

Ypel5 Cas9-CKO Strategy

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Project Overview

Project Name

Ypel5

Project type

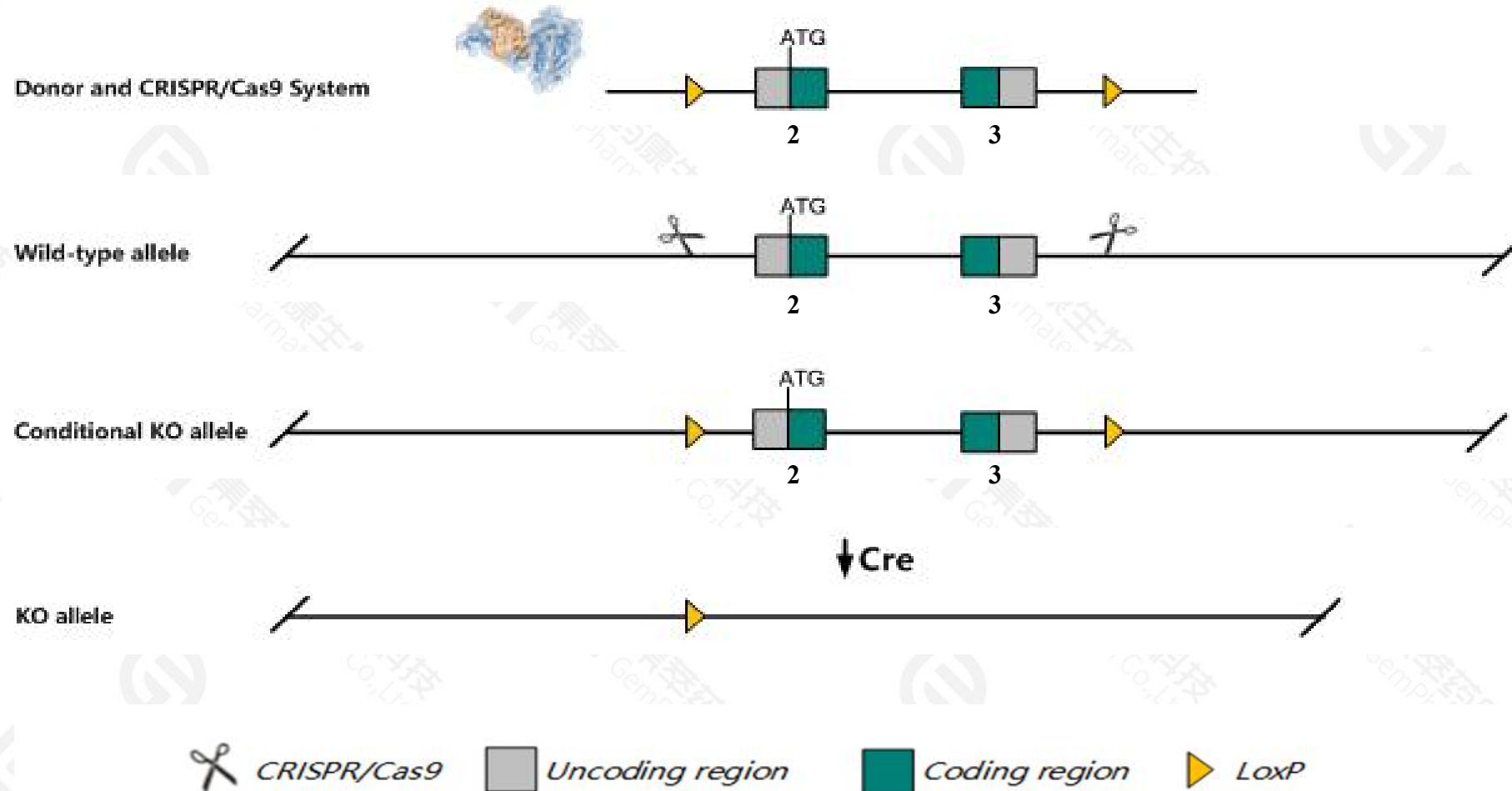
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Ypel5* gene. The schematic diagram is as follows:



- The *Ypel5* gene has 5 transcripts. According to the structure of *Ypel5* gene, exon2-exon3 of *Ypel5*-205(ENSMUST00000233886.2) 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 *Ypel5* 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 *Ypel5* gene is located on the Chr17. 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)

Ypel5 yippee like 5 [Mus musculus (house mouse)]

Gene ID: 383295, updated on 12-Feb-2021

Summary



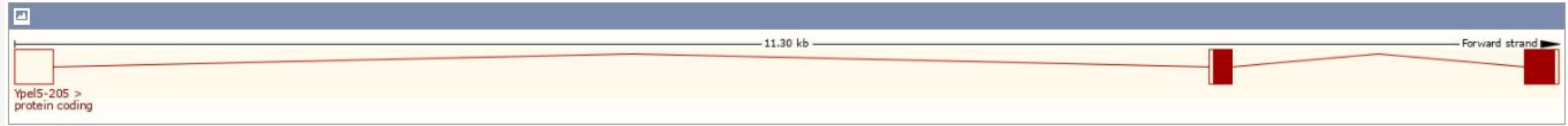
Official Symbol	Ypel5 provided by MGI
Official Full Name	yippee like 5 provided by MGI
Primary source	MGI:MGI:1916937
See related	Ensembl:ENSMUSG00000039770
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	2310076K21Rik, CGI-12, CGI-127
Expression	Ubiquitous expression in frontal lobe adult (RPKM 16.8), cerebellum adult (RPKM 15.5) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

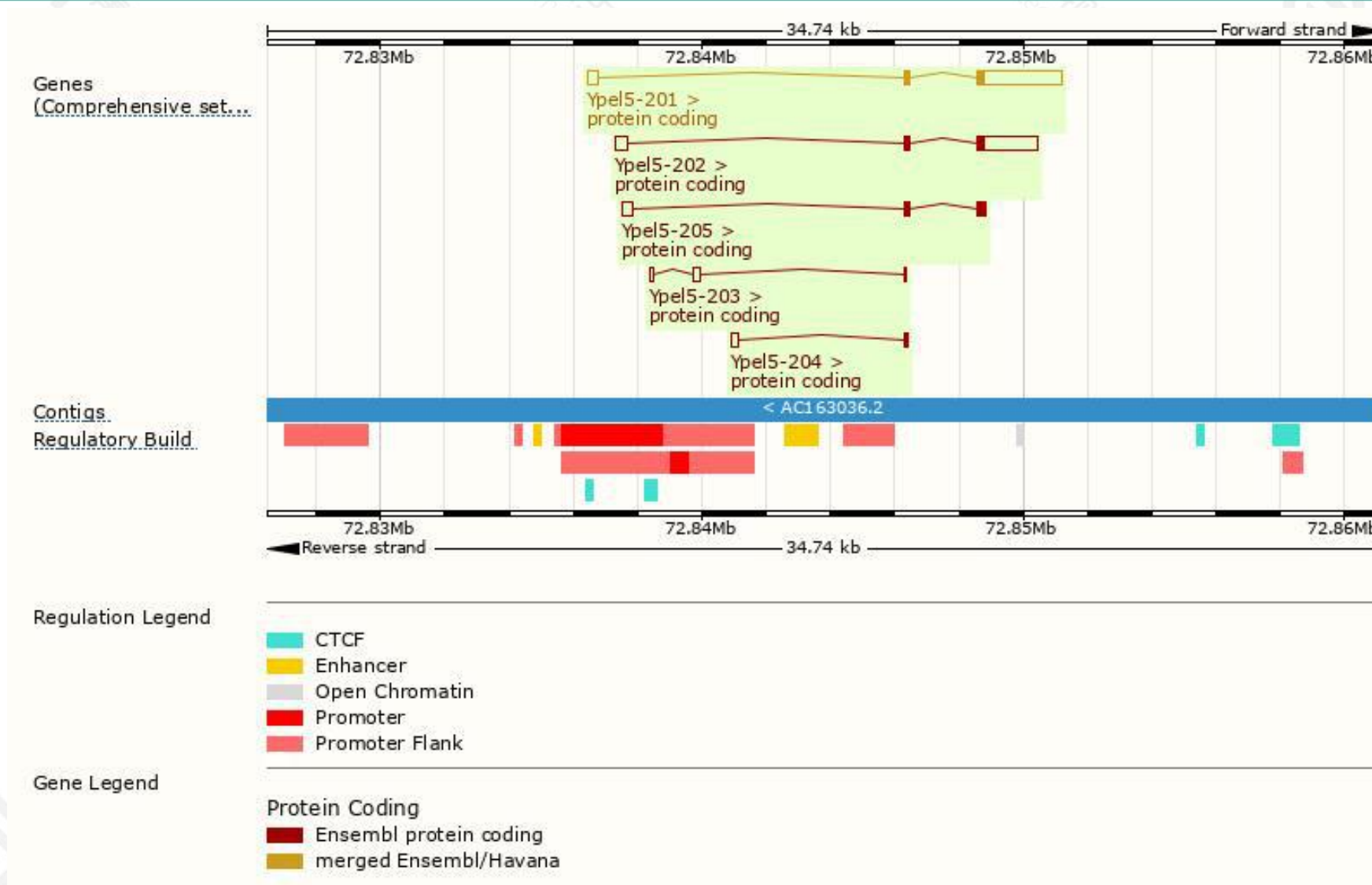
The gene has 5 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ypel5-201	ENSMUST00000045174.7	3120	121aa	Protein coding	CCDS28963		TSL:1 , GENCODE basic , APPRIS P1 ,
Ypel5-202	ENSMUST00000233210.2	2419	121aa	Protein coding	CCDS28963		GENCODE basic , APPRIS P1 ,
Ypel5-205	ENSMUST00000233886.2	704	121aa	Protein coding	CCDS28963		GENCODE basic , APPRIS P1 ,
Ypel5-203	ENSMUST00000233221.2	422	13aa	Protein coding	-		CDS 3' incomplete ,
Ypel5-204	ENSMUST00000233662.2	306	27aa	Protein coding	-		CDS 3' incomplete ,

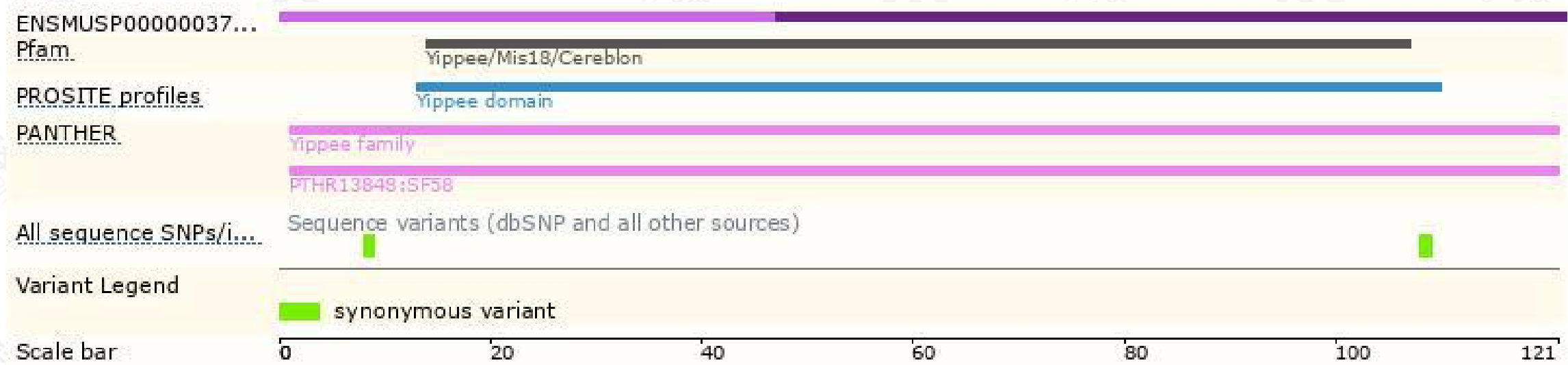
The strategy is based on the design of *Ypel5-205* transcript,the transcription is shown below:



Genomic location distribution



Protein domain



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