Polr3g Cas9-CKO Strategy

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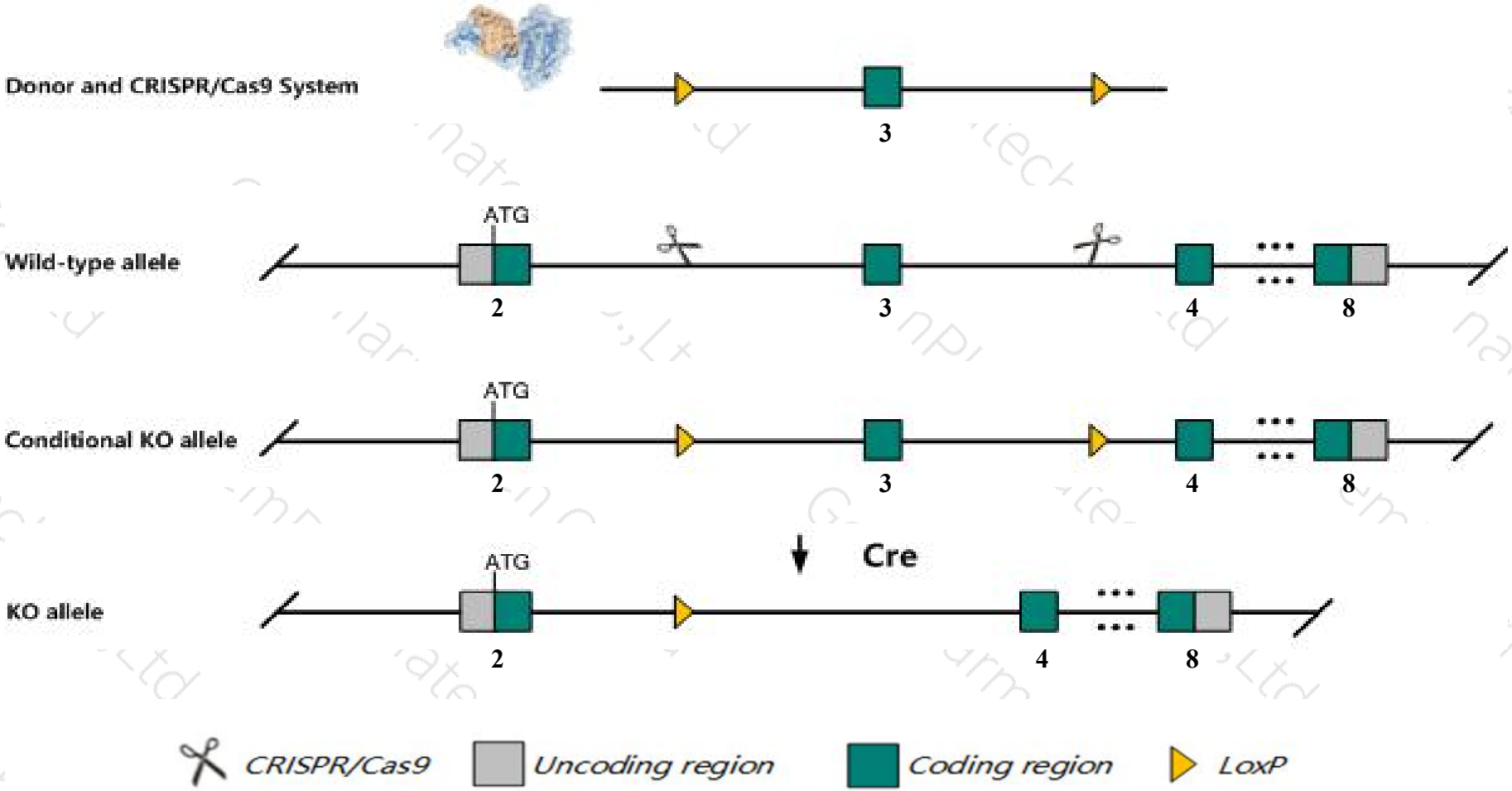
Project Overview

Project Name	***Polr3g***
Project type	**Cas9-CKO**
Strain background	**C57BL/6JGpt**

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Polr3g* gene. The schematic diagram is as follows:



Technical routes

- The *Polr3g* gene has 5 transcripts. According to the structure of *Polr3g* gene, exon3 of *Polr3g-201* (ENSMUST00000048993.11) transcript is recommended as the knockout region.The region contains 130bp coding sequence. Knock out the region will result in disruption of protein function.

- In this project we use CRISPR/Cas9 technology to modify *Polr3g* 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.

Notice

➢ The *Polr3g* gene is located on the Chr13. 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）

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Polr3g polymerase (RNA) III (DNA directed) polypeptide G [Mus musculus (house mouse)]

Gene ID: 67486, updated on 31-Jan-2019

Summary

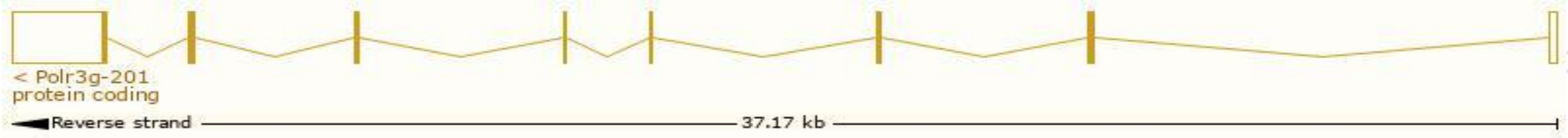
Official Symbol	Polr3g provided by MGI
Official Full Name	polymerase (RNA) III (DNA directed) polypeptide G provided byMGI
Primary source	MGI:MGI:1914736
See related	Ensembl:ENSMUSG00000035834
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	2310047G20Rik, AV275904, RPC32, RPC7
Expression	Ubiquitous expression in placenta adult (RPKM 4.2), CNS E11.5 (RPKM 4.0) and 27 other tissuesSee more
Orthologs	human all

Transcript information（Ensembl）

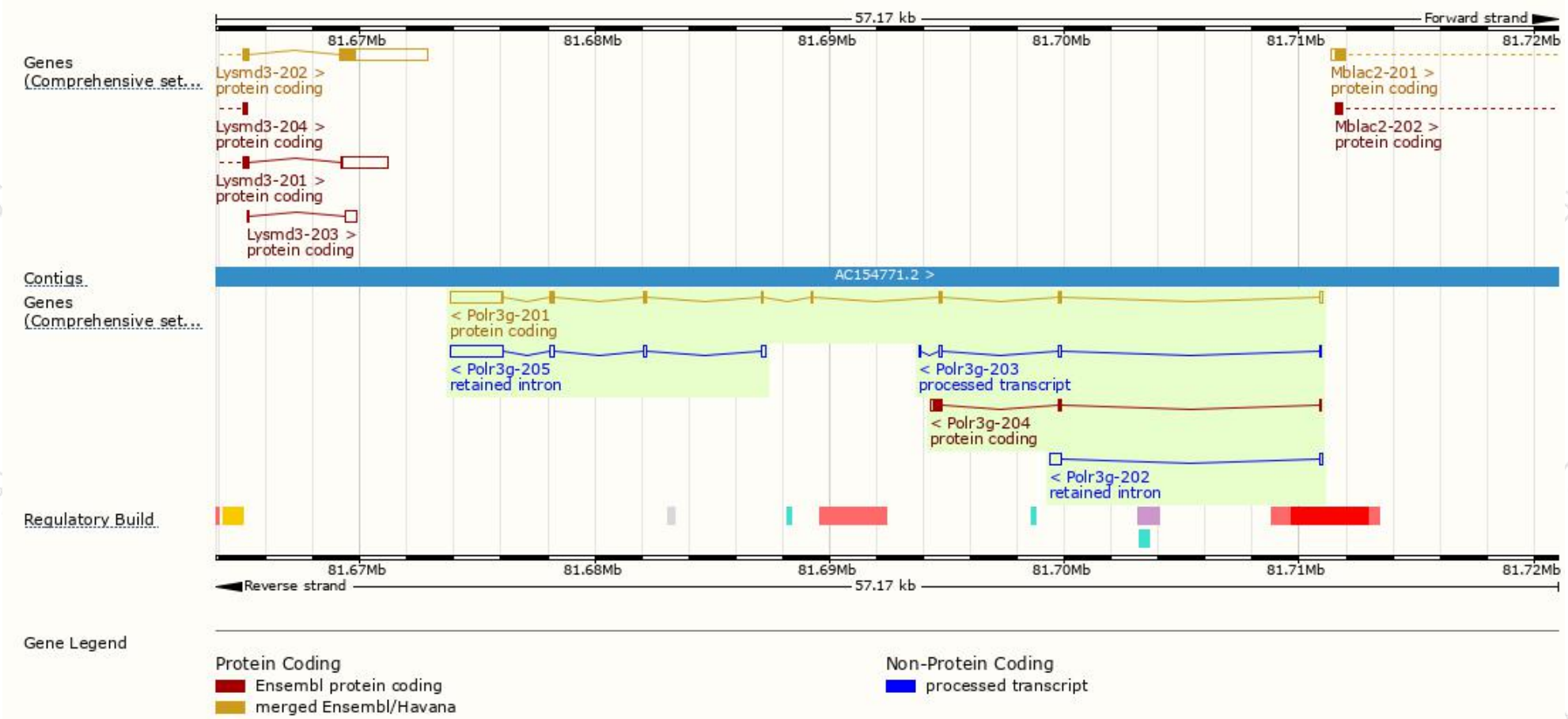
The gene has 5 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Polr3g-201	ENSMUST00000048993.11	3054	223aa	Protein coding	CCDS36738	Q6NXY9	TSL:1 GENCODE basic APPRIS P1
Polr3g-204	ENSMUST00000161920.1	733	160aa	Protein coding	-	Q6NXY9	TSL:1 GENCODE basic
Polr3g-203	ENSMUST00000161093.7	424	No protein	Processed transcript	-	-	TSL:3
Polr3g-205	ENSMUST00000162535.1	2630	No protein	Retained intron	-	-	TSL:1
Polr3g-202	ENSMUST00000160331.1	658	No protein	Retained intron	-	-	TSL:2

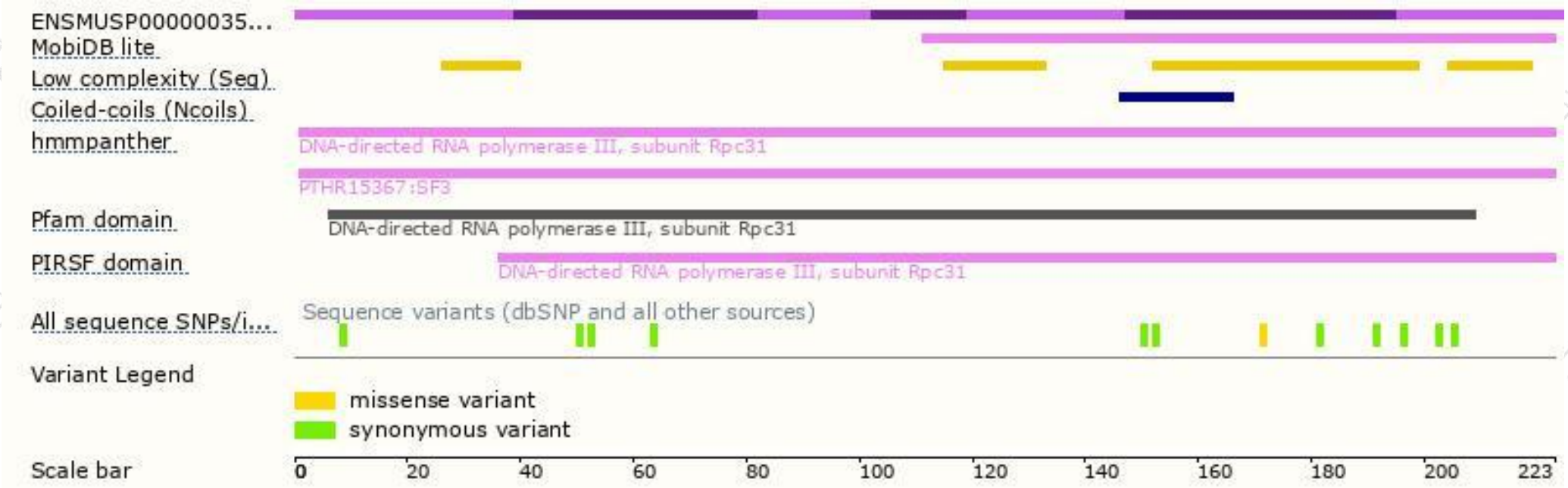
The strategy is based on the design of *Polr3g-201* transcript,The transcription is shown below



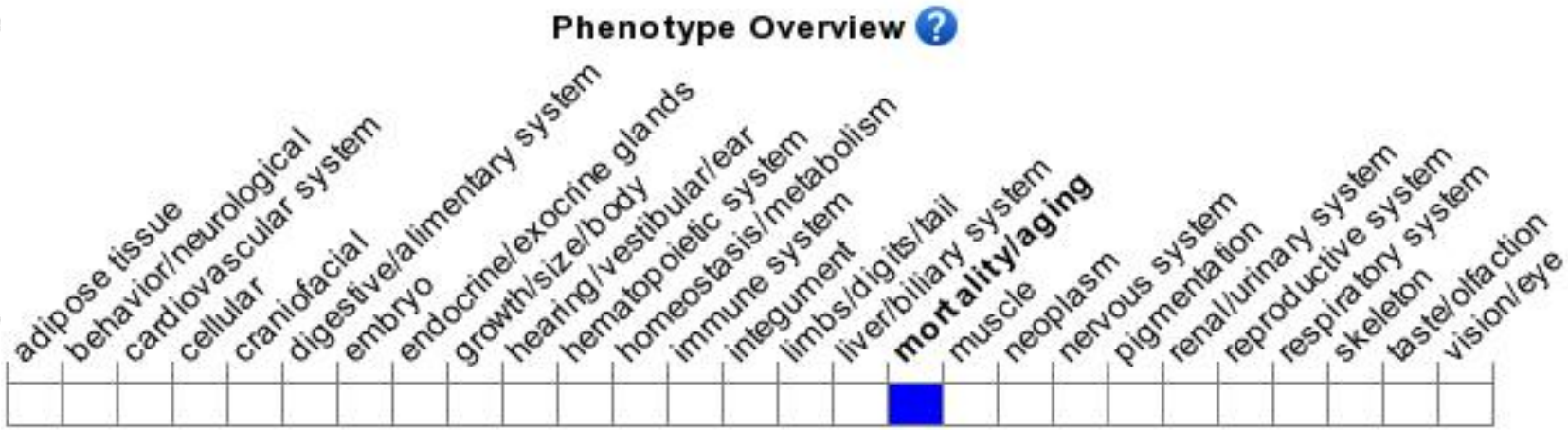
Genomic location distribution



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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