Rxfp1 Cas9-CKO Strategy

Designer: Huan Wang
Reviewer: Huan Fan
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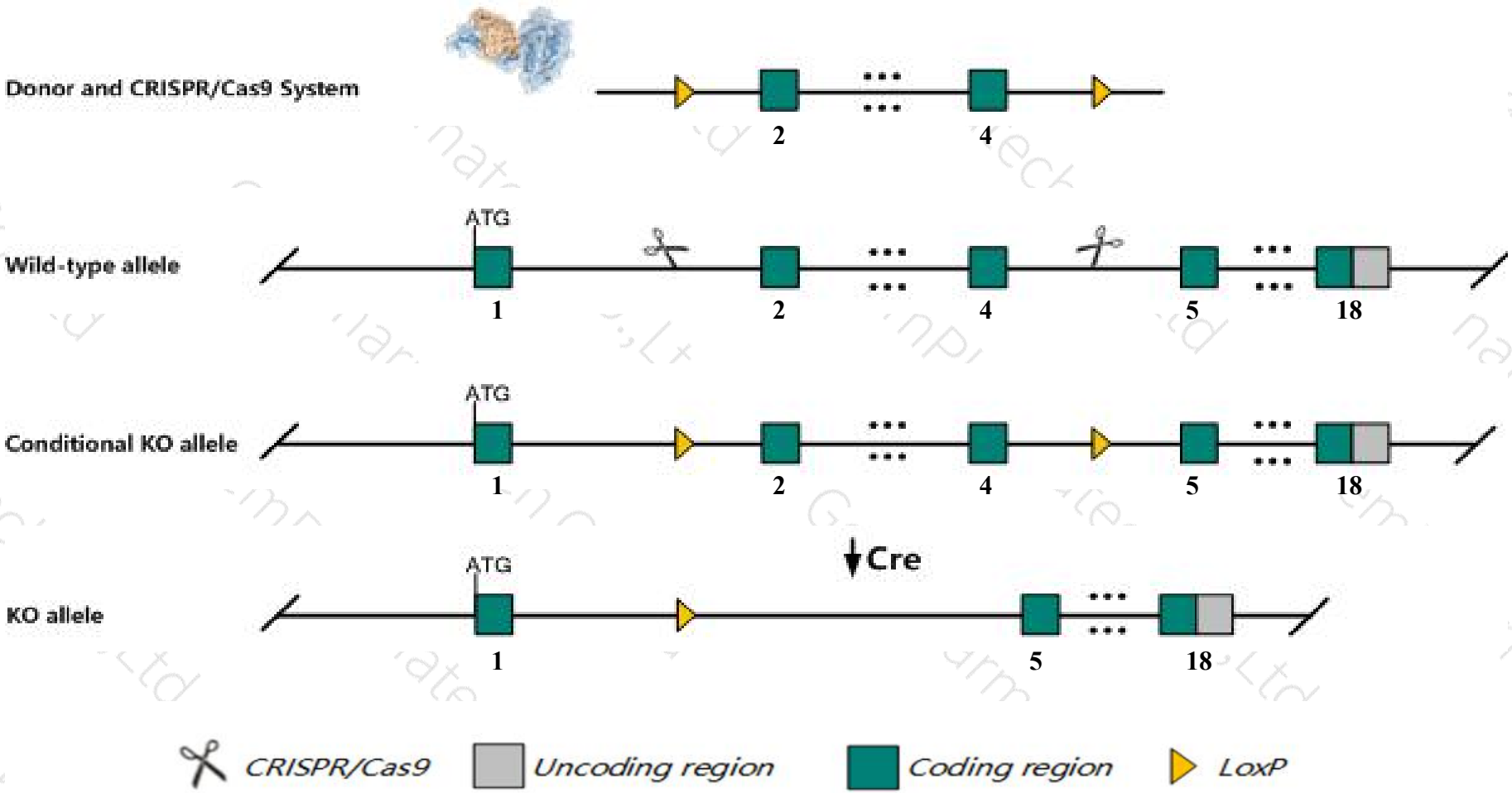
Project Overview

Project Name	***Rxfp1***
Project type	**Cas9-CKO**
Strain background	**C57BL/6JGpt**

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Rxfp1* gene. The schematic diagram is as follows:



Technical routes

- The *Rxfp1* gene has 4 transcripts. According to the structure of *Rxfp1* gene, exon2-exon4 of *Rxfp1-201* (ENSMUST00000078527.12) transcript is recommended as the knockout region.The region contains 343bp coding sequence. Knock out the region will result in disruption of protein function.

- In this project we use CRISPR/Cas9 technology to modify *Rxfp1* 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

- According to the existing MGI data,Mice homozygous for disruptions in this gene display reduced male fertility, particularly at younger ages and early generations. Impaired nipple development prevents nursing by females.

- The *Rxfp1* gene is located on the Chr3. 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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Rxfp1 relaxin/insulin-like family peptide receptor 1 [Mus musculus (house mouse)]

Gene ID: 381489, updated on 31-Jan-2019

Summary

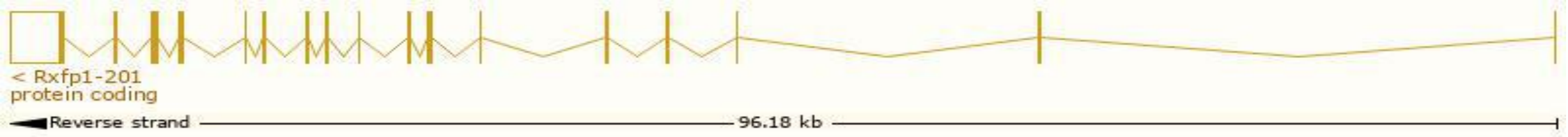
Official Symbol	Rxfp1 provided by MGI
Official Full Name	relaxin/insulin-like family peptide receptor 1 provided by MGI
Primary source	MGI:MGI:2682211
See related	Ensembl:ENSMUSG00000034009
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Gm1018, Lgr7
Expression	Biased expression in subcutaneous fat pad adult (RPKM 9.6), cortex adult (RPKM 2.8) and 4 other tissues See more
Orthologs	human all

Transcript information（Ensembl）

The gene has 4 transcripts,all transcripts are shown below:

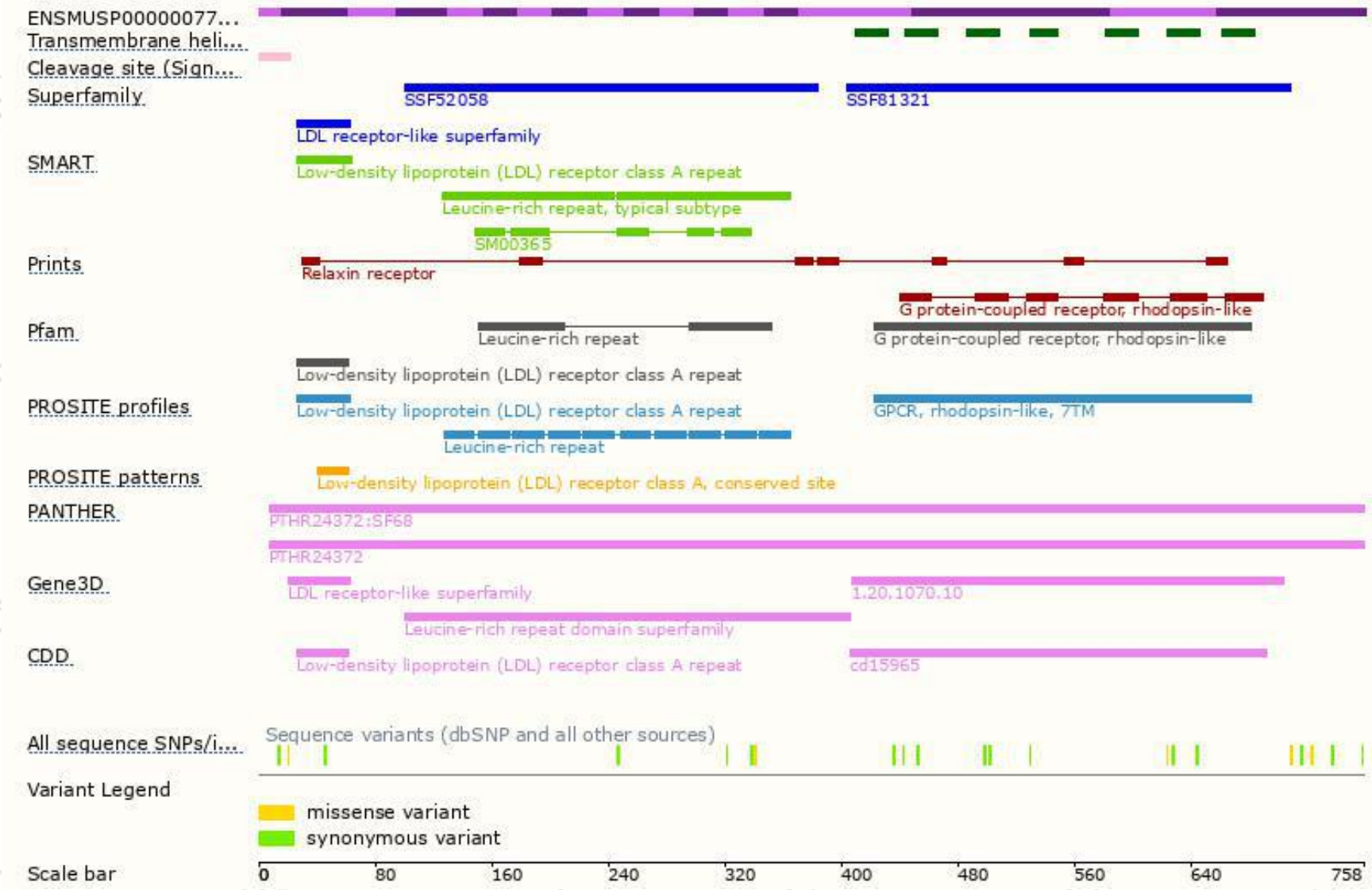
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Rxfp1-201	ENSMUST00000078527.12	5382	758aa	Protein coding	CCDS17420	Q6R6I7	TSL:1 GENCODE basic APPRIS P1
Rxfp1-202	ENSMUST00000182491.6	4983	102aa	Nonsense mediated decay	-	S4R2B5	TSL:1
Rxfp1-204	ENSMUST00000183199.1	1883	No protein	Retained intron	-	-	TSL:NA
Rxfp1-203	ENSMUST00000183040.2	1388	No protein	lncRNA	-	-	TSL:1

The strategy is based on the design of *Rxfp1-201* transcript,The transcription is shown below

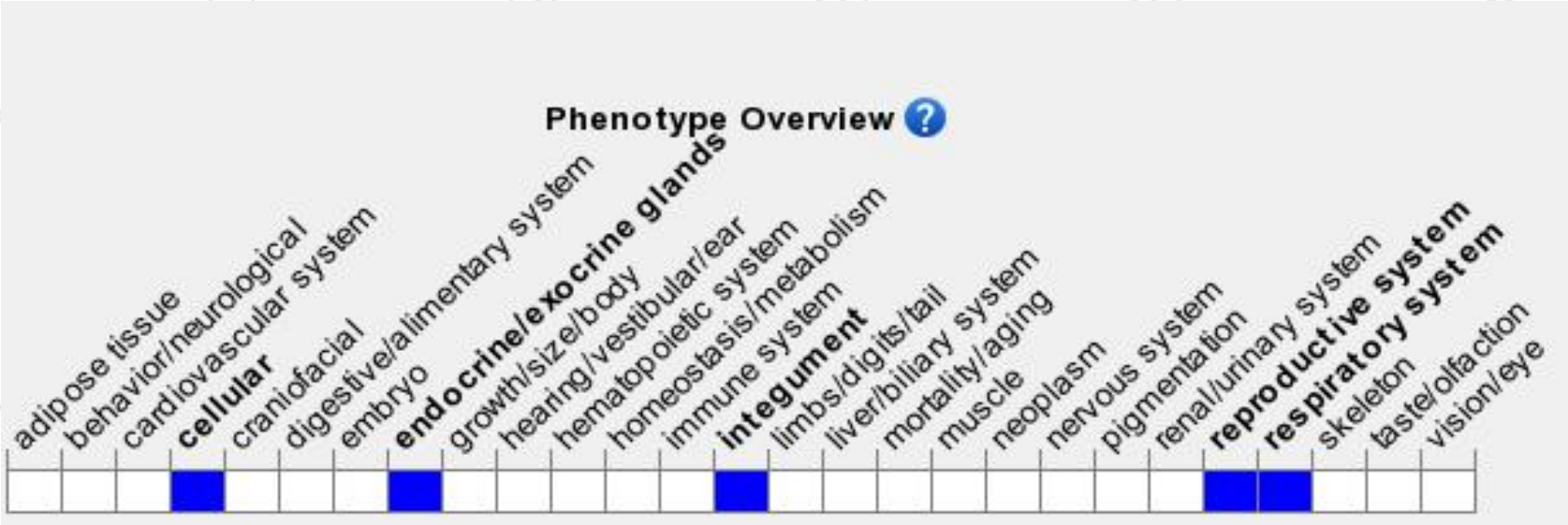


Genomic location distribution

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/).

According to the existing MGI data,Mice homozygous for disruptions in this gene display reduced male fertility, particularly at younger ages and early generations. Impaired nipple development prevents nursing by females.

集萃药康
GemPharmatech

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

Tel: 400-9660890

集萃药康生物科技
GemPharmatech Co.,Ltd