

# Prrc2b Cas9-KO Strategy

**Designer:** 

Reviewer

**Design Date:** 

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# **Project Overview**



**Project Name** 

Prrc2b

**Project type** 

Cas9-KO

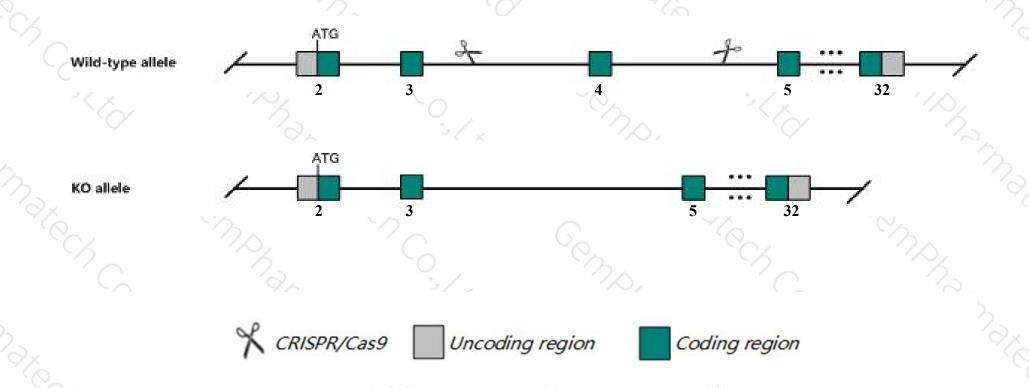
Strain background

C57BL/6JGpt

# **Knockout strategy**



This model will use CRISPR/Cas9 technology to edit the *Prrc2b* gene. The schematic diagram is as follows:



### **Technical routes**



- ➤ The *Prrc2b* gene has 10 transcripts. According to the structure of *Prrc2b* gene, exon4 of *Prrc2b-202*(ENSMUST00000069817.14) transcript is recommended as the knockout region. The region contains 103bp coding sequence Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Prrc2b* gene. The brief process is as follows: CRISPR/Cas9 system

### **Notice**



- > The *Prrc2b* 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.
- Transcript *Prrc2b*-204,210 may not be affected.
- 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.

### Gene information (NCBI)



#### Prrc2b proline-rich coiled-coil 2B [ Mus musculus (house mouse) ]

Gene ID: 227723, updated on 12-Aug-2019

#### Summary

☆ ?

Official Symbol Prrc2b provided by MGI

Official Full Name proline-rich coiled-coil 2B provided by MGI

Primary source MGI:MGI:1923304

See related Ensembl: ENSMUSG00000039262

Gene type protein coding
RefSeq status VALIDATED
Organism <u>Mus musculus</u>

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

Murinae; Mus; Mus

Also known as Bat2l; Bat2l1; Al173903; mKIAA0515; D430039P21; 5830434P21Rik

Expression Ubiquitous expression in whole brain E14.5 (RPKM 35.1), ovary adult (RPKM 32.2) and 28 other tissues See more

Orthologs <u>human</u> all

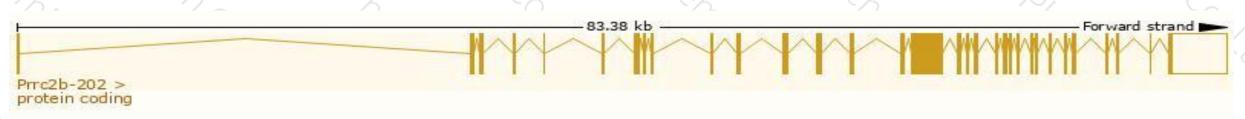
# Transcript information (Ensembl)



The gene has 10 transcripts, all transcripts are shown below:

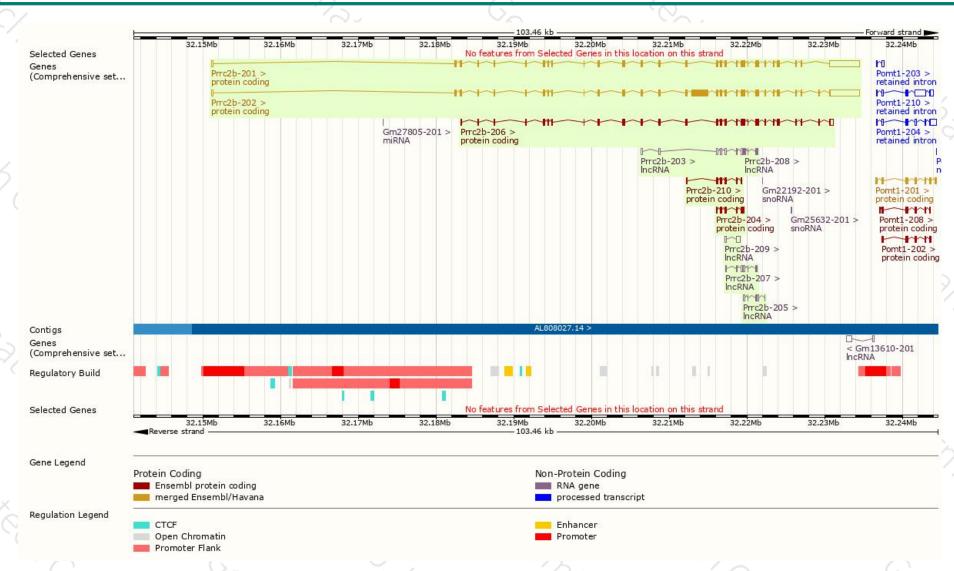
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Prrc2b-202	ENSMUST00000069817.14	10671	2230aa	Protein coding	CCDS50565	F8WHT3	TSL:1 GENCODE basic APPRIS ALT2
Prrc2b-201	ENSMUST00000036691.13	8612	1486aa	Protein coding	CCDS15907	Q7TPM1	TSL:1 GENCODE basic APPRIS P3
Prrc2b-206	ENSMUST00000132459.7	4837	1472aa	Protein coding	120	F7AYW2	CDS 5' incomplete TSL:1
Prrc2b-210	ENSMUST00000156313.7	751	<u>250aa</u>	Protein coding	750	F6V431	5' and 3' truncations in transcript evidence prevent annotation of the start and the end of the CDS. CDS 5' and 3' incomplete TSL:
Prrc2b-204	ENSMUST00000128936.1	696	<u>232aa</u>	Protein coding	120	F6Q3E1	5' and 3' truncations in transcript evidence prevent annotation of the start and the end of the CDS. CDS 5' and 3' incomplete TSL:
Prrc2b-203	ENSMUST00000123270.7	1258	No protein	Processed transcript		199	TSL:3
Prrc2b-207	ENSMUST00000140015.7	737	No protein	Processed transcript	120	94	TSL:3
Prrc2b-205	ENSMUST00000129626.7	623	No protein	Processed transcript	757	ės –	TSL:2
Prrc2b-209	ENSMUST00000142219.1	604	No protein	Processed transcript	(2)	85	TSL:5
Prrc2b-208	ENSMUST00000141053.1	383	No protein	Processed transcript		199	TSL:3

The strategy is based on the design of *Prrc2b-202* transcript, The transcription is shown below



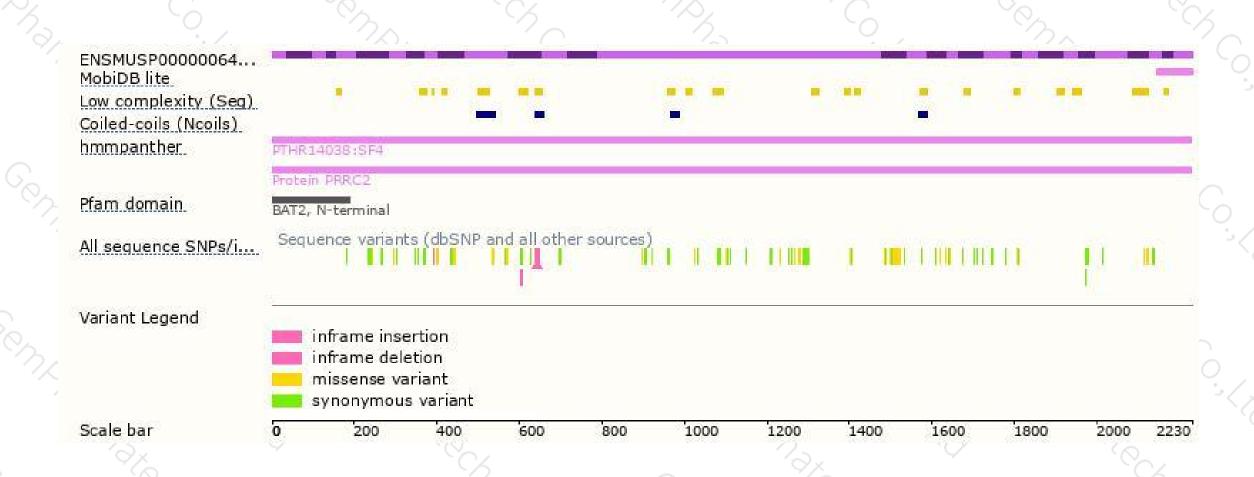
### Genomic location distribution





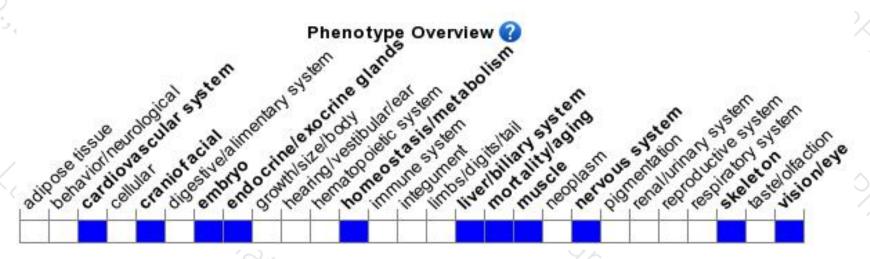
### Protein domain





# 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





