

# *Ppp3cb* Cas9-KO Strategy

**Designer:**

**Huimin Su**

**Reviewer:**

**Ruirui Zhang**

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# Project Overview

**Project Name**

*Ppp3cb*

**Project type**

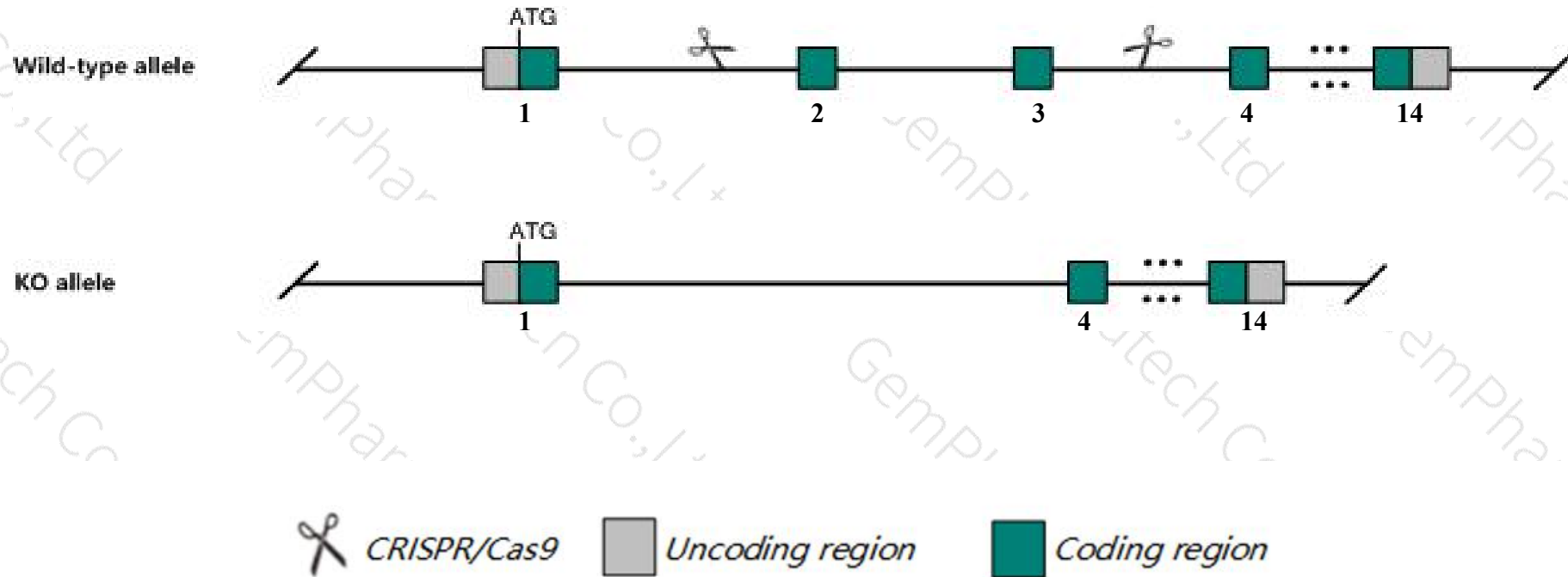
**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Ppp3cb* gene. The schematic diagram is as follows:



- The *Ppp3cb* gene has 9 transcripts. According to the structure of *Ppp3cb* gene, exon2-exon3 of *Ppp3cb*-202 (ENSMUST00000159027.7) transcript is recommended as the knockout region. The region contains 326bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Ppp3cb* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Homozygous null mice have small hearts and thymi, and reduced body weight. Cardiac function is normal, but mice lack a cardiac hypertrophic response to pressure overload, angiotensin II, or isopreteronol. Thymi are hypoplastic, with abnormal T cell development and reduced numbers of T cells.
- The *Ppp3cb* gene is located on the Chr14. 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.



# Gene information (NCBI)

## Ppp3cb protein phosphatase 3, catalytic subunit, beta isoform [ *Mus musculus* (house mouse) ]

Gene ID: 19056, updated on 12-Aug-2019

### Summary

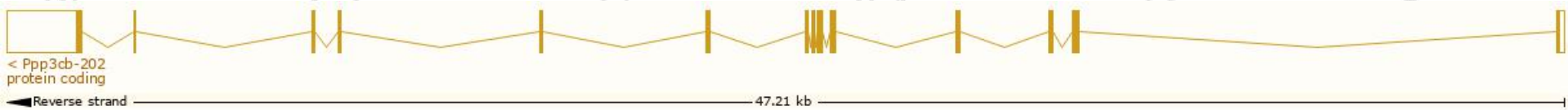
<b>Official Symbol</b>	Ppp3cb provided by <a href="#">MGI</a>
<b>Official Full Name</b>	protein phosphatase 3, catalytic subunit, beta isoform provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:107163</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000021816</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	VALIDATED
<b>Organism</b>	<a href="#">Mus musculus</a>
<b>Lineage</b>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
<b>Also known as</b>	Cnab; Calnb; CnAbeta; 1110063J16Rik
<b>Expression</b>	Broad expression in CNS E18 (RPKM 36.0), frontal lobe adult (RPKM 32.9) and 20 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

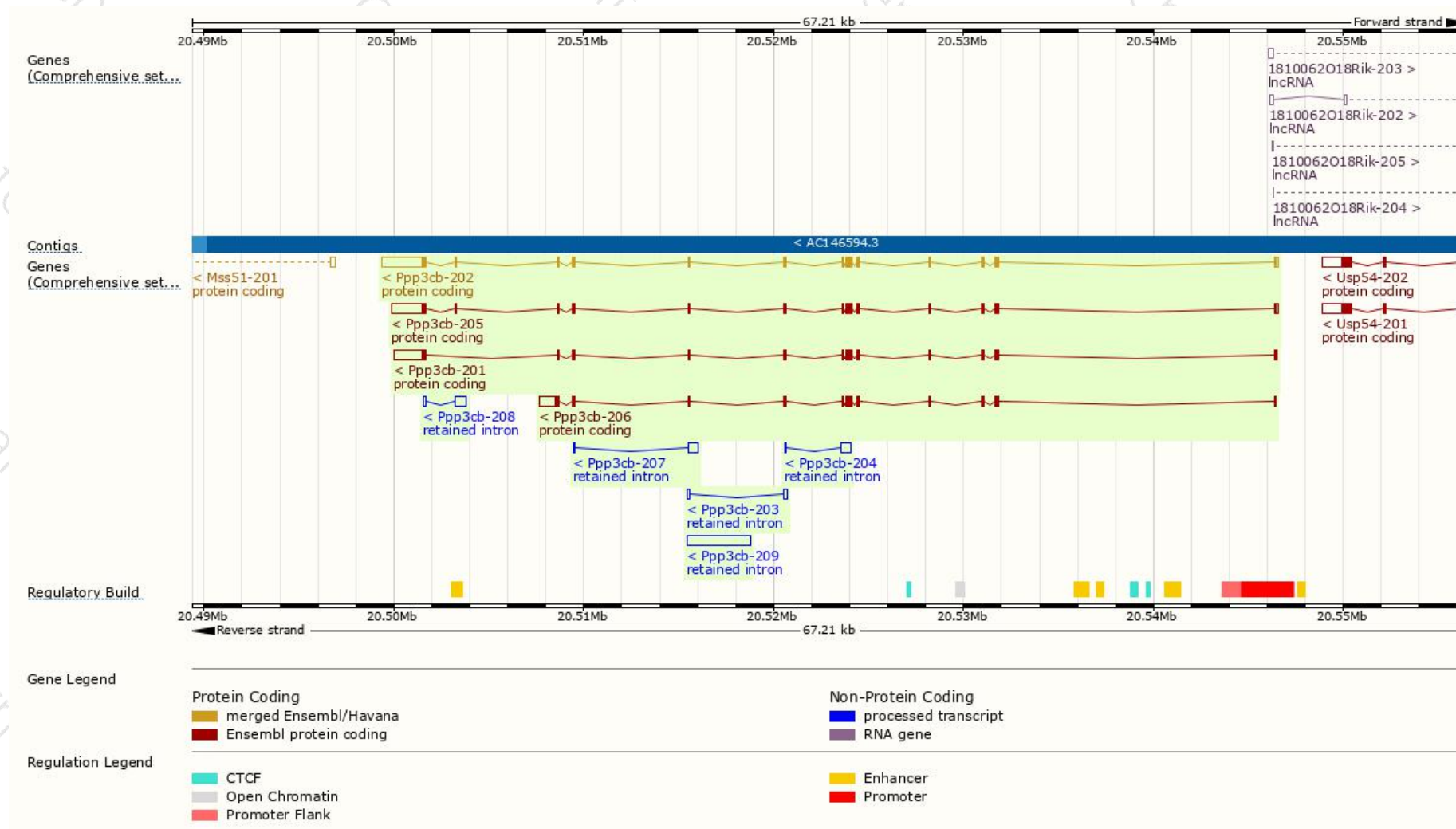
The gene has 9 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ppp3cb-202	<a href="#">ENSMUST00000159027.7</a>	3831	<a href="#">525aa</a>	Protein coding	<a href="#">CCDS26846</a>	<a href="#">P48453</a>	TSL:1 GENCODE basic APPRIS P3
Ppp3cb-205	<a href="#">ENSMUST00000161445.7</a>	3298	<a href="#">524aa</a>	Protein coding	<a href="#">CCDS79266</a>	<a href="#">E0CZ78</a>	TSL:1 GENCODE basic APPRIS ALT1
Ppp3cb-201	<a href="#">ENSMUST00000022355.10</a>	3074	<a href="#">515aa</a>	Protein coding	<a href="#">CCDS79267</a>	<a href="#">G3X8U7</a>	TSL:5 GENCODE basic APPRIS ALT1
Ppp3cb-206	<a href="#">ENSMUST00000161989.1</a>	2322	<a href="#">497aa</a>	Protein coding	-	<a href="#">E9Q6P2</a>	TSL:1 GENCODE basic
Ppp3cb-209	<a href="#">ENSMUST00000223854.1</a>	3360	No protein	Retained intron	-	-	-
Ppp3cb-208	<a href="#">ENSMUST00000162217.1</a>	635	No protein	Retained intron	-	-	TSL:2
Ppp3cb-207	<a href="#">ENSMUST00000162107.1</a>	546	No protein	Retained intron	-	-	TSL:3
Ppp3cb-204	<a href="#">ENSMUST00000161332.1</a>	536	No protein	Retained intron	-	-	TSL:3
Ppp3cb-203	<a href="#">ENSMUST00000160812.1</a>	321	No protein	Retained intron	-	-	TSL:2

The strategy is based on the design of *Ppp3cb-202* transcript,The transcription is shown below

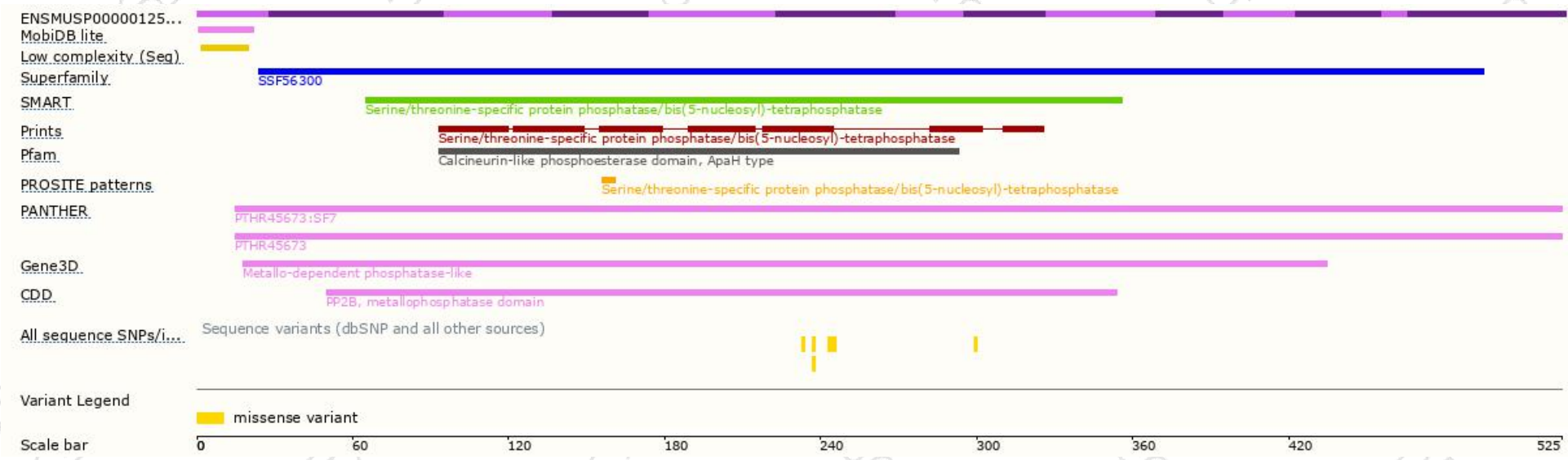


# Genomic location distribution

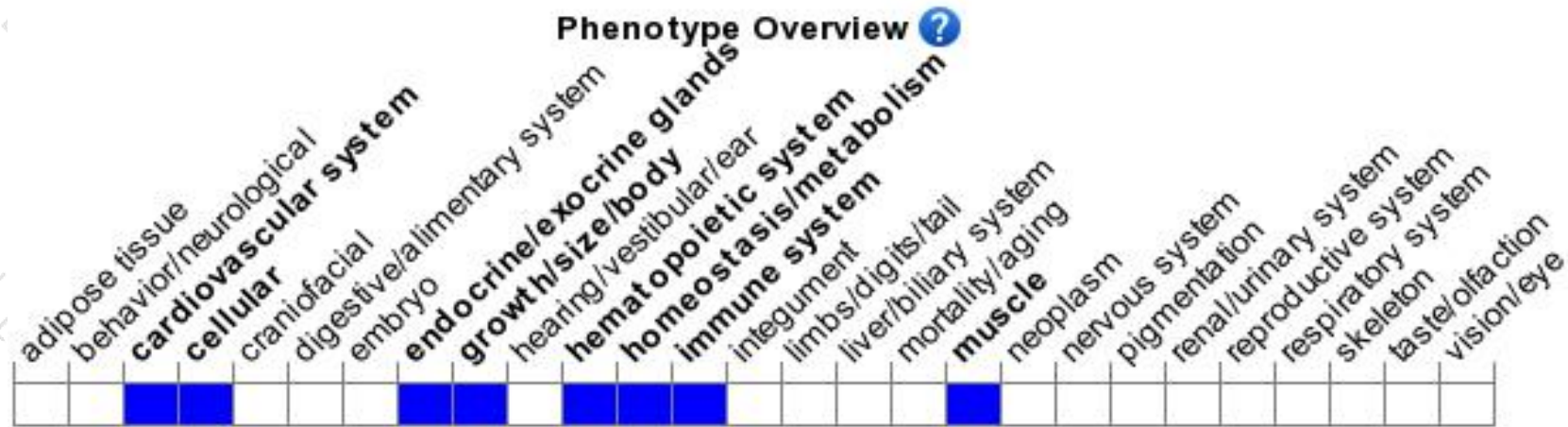




# Protein domain



# Mouse phenotype description(MGI )



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

According to the existing MGI data, Homozygous null mice have small hearts and thymi, and reduced body weight. Cardiac function is normal, but mice lack a cardiac hypertrophic response to pressure overload, angiotensin II, or isopreteronol. Thymi are hypoplastic, with abnormal T cell development and reduced numbers of T cells.

If you have any questions, you are welcome to inquire.

Tel: 400-9660890

