

# *Hepacam2* Cas9-CKO Strategy

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# Project Overview

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**Project Name**

*Hepacam2*

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**Project type**

**Cas9-CKO**

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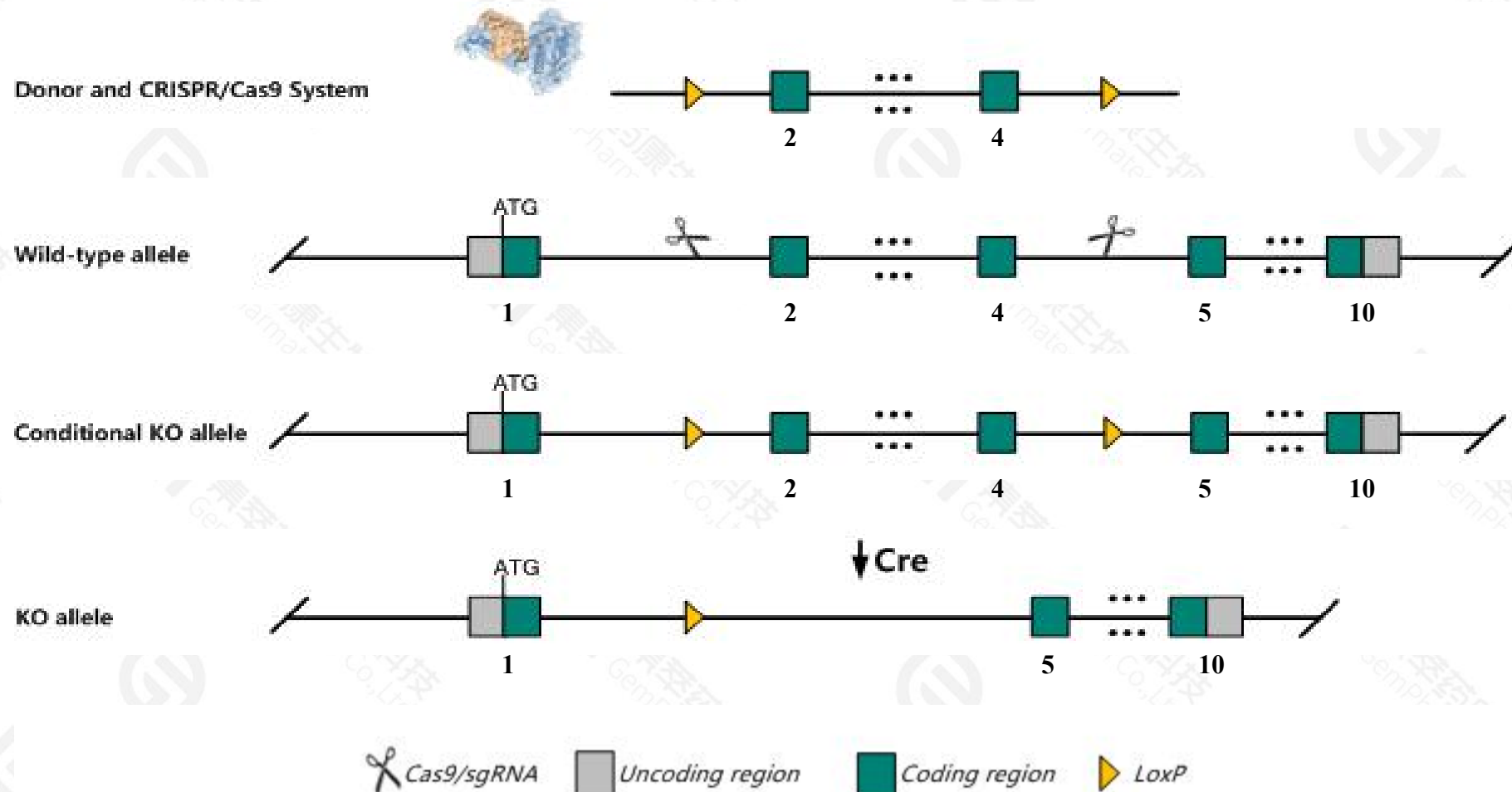
**Strain background**

**C57BL/6JGpt**

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# Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Hepacam2* gene. The schematic diagram is as follows:



- The *Hepacam2* gene has 5 transcripts. According to the structure of *Hepacam2* gene, exon2-exon4 of *Hepacam2*-201(ENSMUST00000049985.14) transcript is recommended as the knockout region. The region contains most 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 *Hepacam2* gene. The brief process is as follows: sgRNA was transcribed in vitro, donor vector was constructed. Cas9, sgRNA 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 was 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.

- The KO region deletes most of the coding sequence, but does not result in frameshift.
- The *Hepacam2* gene is located on the Chr6. 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)

## Hepacam2 HEPACAM family member 2 [ *Mus musculus* (house mouse) ]

[Download Datasets](#)

Gene ID: 101202, updated on 17-Feb-2021

### Summary

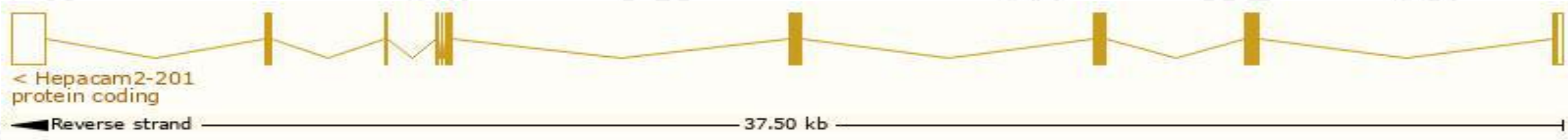
Official Symbol	Hepacam2 provided by <a href="#">MGI</a>
Official Full Name	HEPACAM family member 2 provided by <a href="#">MGI</a>
Primary source	<a href="#">MGI:MGI:2141520</a>
See related	<a href="#">Ensembl:ENSMUSG00000044156</a>
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	<a href="#">Mus musculus</a>
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	AI987662
Expression	Biased expression in colon adult (RPKM 18.5), large intestine adult (RPKM 13.0) and 5 other tissues <a href="#">See more</a>
Orthologs	<a href="#">human</a> <a href="#">all</a>
<b>NEW</b>	Try the new <a href="#">Gene table</a>
	Try the new <a href="#">Transcript table</a>

# Transcript information (Ensembl)

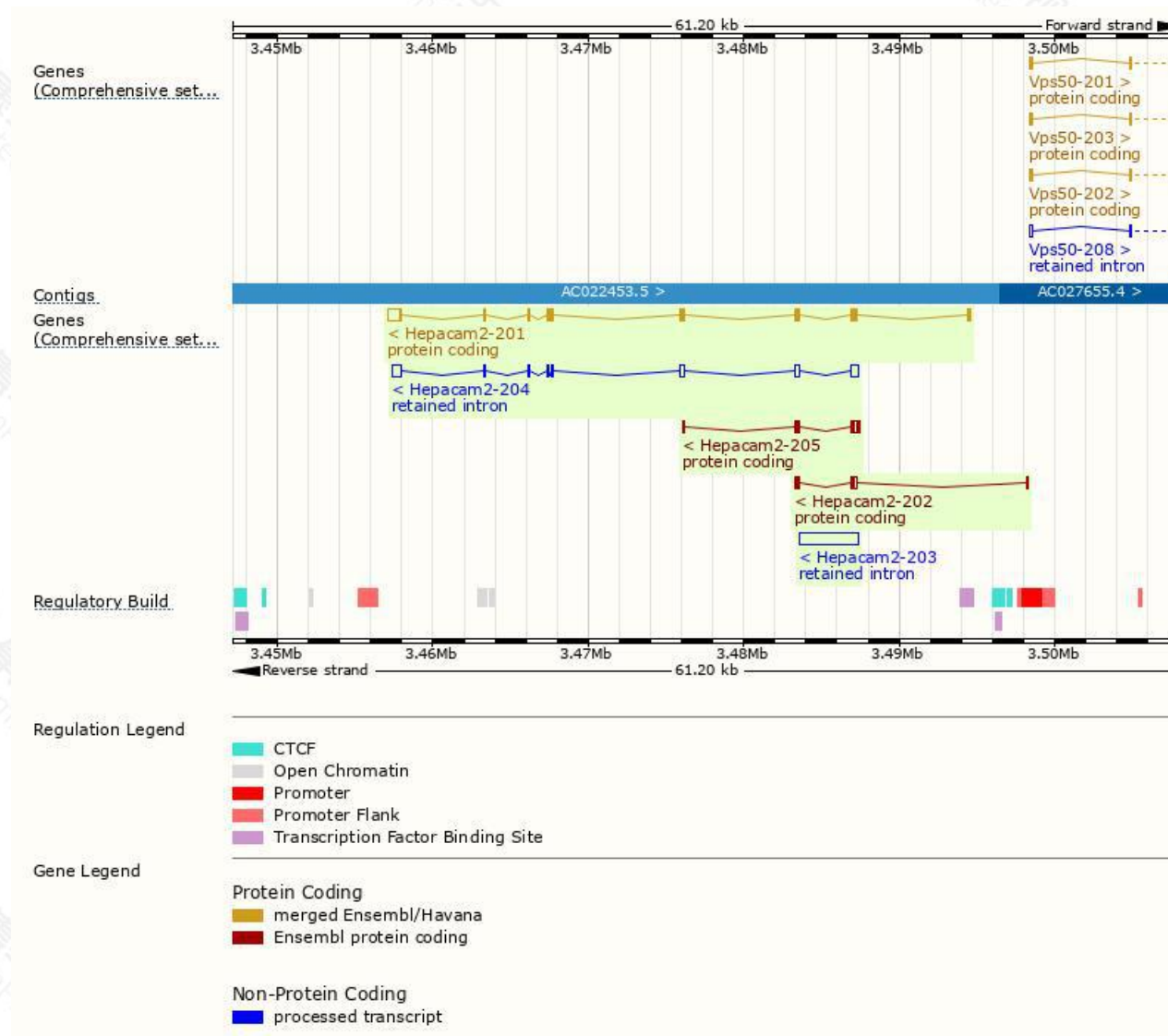
The gene has 5 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Hepacam2-201	<a href="#">ENSMUST00000049985.14</a>	2328	<a href="#">463aa</a>	Protein coding	<a href="#">CCDS39414</a>	<a href="#">Q4VAH7</a>	TSL:1 GENCODE basic APPRIS P1
Hepacam2-205	<a href="#">ENSMUST00000201607.3</a>	796	<a href="#">206aa</a>	Protein coding	-	<a href="#">A0A0J9YTS9</a>	CDS 3' incomplete TSL:5
Hepacam2-202	<a href="#">ENSMUST00000200854.1</a>	690	<a href="#">153aa</a>	Protein coding	-	<a href="#">V9GX19</a>	CDS 3' incomplete TSL:3
Hepacam2-203	<a href="#">ENSMUST00000200972.1</a>	3786	No protein	Retained intron	-	-	TSL:NA
Hepacam2-204	<a href="#">ENSMUST00000201276.1</a>	1945	No protein	Retained intron	-	-	TSL:5

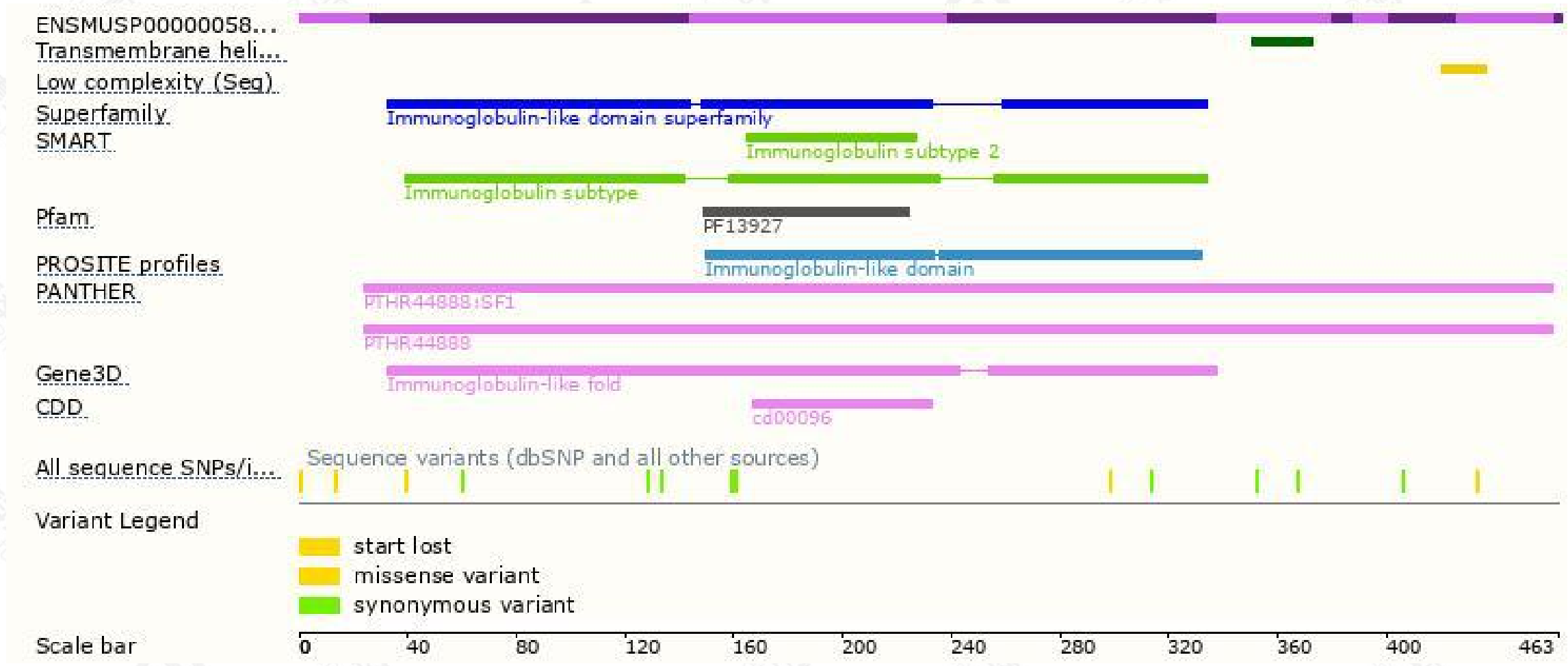
The strategy is based on the design of *Hepacam2-201* transcript,the transcription is shown below:



# Genomic location distribution



# Protein domain



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

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