

Rybp Cas9-CKO Strategy

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Reviewer: Huimin Su

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Project Overview



Project Name Rybp

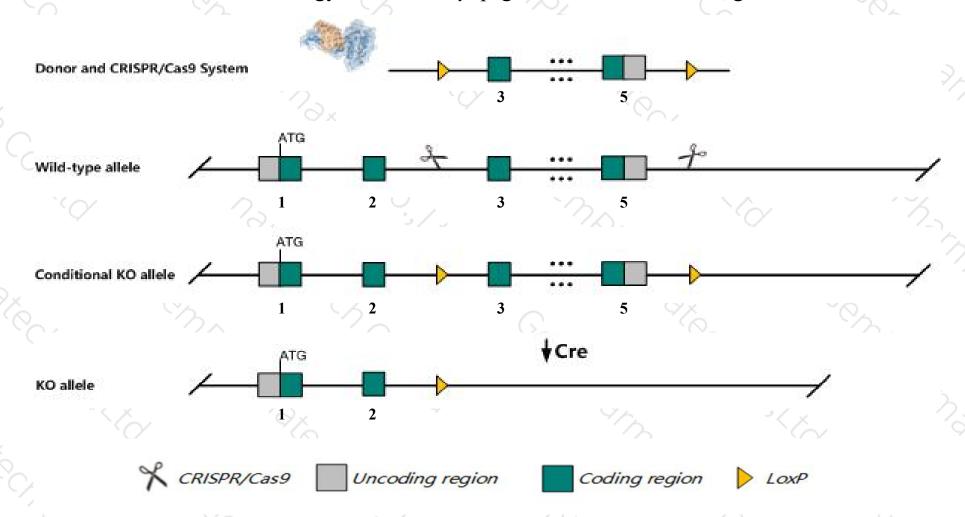
Project type Cas9-CKO

Strain background C57BL/6JGpt

Conditional Knockout strategy



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



Technical routes



- > The *Rybp* gene has 3 transcripts. According to the structure of *Rybp* gene, exon3-exon5 of *Rybp-201*(ENSMUST00000101118.3) 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 *Rybp* 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, homozygous null mice initiate implantation but fail to complete decidualization and display early embryonic lethality. a subset of heterozygotes show forebrain overgrowth, delayed neural tube closure, exencephaly, failed optic fissure closure, malformed lenses, and retinal/optic nerve colobomas.
- > The *Rybp* gene is located on the Chr6. 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)



Rybp RING1 and YY1 binding protein [Mus musculus (house mouse)]

Gene ID: 56353, updated on 13-Mar-2020

Summary

☆ ?

Official Symbol Rybp provided by MGI

Official Full Name RING1 and YY1 binding protein provided by MGI

Primary source MGI:MGI:1929059

See related Ensembl: ENSMUSG00000072872

Gene type protein coding
RefSeq status PROVISIONAL
Organism Mus musculus

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;

Muroidea; Muridae; Murinae; Mus; Mus

Also known as 2410018J24Rik, DEDAF, YEAF1

Expression Ubiquitous expression in placenta adult (RPKM 21.3), CNS E18 (RPKM 18.2) and 27 other tissuesSee more

Orthologs <u>human</u> all

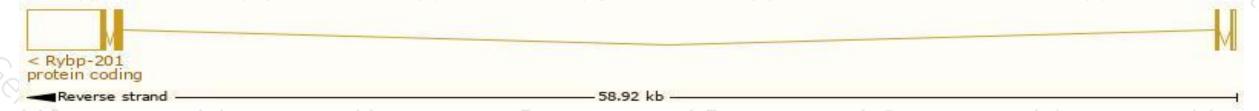
Transcript information (Ensembl)



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

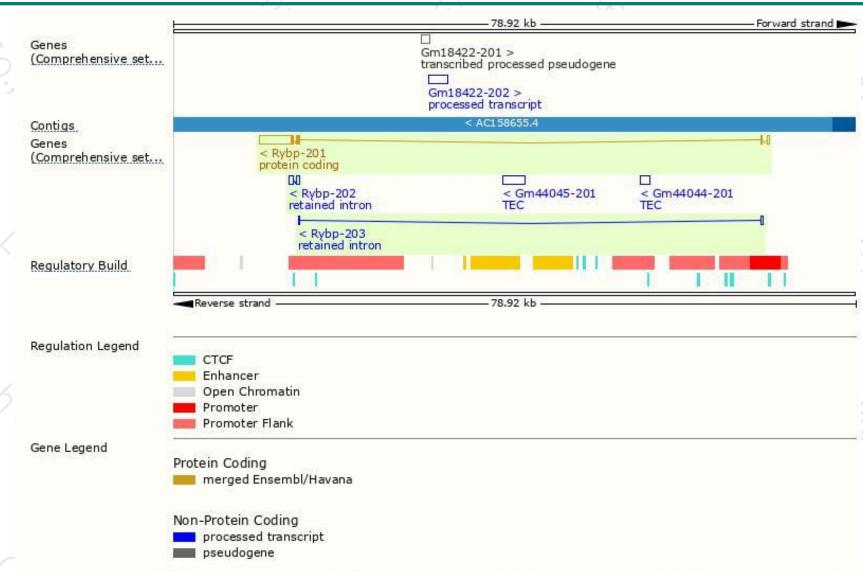
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Floor
Name	Transcript ID	pp	Protein	віотуре	CCDS	UniProt	Flags
Rybp-201	ENSMUST00000101118.3	4503	228aa	Protein coding	CCDS39579	Q8CCI5	TSL:1 GENCODE basic APPRIS is a system to annotate alternatively spliced transcripts based on a range of computational methods to identify the most functionally important transcript(s) of a gene. APPRIS
Rybp-202	ENSMUST00000203410.1	680	No protein	Retained intron	- 8	· .	TSL:2
Rybp-203	ENSMUST00000205164.1	412	No protein	Retained intron	20	-	TSL:2

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



Genomic location distribution





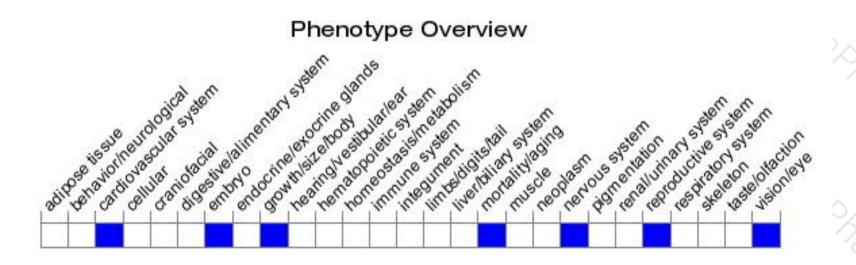
Protein domain





Mouse phenotype description(MGI)





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

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





