Ackr3 Cas9-KO Strategy

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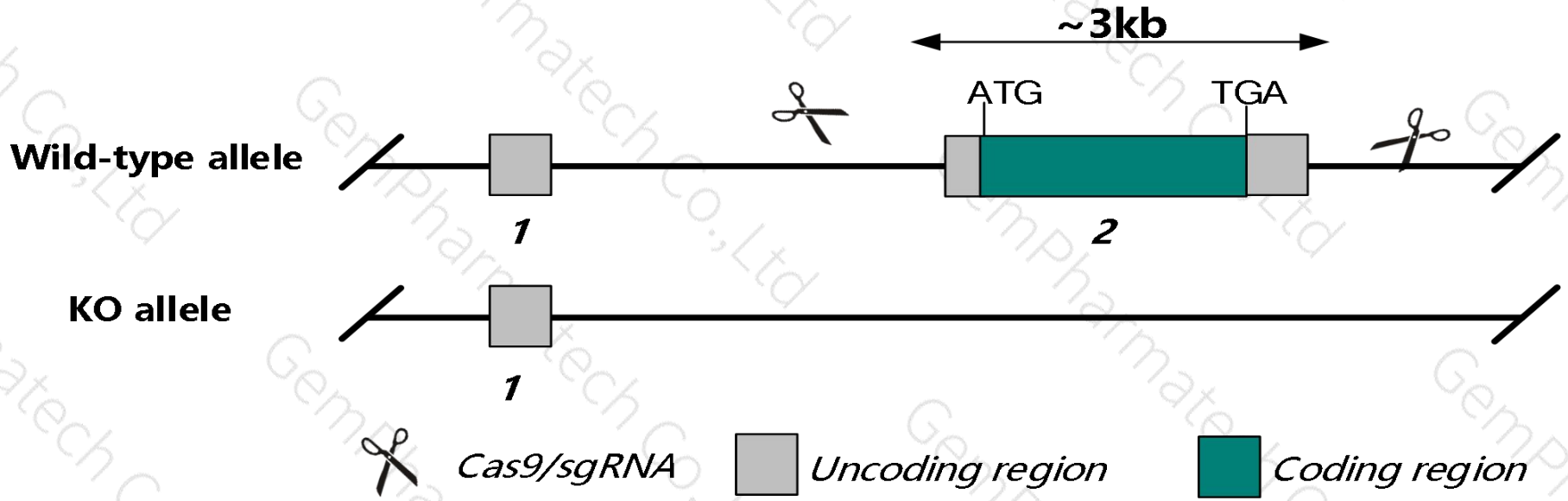
Project Overview

Project Name	***Ackr3***
Project type	**Cas9-KO**
Strain background	**C57BL/6J**

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Ackr3* gene. The schematic diagram is as follows:



Technical routes

- The *Ackr3* gene has 2 transcripts. According to the structure of *Ackr3* gene, exon2 of *Ackr3-201* (ENSMUST00000065587.4) transcript is recommended as the knockout region.The region contains start codon ATG. Knock out the region will result in disruption of protein function.

- In this project we use CRISPR/Cas9 technology to modify *Ackr3* gene. The brief process is as follows: sgRNA was transcribed in vitro.Cas9 and sgRNA were microinjected into the fertilized eggs of C57BL/6J 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/6J mice.

Notice

- According to the existing MGI data,Most homozygous null mutations result in perinatal lethality with cardiac defects including semilunar valve defects.
- This strategy knocks out the initiation codon (ATG) of the Ackr3 gene and may recognize the new initiation codon (ATG), forming an unknown protein.
- The *Ackr3* gene is located on the Chr1. 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information（NCBI）

集萃药康 GemPharmatech

Ackr3 atypical chemokine receptor 3 [*Mus musculus* (house mouse)]

Gene ID: 12778, updated on 23-Apr-2019

Summary

Official Symbol	Ackr3 provided by MGI
Official Full Name	atypical chemokine receptor 3 provided by MGI
Primary source	MGI:MGI:109562
See related	Ensembl:ENSMUSG00000044337
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	Rdc1; Cxcr7; RDC-1; CXC-R7; CXCR-7; Cmkor1; AW541270
Expression	Broad expression in adrenal adult (RPKM 169.3), heart adult (RPKM 116.2) and 20 other tissues See more
Orthologs	human all

Genomic context

Location: 1 D; 1 45.28 cM

See Ackr3 in Genome Data Viewer

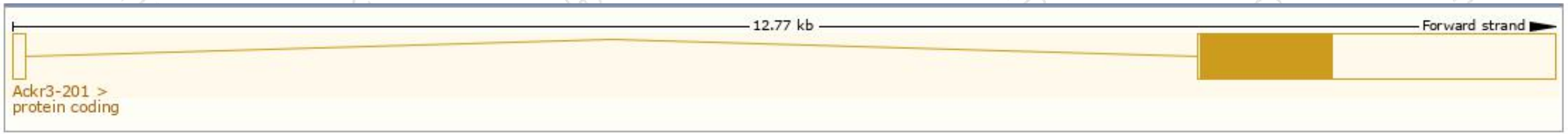
Exon count: 3

Transcript information（Ensembl）

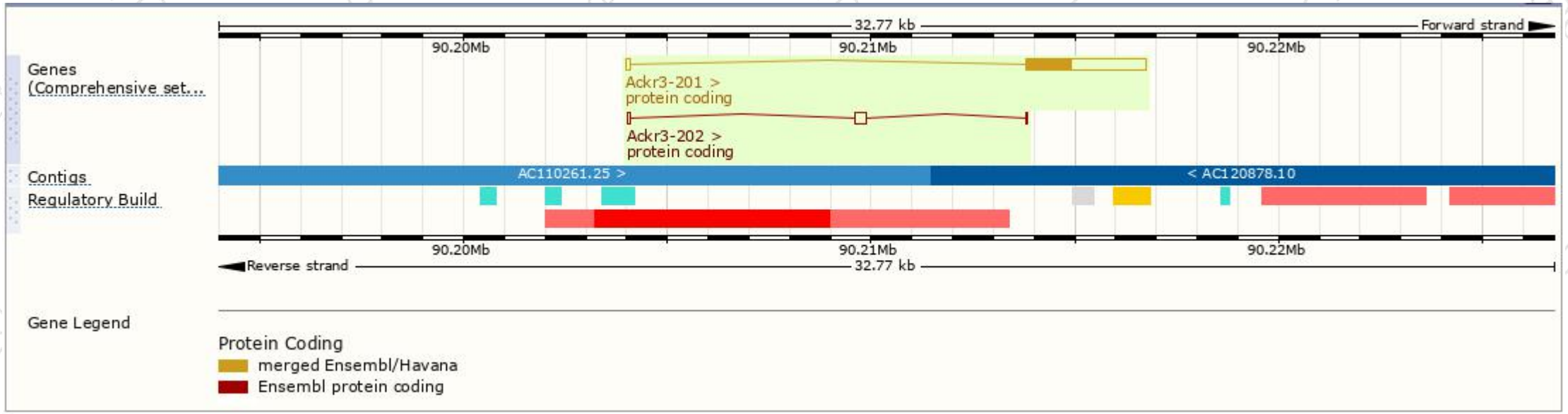
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ackr3-201	ENSMUST00000065587.4	3065	362aa	Protein coding	CCDS15152	P56485	TSL:1 GENCODE basic APPRIS P1
Ackr3-202	ENSMUST00000159654.1	404	4aa	Protein coding	-	-	CDS 3' incomplete TSL:3

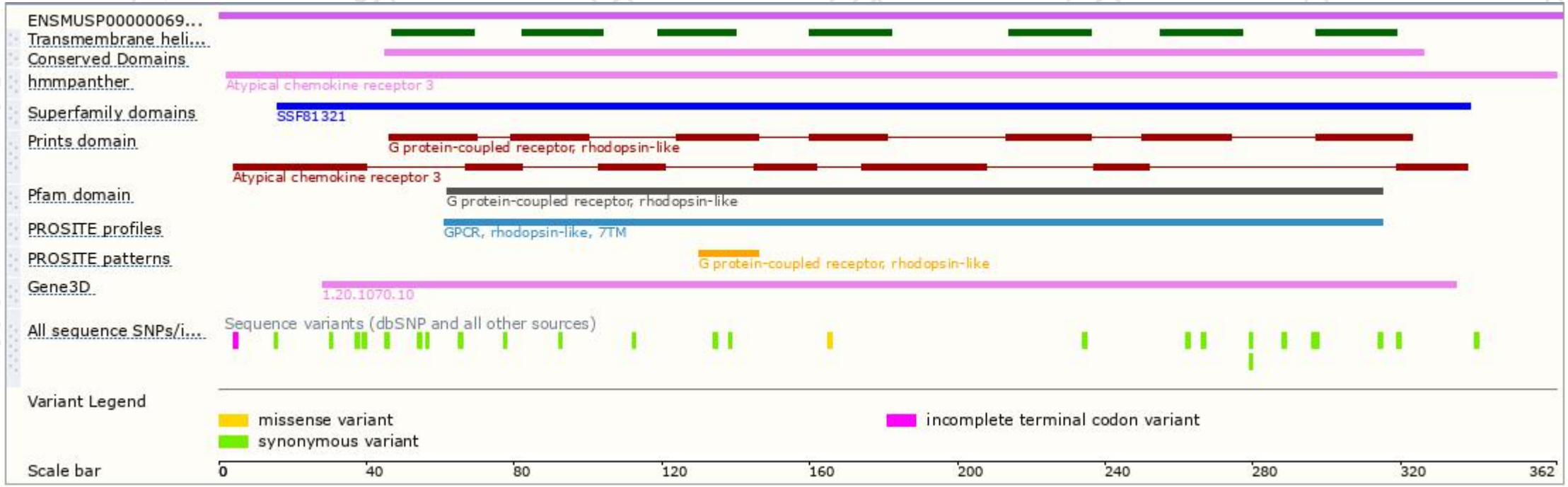
The strategy is based on the design of *Ackr3-201* transcript,The transcription is shown below



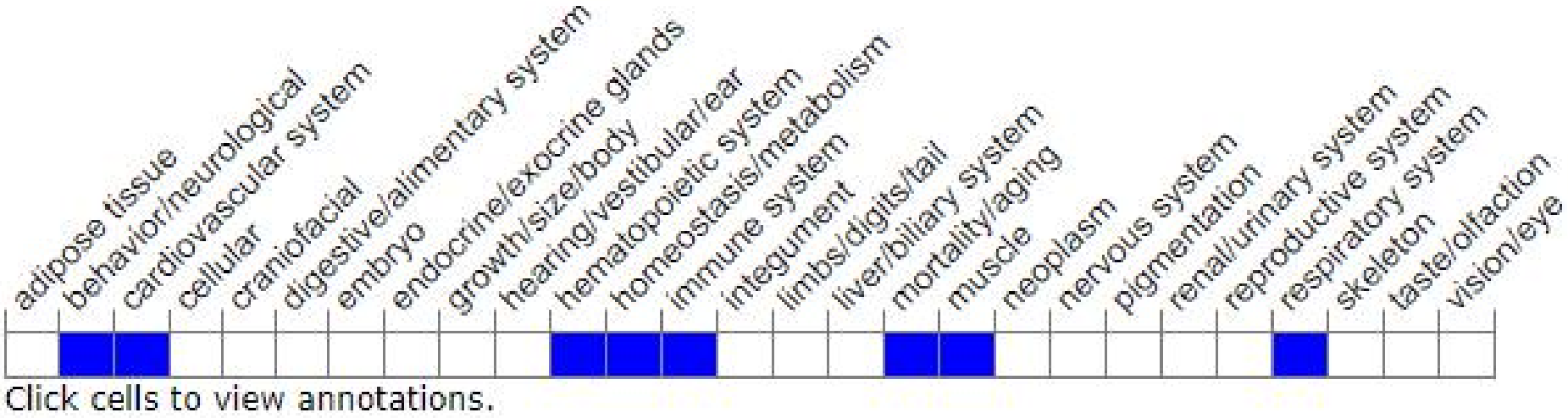
Genomic location distribution



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/marker/MGI:109562).

Most homozygous null mutations result in perinatal lethality with cardiac defects including semilunar valve defects.

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

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