

# *Prtg* Cas9-KO Strategy

Designer: Xueting Zhang  
reviewer: Yanhua Shen  
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# Project Overview

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**Project Name**

*Prtg*

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**Project type**

**Cas9-KO**

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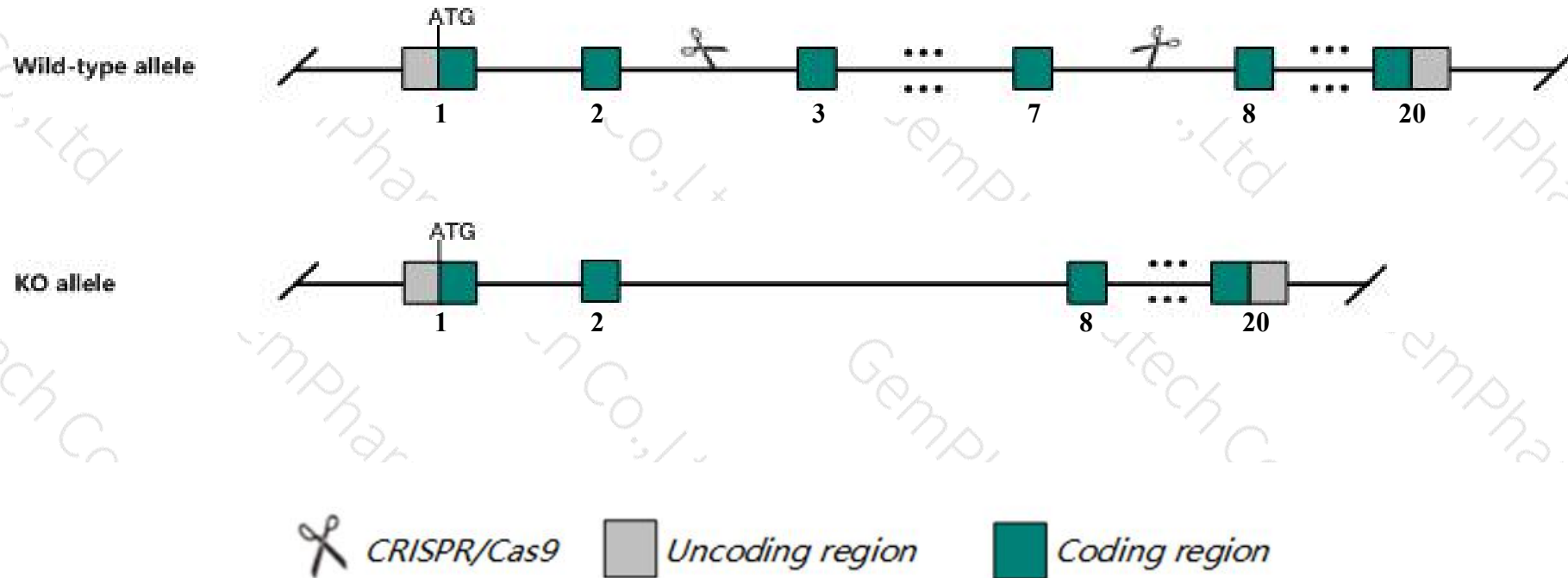
**Strain background**

**C57BL/6JGpt**

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# Knockout strategy

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



- The *Prtg* gene has 2 transcripts. According to the structure of *Prtg* gene, exon3-exon7 of *Prtg*-201 (ENSMUST00000055535.8) transcript is recommended as the knockout region. The region contains 736bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Prtg* gene. The brief process is as follows: CRISPR/Cas9 system w

- Transcript *Prtg*-202 may not be affected.
- The *Prtg* gene is located on the Chr9. 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)

## Prtg protogenin [ *Mus musculus* (house mouse) ]

Gene ID: 235472, updated on 10-Oct-2019

### Summary

- Official Symbol** Prtg provided by [MGI](#)
- Official Full Name** protogenin provided by [MGI](#)
- Primary source** [MGI:MGI:2444710](#)
- See related** [Ensembl:ENSMUSG00000036030](#)
- 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** Igdcc5; A230098A12Rik
- Expression** Biased expression in CNS E11.5 (RPKM 1.3), placenta adult (RPKM 0.8) and 12 other tissues [See more](#)
- Orthologs** [human](#) [all](#)

### Genomic context

**Location:** 9; 9 D See Prtg in [Genome Data Viewer](#)

**Exon count:** 20

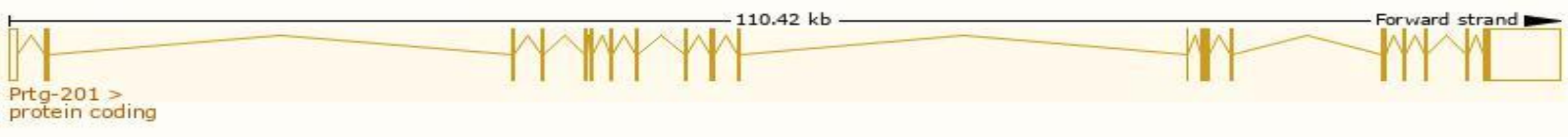
| Annotation release  | Status            | Assembly                                       | Chr | Location                         |
|---------------------|-------------------|------------------------------------------------|-----|----------------------------------|
| <a href="#">108</a> | current           | GRCm38.p6 ( <a href="#">GCF_000001635.26</a> ) | 9   | NC_000075.6 (72807274..72917307) |
| Build 37.2          | previous assembly | MGSCv37 ( <a href="#">GCF_000001635.18</a> )   | 9   | NC_000075.5 (72655081..72765114) |

# Transcript information (Ensembl)

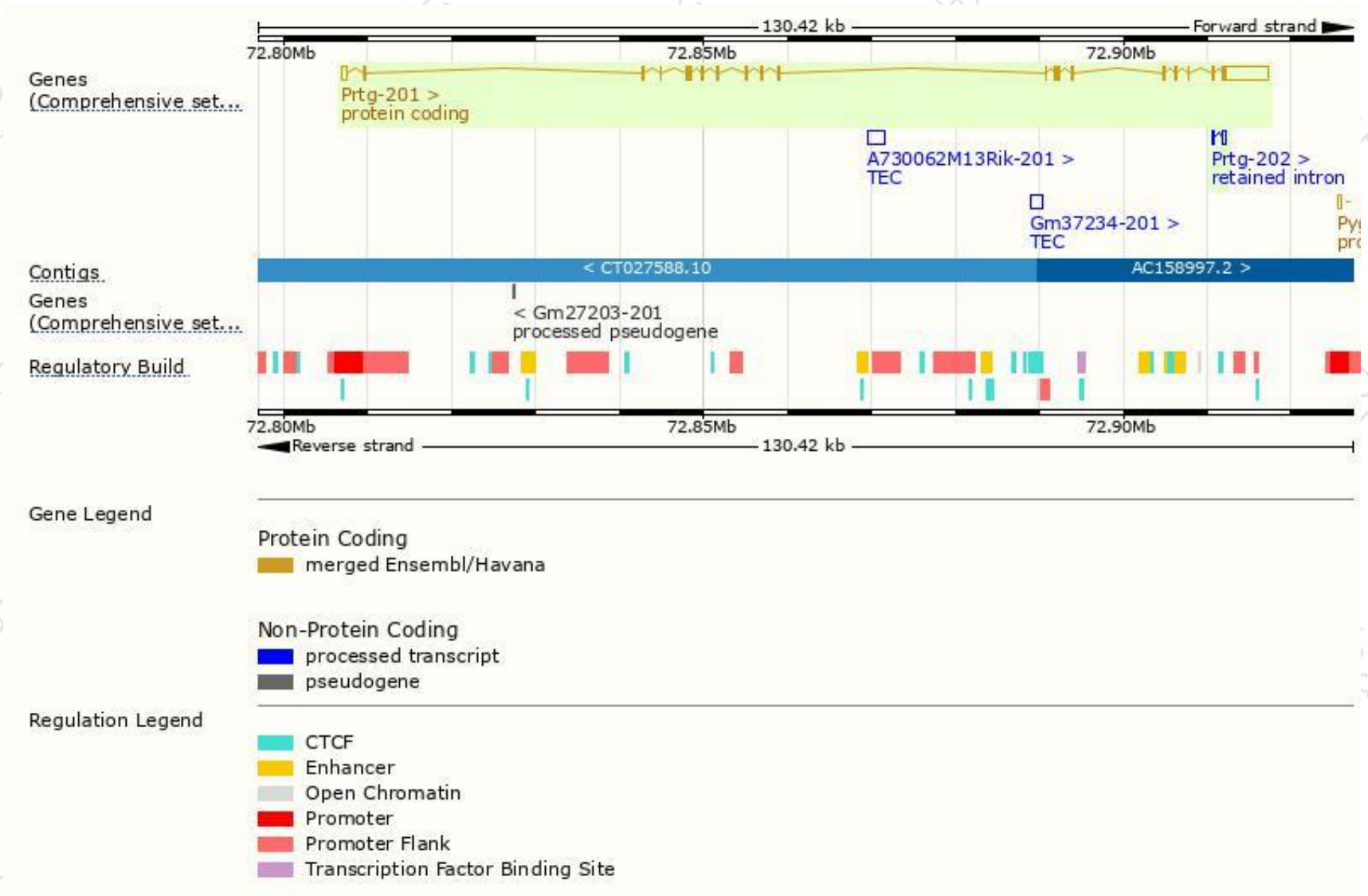
The gene has 2 transcripts,all transcripts are shown below:

| Name     | Transcript ID                        | bp   | Protein                | Biotype         | CCDS                      | UniProt                | Flags                         |
|----------|--------------------------------------|------|------------------------|-----------------|---------------------------|------------------------|-------------------------------|
| Prtg-201 | <a href="#">ENSMUST00000055535.8</a> | 9148 | <a href="#">1191aa</a> | Protein coding  | <a href="#">CCDS23331</a> | <a href="#">Q2EY15</a> | TSL:1 GENCODE basic APPRIS P1 |
| Prtg-202 | <a href="#">ENSMUST00000184274.1</a> | 517  | No protein             | Retained intron | -                         | -                      | TSL:2                         |

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

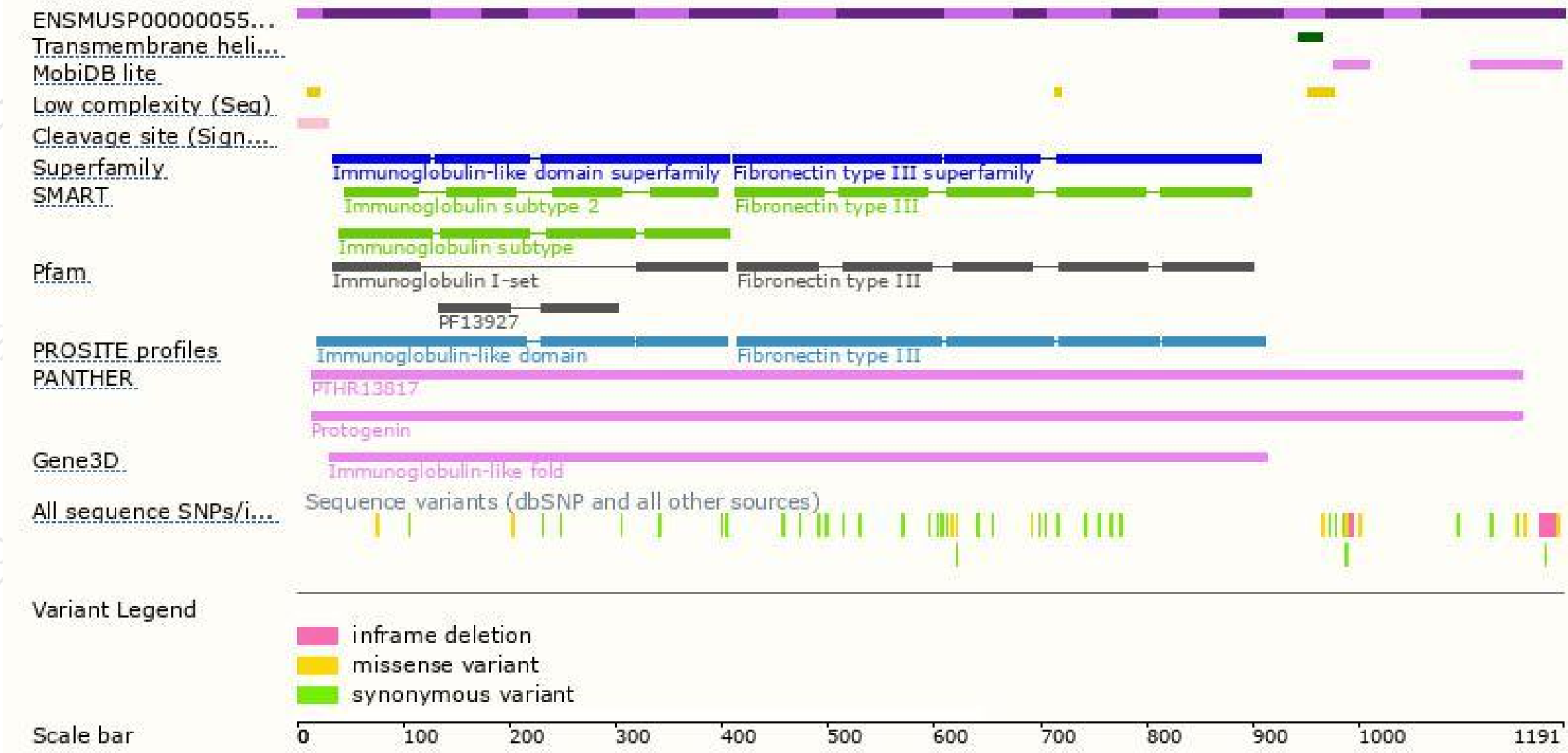


# Genomic location distribution





# Protein domain



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

Tel: 400-9660890

