

Polr2c **Cas9-CKO Strategy**

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Project Overview

Project Name

Polr2c

Project type

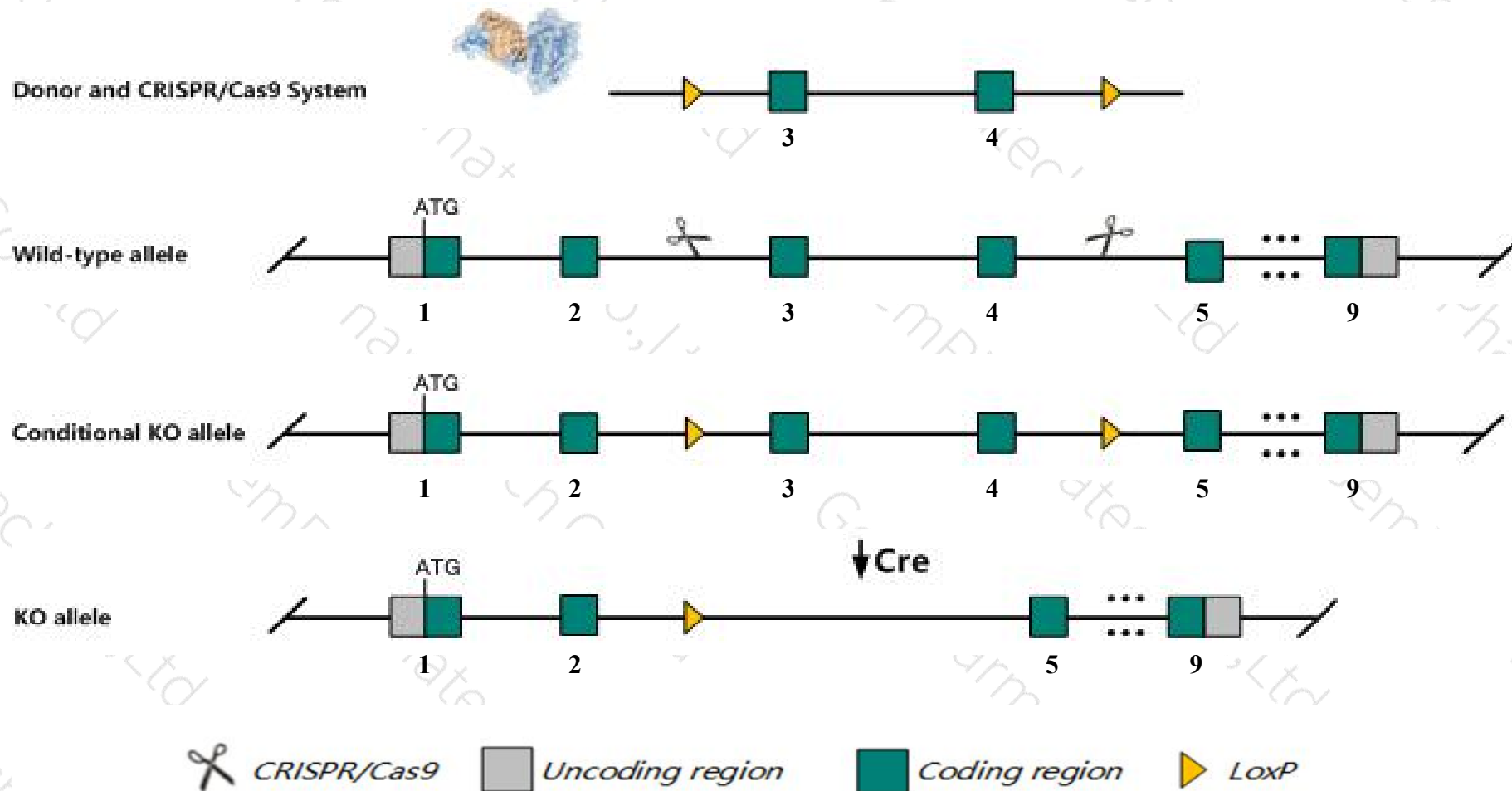
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Polr2c* gene. The schematic diagram is as follows:



- The *Polr2c* gene has 7 transcripts. According to the structure of *Polr2c* gene, exon3-exon4 of *Polr2c*-201 (ENSMUST00000109521.3) transcript is recommended as the knockout region. The region contains 122bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Polr2c* 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 *Polr2c* gene is located on the Chr8. 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)

Polr2c polymerase (RNA) II (DNA directed) polypeptide C [*Mus musculus* (house mouse)]

Gene ID: 20021, updated on 5-Apr-2020

Summary



Official Symbol	Polr2c provided by MGI
Official Full Name	polymerase (RNA) II (DNA directed) polypeptide C provided by MGI
Primary source	MGI:MGI:109299
See related	Ensembl:ENSMUSG00000031783
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	RPB3; 33kDa; Rpo2-3; mRBP31
Expression	Ubiquitous expression in kidney adult (RPKM 36.8), CNS E14 (RPKM 33.0) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

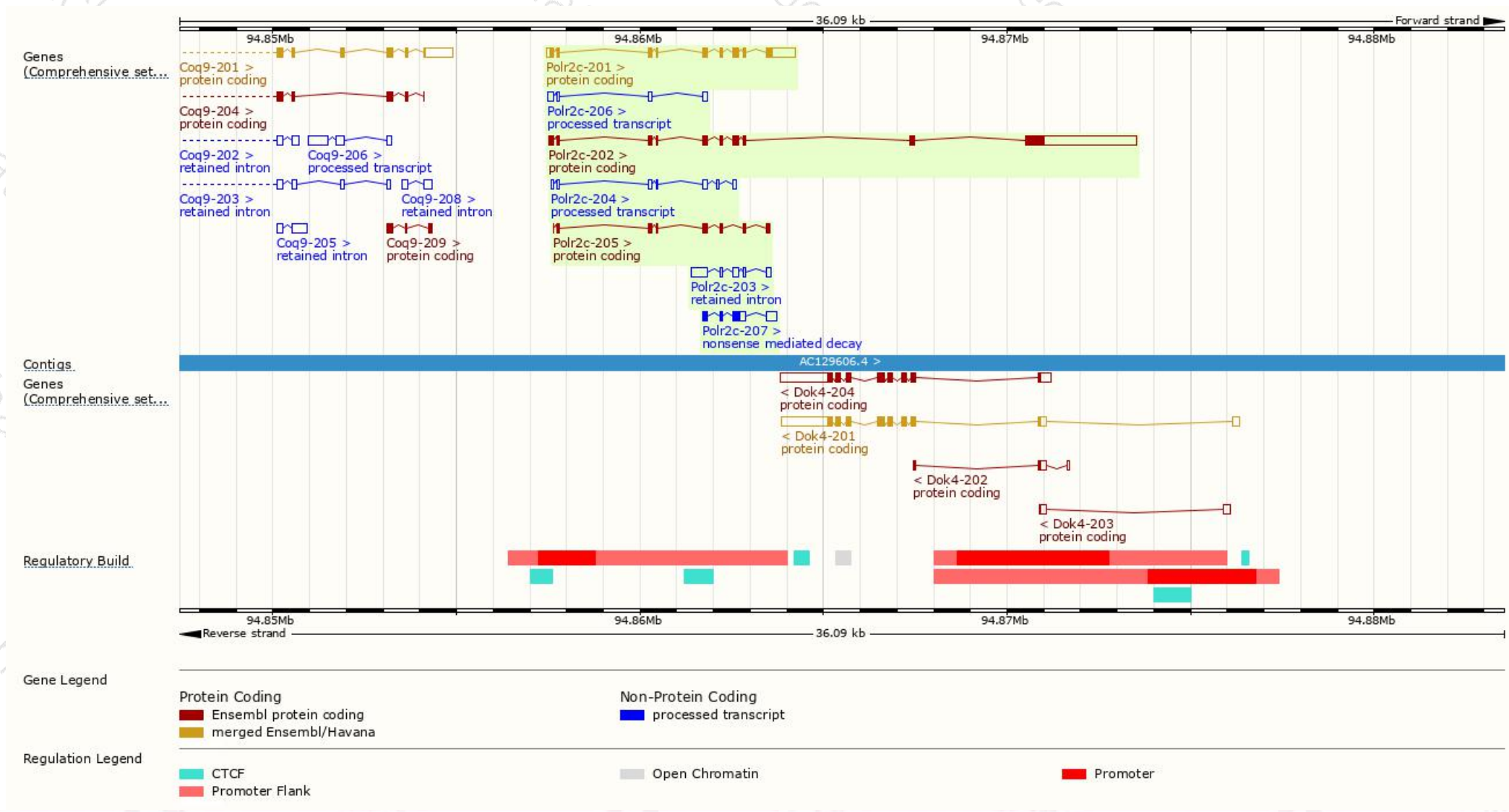
The gene has 7 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Polr2c-201	ENSMUST00000109521.3	1580	275aa	Protein coding	CCDS40442	P97760	TSL:1 GENCODE basic APPRIS P1
Polr2c-202	ENSMUST00000211939.1	3881	439aa	Protein coding	-	A0A1D5RML8	TSL:2 GENCODE basic
Polr2c-205	ENSMUST00000212124.1	528	170aa	Protein coding	-	A0A1D5RLD4	CDS 5' incomplete TSL:3
Polr2c-207	ENSMUST00000212848.1	830	128aa	Nonsense mediated decay	-	A0A1D5RLM9	CDS 5' incomplete TSL:3
Polr2c-204	ENSMUST00000212120.1	530	No protein	Processed transcript	-	-	TSL:3
Polr2c-206	ENSMUST00000212744.1	376	No protein	Processed transcript	-	-	TSL:3
Polr2c-203	ENSMUST00000211957.1	855	No protein	Retained intron	-	-	TSL:2

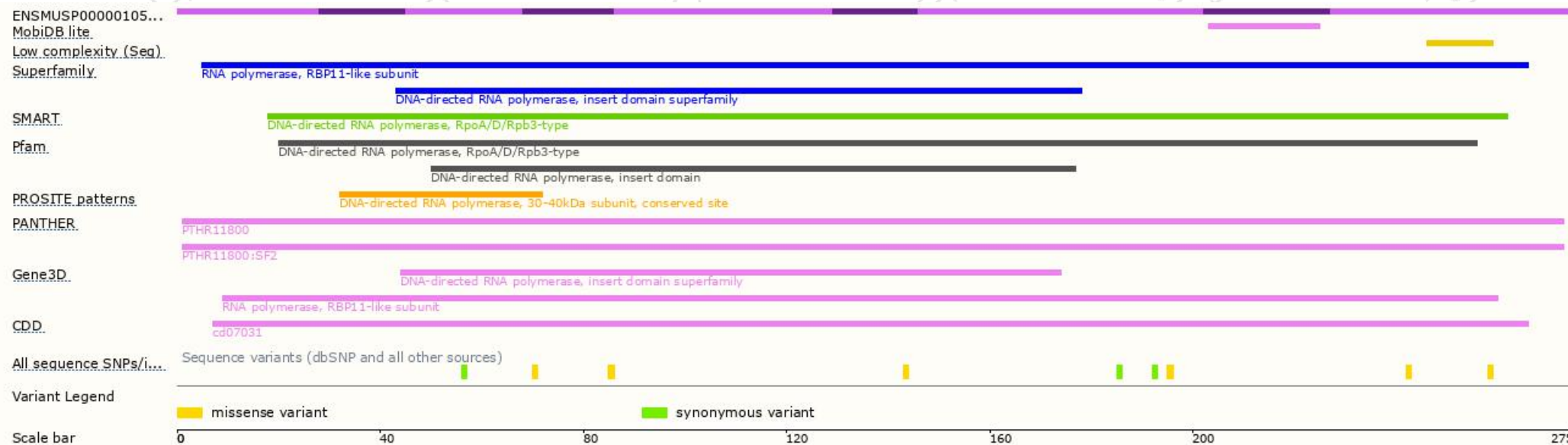
The strategy is based on the design of *Polr2c-201* transcript, the transcription is shown below



Genomic location distribution



Protein domain



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

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