

Panx3 Cas9-CKO Strategy

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Reviewer:

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Project Overview

Project Name

Panx3

Project type

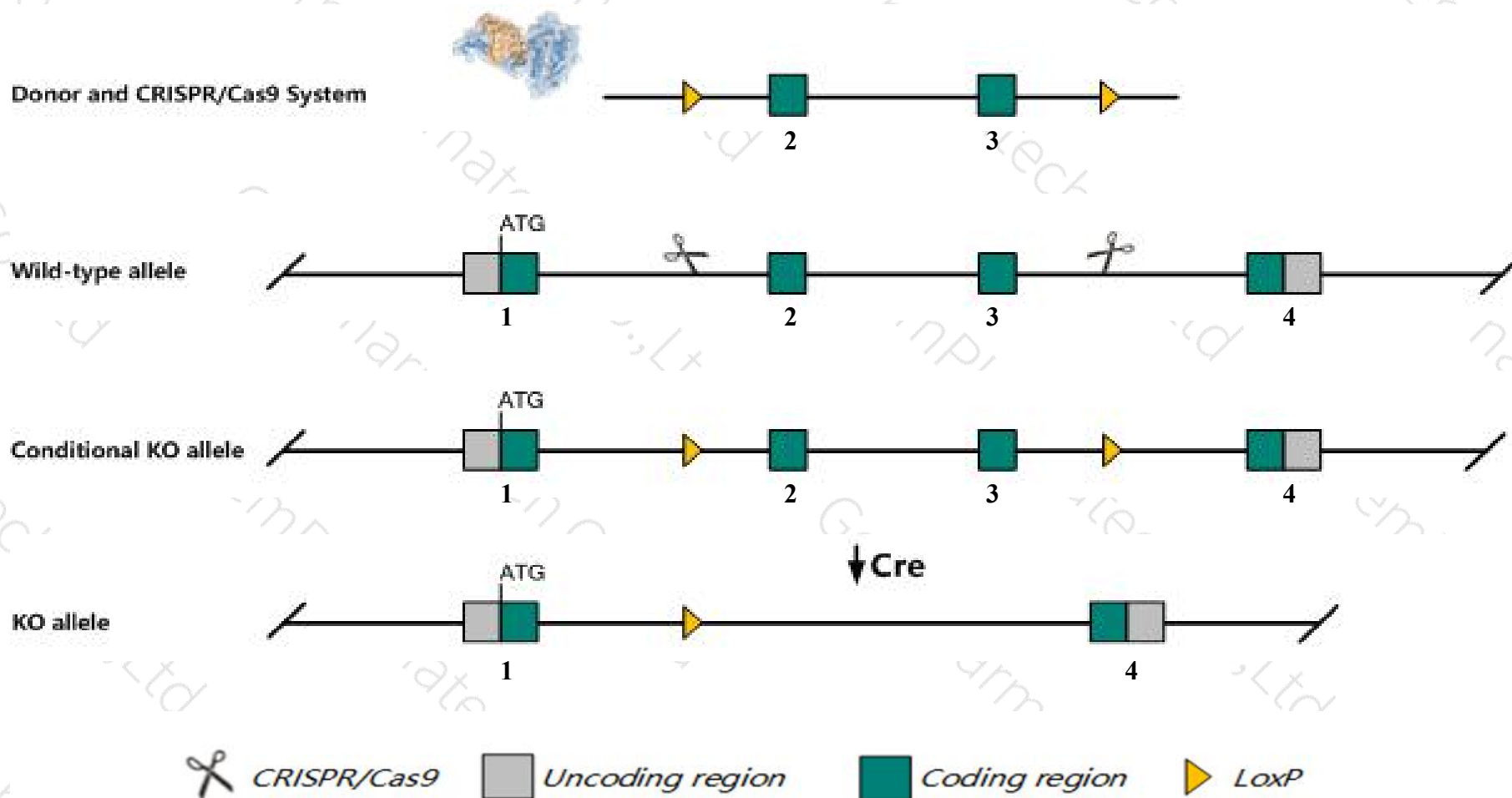
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Panx3* gene. The schematic diagram is as follows:



- The *Panx3* gene has 3 transcripts. According to the structure of *Panx3* gene, exon2-exon3 of *Panx3-201* (ENSMUST00000011262.3) transcript is recommended as the knockout region. The region contains 358bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Panx3* 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, Mice homozygous for a knock-out allele exhibit protection against surgically induced osteoarthritis.
- The *Panx3* 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)

Panx3 pannexin 3 [*Mus musculus* (house mouse)]

Gene ID: 208098, updated on 12-Aug-2019

Summary

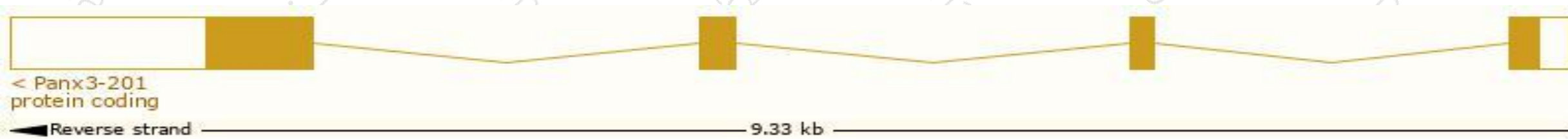
Official Symbol	Panx3 provided by MGI
Official Full Name	pannexin 3 provided by MGI
Primary source	MGI:MGI:1918881
See related	Ensembl:ENSMUSG00000011118
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	3230401P04; 4833413G11Rik
Expression	Biased expression in limb E14.5 (RPKM 14.6) and CNS E14 (RPKM 1.6) See more
Orthologs	human all

Transcript information (Ensembl)

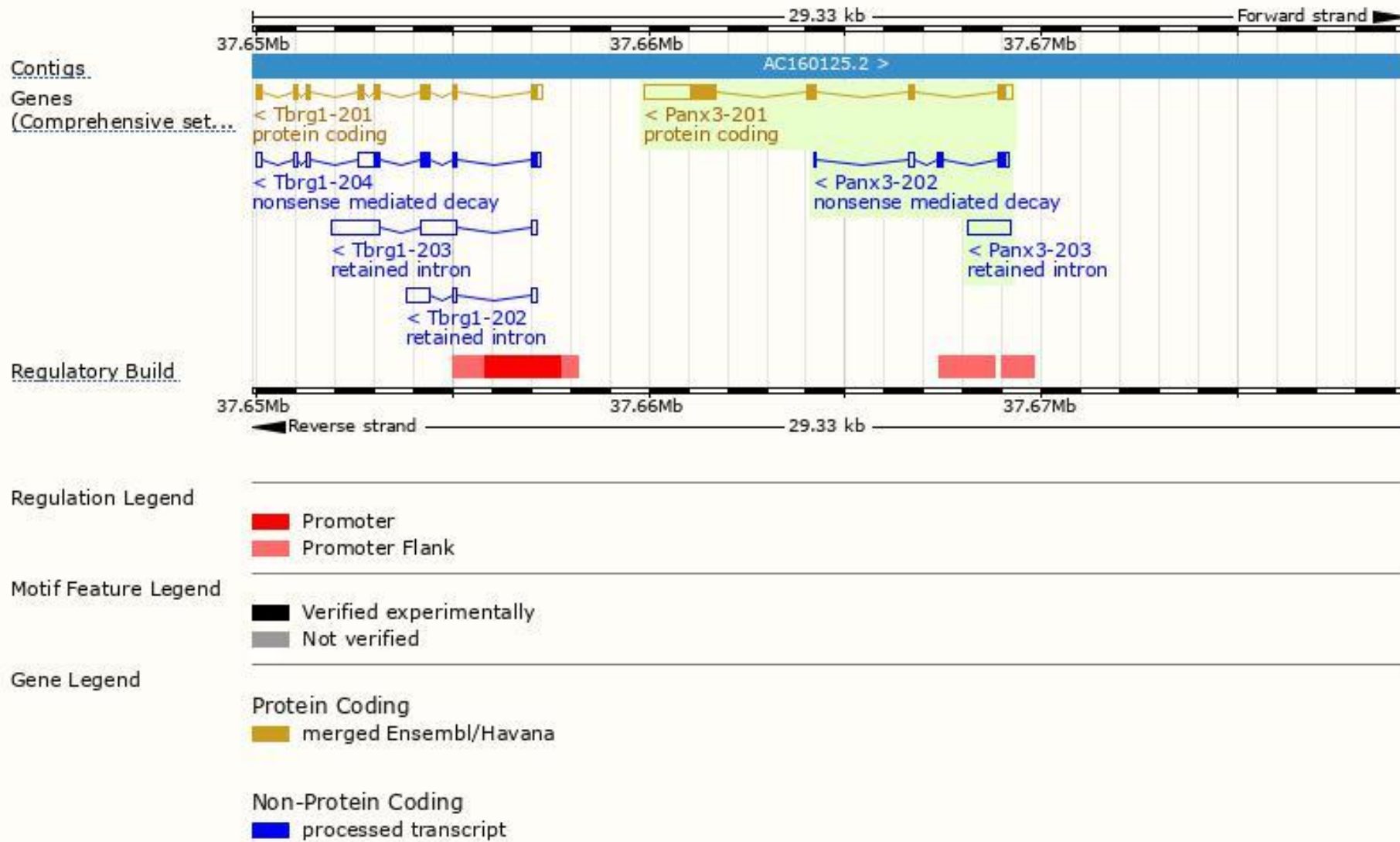
The gene has 3 transcripts, all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Panx3-201	ENSMUST00000011262.3	2531	392aa	Protein coding	CCDS22984	Q6IMP0 Q8CEG0	TSL:1 GENCODE basic APPRIS P1
Panx3-202	ENSMUST00000142228.1	603	97aa	Nonsense mediated decay	-	D6RDS8	TSL:5
Panx3-203	ENSMUST00000215047.1	1078	No protein	Retained intron	-	-	TSL:NA

The strategy is based on the design of *Panx3-201* transcript, The transcription is shown below



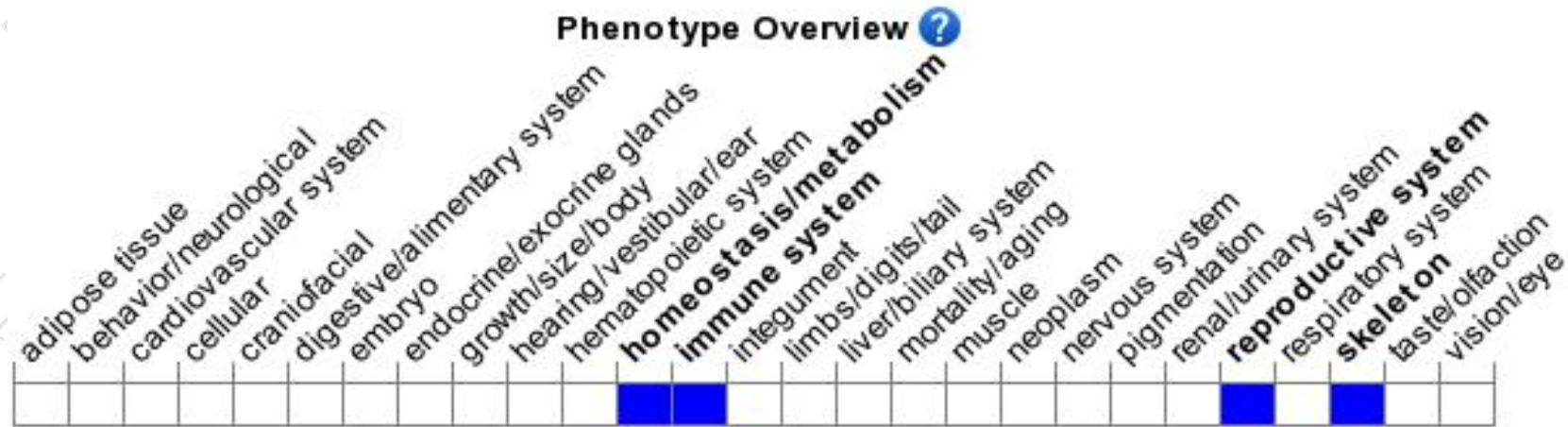
Genomic location distribution



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/>).

According to the existing MGI data, Mice homozygous for a knock-out allele exhibit protection against surgically induced osteoarthritis.

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

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