

# *Ptger4* Cas9-KO Strategy

Designer: Yun Li

Reviewer: Longyun Hu

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# Project Overview

**Project Name**

*Ptger4*

**Project type**

**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Ptger4* gene. The schematic diagram is as follows:



- The *Ptger4* gene has 3 transcripts. According to the structure of *Ptger4* gene, exon1-exon3 of *Ptger4-202* (ENSMUST00000120563.1) 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 *Ptger4* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Most homozygous targeted null mutants die shortly after birth due to failed closure of the ductus arteriosis. Survivors show decreased migration of Langerhans cells to lymph nodes, contact hypersensitivity and decreased incidence of induced arthritis.
- The *Ptger4* gene is located on the Chr15. 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)

## Ptger4 prostaglandin E receptor 4 (subtype EP4) [Mus musculus (house mouse)]

Gene ID: 19219, updated on 16-Mar-2019

### Summary



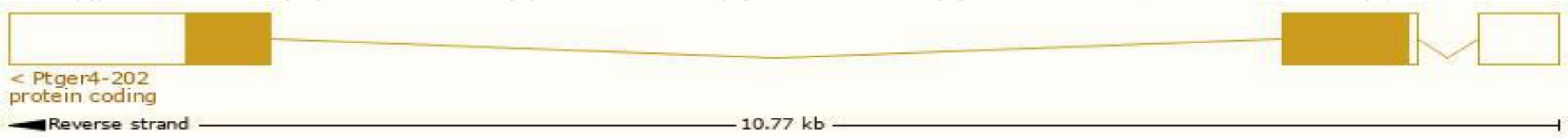
<b>Official Symbol</b>	Ptger4 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	prostaglandin E receptor 4 (subtype EP4) provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:104311</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000039942</a>
<b>Gene type</b>	protein 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>	EP4, Ptgerp4
<b>Expression</b>	Biased expression in small intestine adult (RPKM 27.1), colon adult (RPKM 22.4) and 14 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

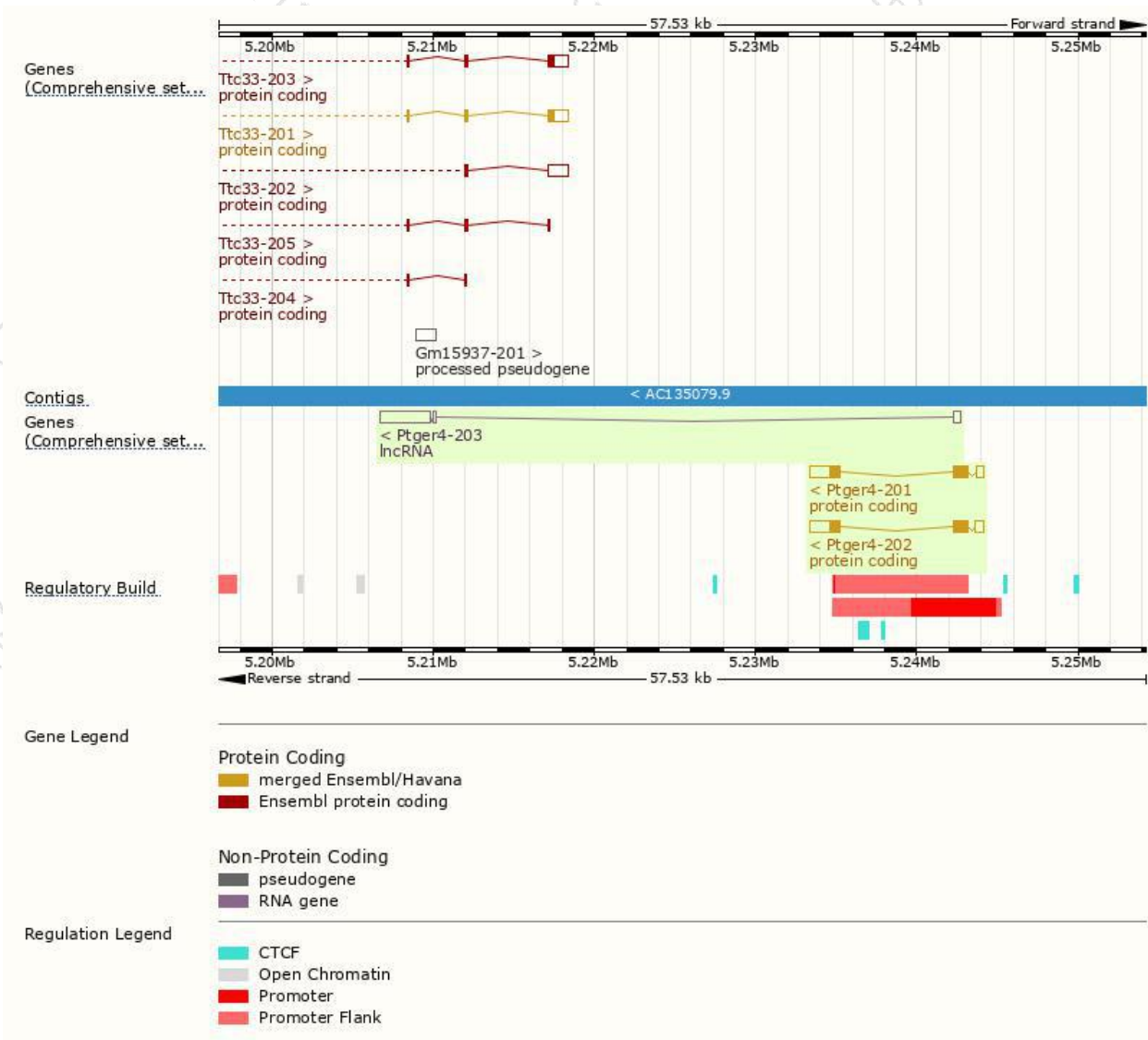
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ptger4-202	<a href="#">ENSMUST00000120563.1</a>	3312	<a href="#">488aa</a>	Protein coding	<a href="#">CCDS49575</a>	<a href="#">Q91VE4</a>	TSL:1 GENCODE basic APPRIS ALT2
Ptger4-201	<a href="#">ENSMUST00000047379.14</a>	3264	<a href="#">513aa</a>	Protein coding	<a href="#">CCDS27365</a>	<a href="#">P32240</a>	TSL:1 GENCODE basic APPRIS P3
Ptger4-203	<a href="#">ENSMUST00000133966.1</a>	3771	No protein	Processed transcript	-	-	TSL:1

The strategy is based on the design of *Ptger4-202* transcript,The transcription is shown below

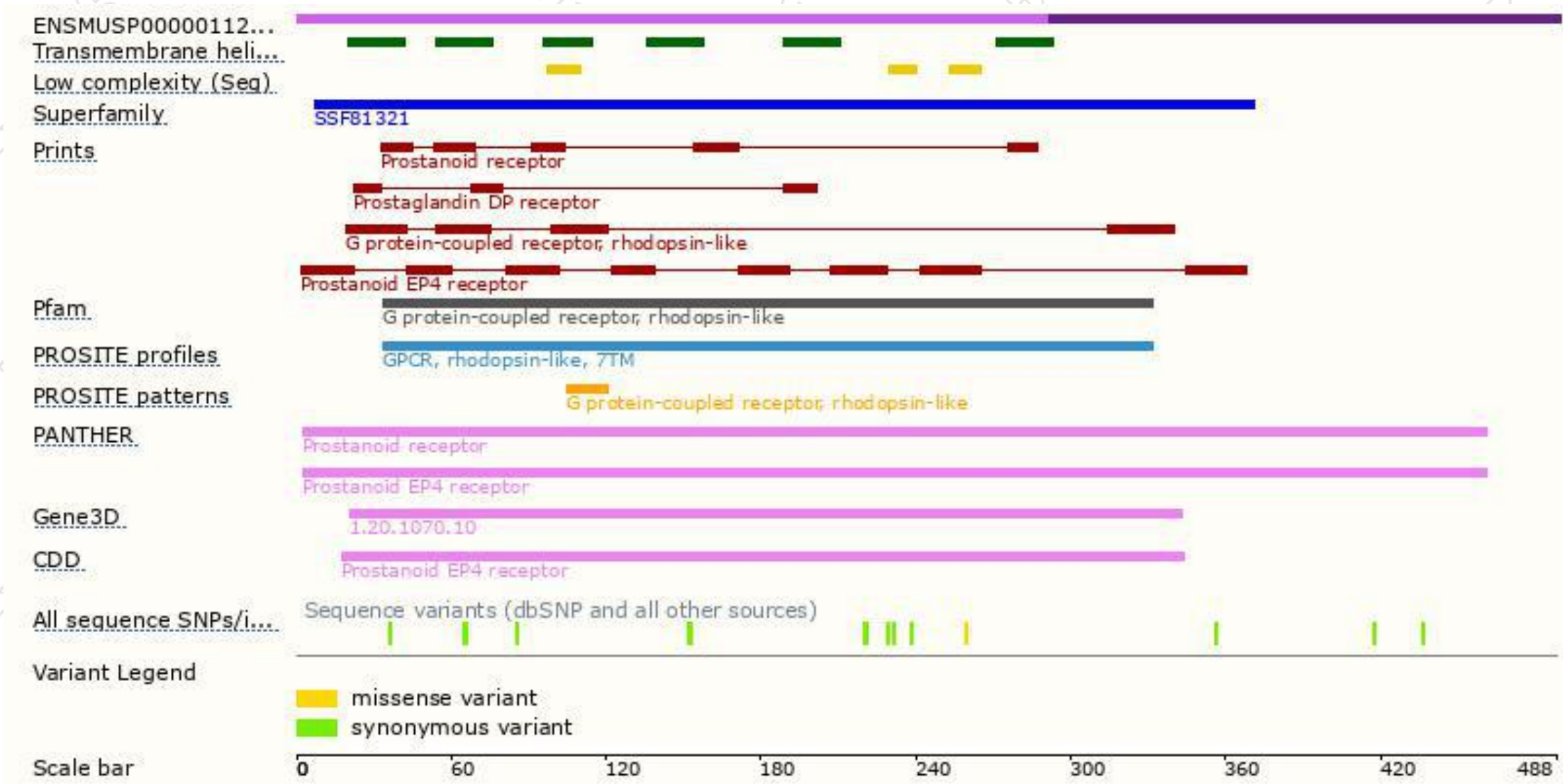


# Genomic location distribution

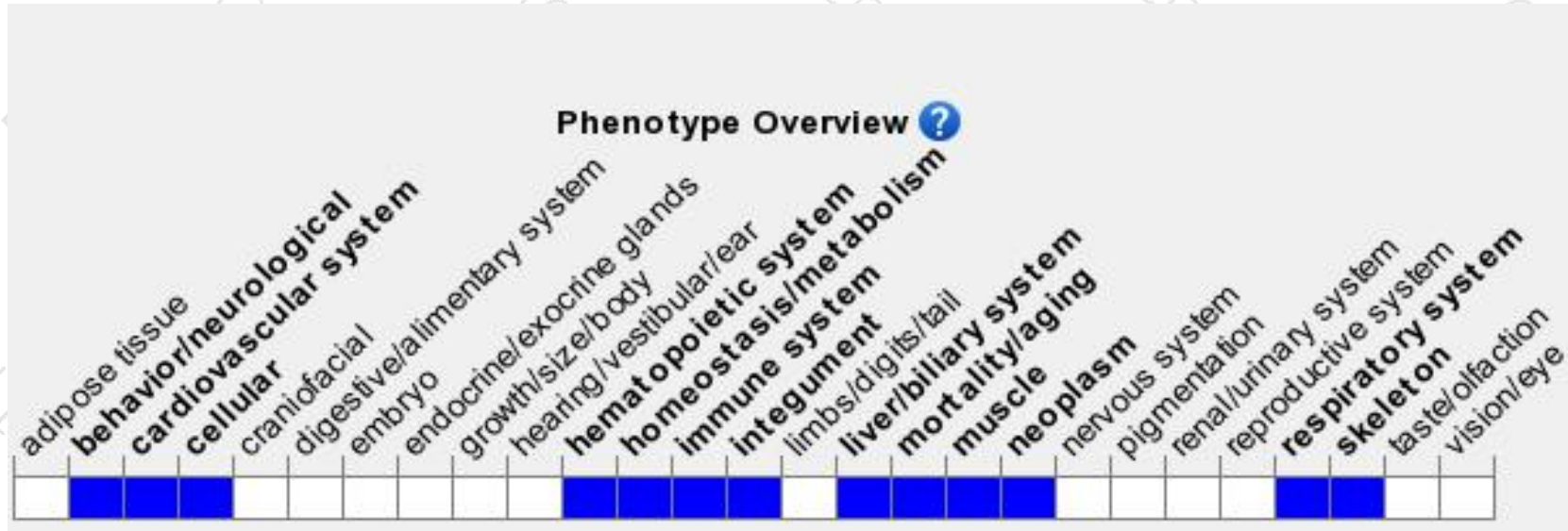




# Protein domain



# 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, Most homozygous targeted null mutants die shortly after birth due to failed closure of the ductus arteriosus. Survivors show decreased migration of Langerhans cells to lymph nodes, contact hypersensitivity and decreased incidence of induced arthritis.

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

Tel: 400-9660890

