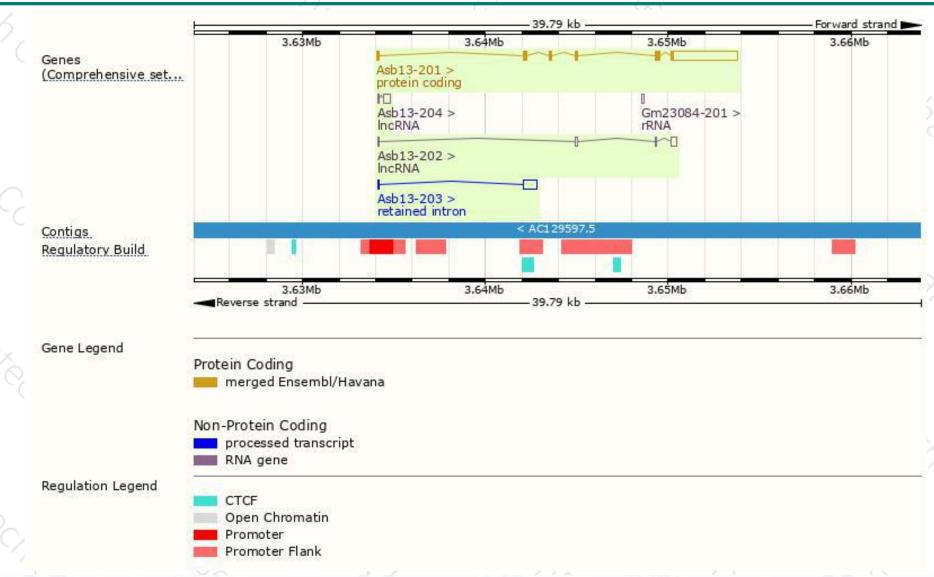
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Asb13-201	ENSMUST00000042288.7	4453	278aa	Protein coding	CCDS36582	Q8VBX0	TSL:1 GENCODE basic APPRIS P1
Asb13-203	ENSMUST00000141967.1	838	No protein	Retained intron	. 8	-	TSL:3
Asb13-202	ENSMUST00000141147.1	576	No protein	IncRNA	26	825	TSL:3
Asb13-204	ENSMUST00000156595.1	456	No protein	IncRNA	-	727	TSL:1

The strategy is based on the design of Asb13-201 transcript, The transcription is shown below



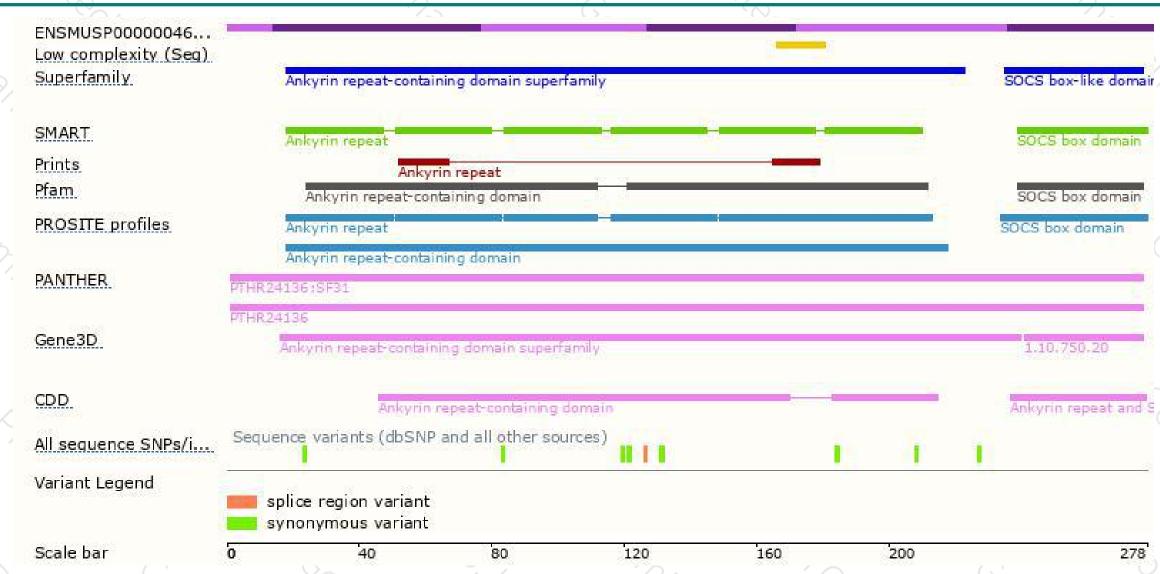
### Genomic location distribution





### Protein domain







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





