

Gp5 Cas9-CKO Strategy

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Reviewer:

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Design Date:

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Project Overview

Project Name

Gp5

Project type

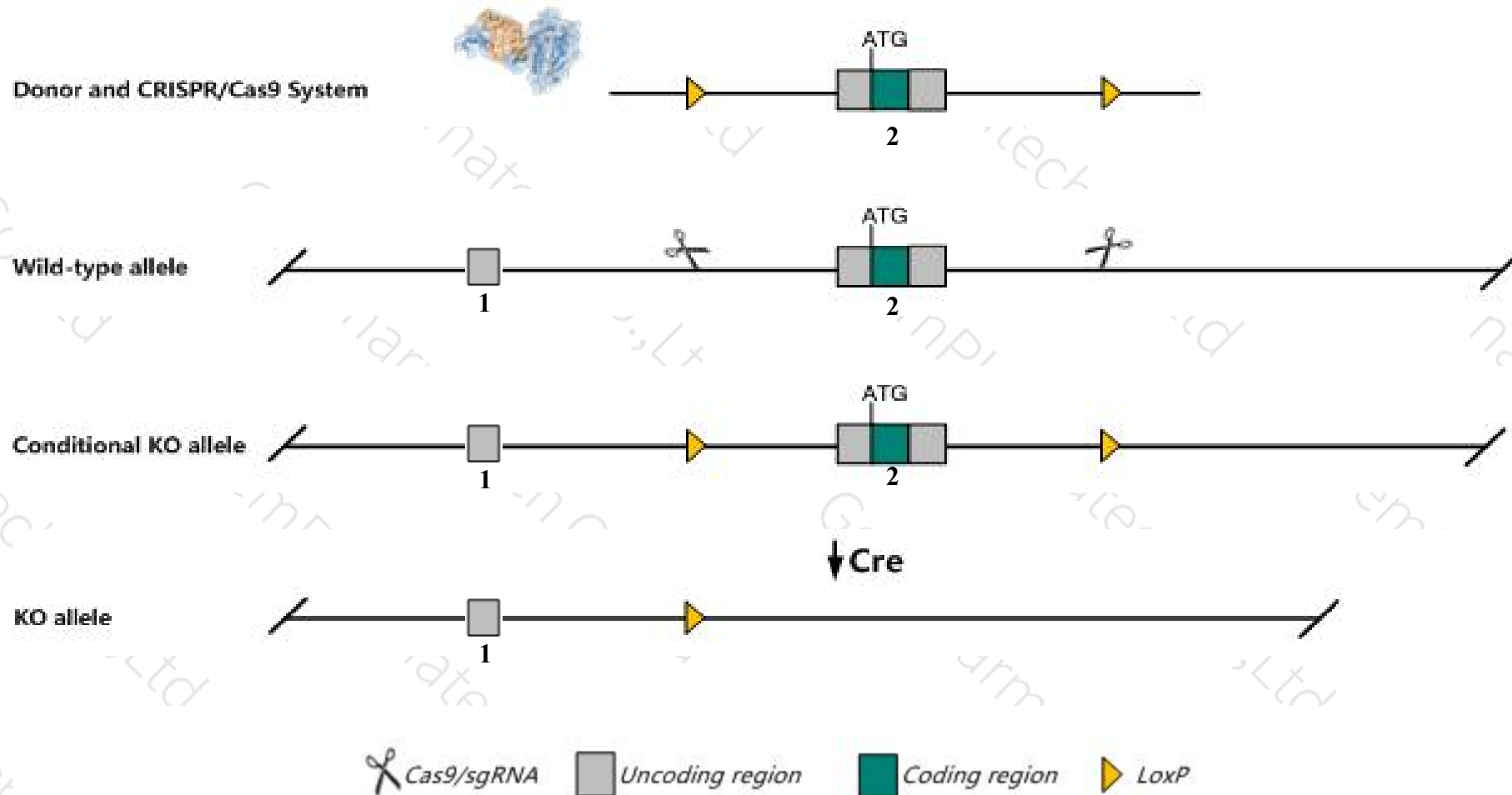
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Gp5* gene. The schematic diagram is as follows:



- The *Gp5* gene has 1 transcript. According to the structure of *Gp5* gene, exon2 of *Gp5-201* (ENSMUST00000061190.7) transcript is recommended as the knockout region. The region contains all 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 *Gp5* 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.

- According to the existing MGI data, Homozygotes for one null allele develop normally with no spontaneous bleeding while their platelets show normal thrombin responsiveness and lack a Bernard-Soulier phenotype. In contrast, homozygotes for a second null allele show a shorter bleeding time and platelet hyperresponsiveness to thrombin.
- The *Gp5* gene is located on the Chr16. 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)

Gp5 glycoprotein 5 (platelet) [*Mus musculus* (house mouse)]

Gene ID: 14729, updated on 12-Aug-2019

Summary

Official Symbol Gp5 provided by [MGI](#)
Official Full Name glycoprotein 5 (platelet) provided by [MGI](#)
Primary source [MGI:MGI:1096363](#)
See related [Ensembl:ENSMUSG00000047953](#)
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 GPV
Expression Biased expression in spleen adult (RPKM 14.8), liver E14.5 (RPKM 6.9) and 5 other tissues [See more](#)
Orthologs [human](#) [all](#)

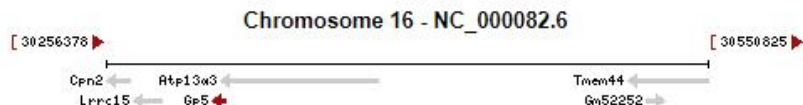
Genomic context

Location: 16; 16 B2

[See Gp5 in Genome Data Viewer](#)

Exon count: 2

Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	16	NC_000082.6 (30307685..30310781, complement)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	16	NC_000082.5 (30307779..30310865, complement)

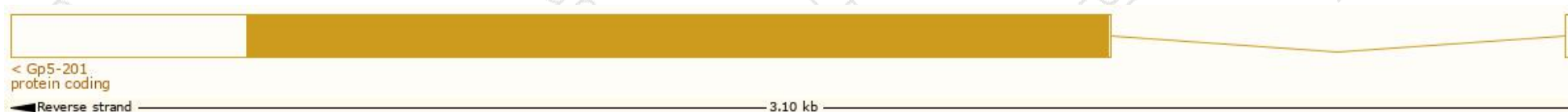


Transcript information (Ensembl)

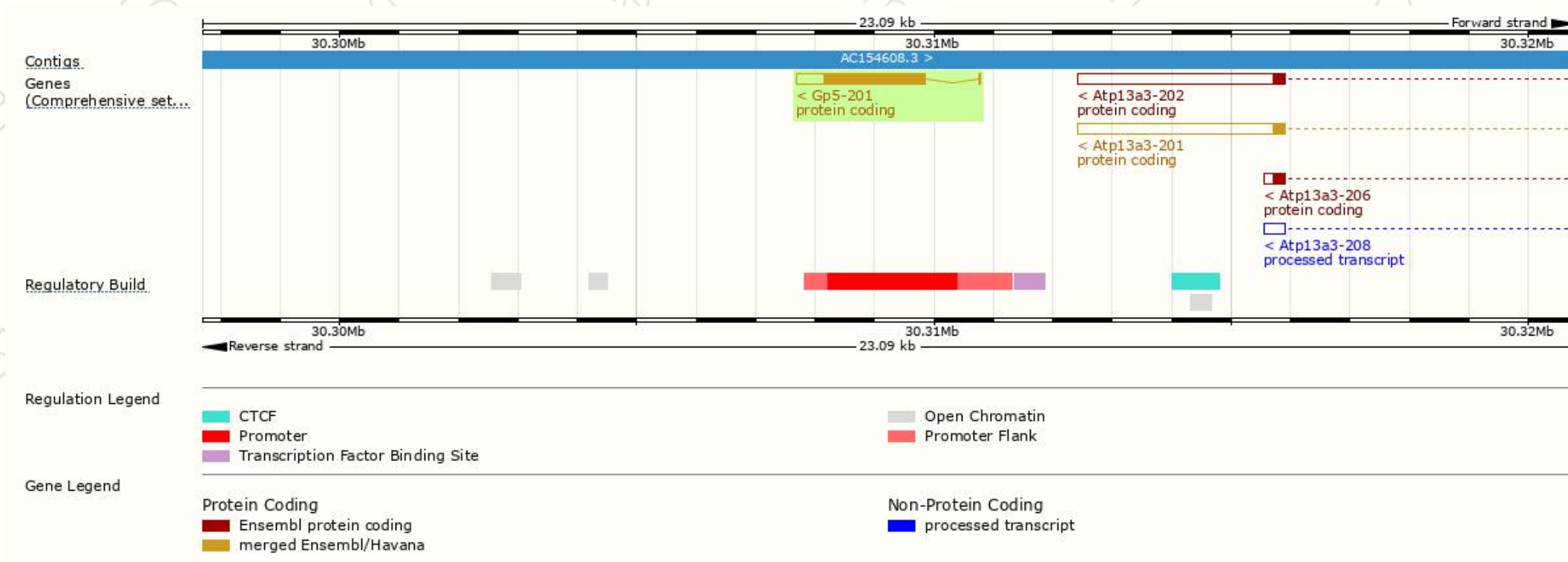
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Gp5-201	ENSMUST00000061190.7	2197	567aa	Protein coding	CCDS28100	Q9QZU3	TSL:1 GENCODE basic APPRIS P1

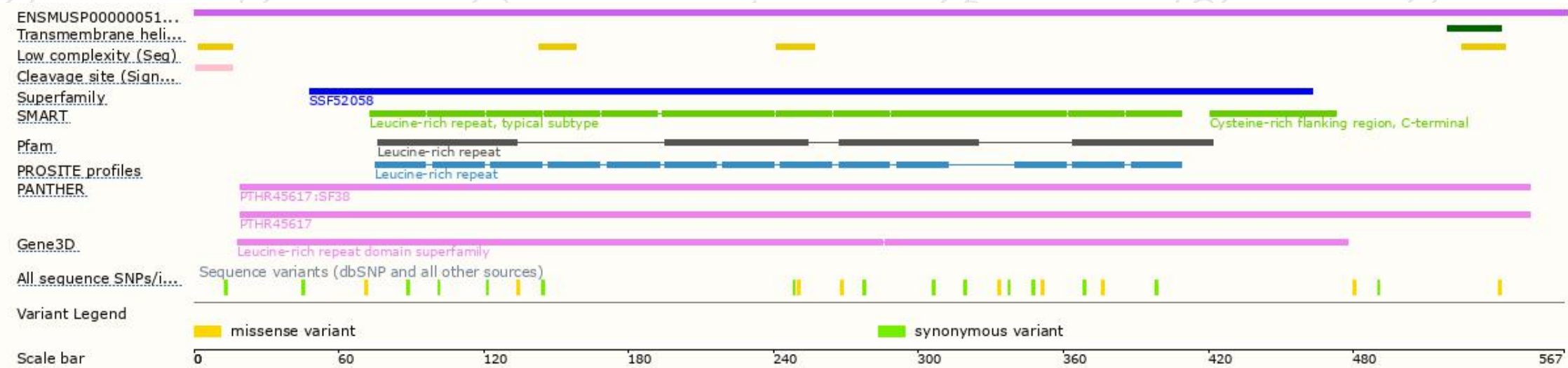
The strategy is based on the design of *Gp5-201* transcript, The transcription is shown below



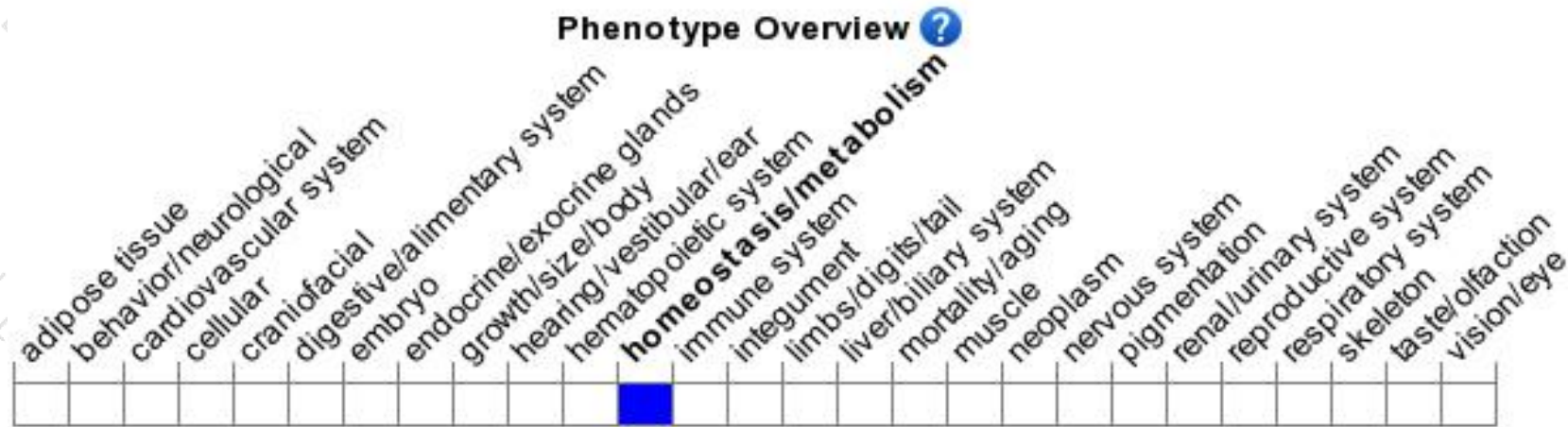
Genomic location distribution



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.

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