

Fpr2 Cas9-KO Strategy

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Design Date: 2019-7-30

Project Overview



Project Name

Fpr2

Project type

Cas9-KO

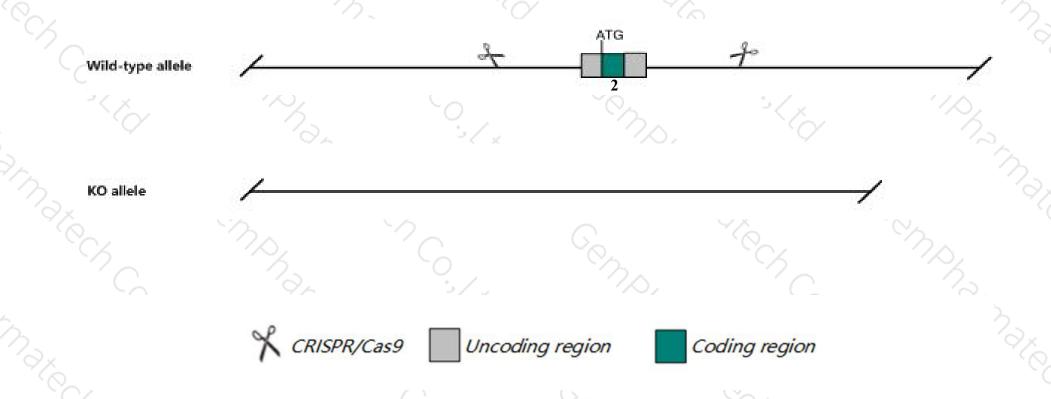
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Fpr2 gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Fpr2* gene has 2 transcripts. According to the structure of *Fpr2* gene, exon2 of *Fpr2-201* (ENSMUST00000064068.4) 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 *Fpr2* gene. The brief process is as follows: CRISPR/Cas9 system v

Notice



- > According to the existing MGI data, Mice homozygous for a targeted reporter allele exhibit altered leukocyte responses and experimentally induced inflammation.
- \succ The KO region contains functional region of the *Fpr3* gene. Knockout the region may affect the function of *Fpr3* gene.
- The *Fpr2* 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)



Fpr2 formyl peptide receptor 2 [Mus musculus (house mouse)]

Gene ID: 14289, updated on 12-Mar-2019

Summary

☆ ?

Official Symbol Fpr2 provided by MGI

Official Full Name formyl peptide receptor 2 provided by MGI

Primary source MGI:MGI:1278319

See related Ensembl: ENSMUSG00000052270

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 E330010107Rik, Fpr-rs2

Expression Biased expression in liver E18 (RPKM 10.5), lung adult (RPKM 3.4) and 3 other tissuesSee more

Orthologs human all

Transcript information (Ensembl)



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

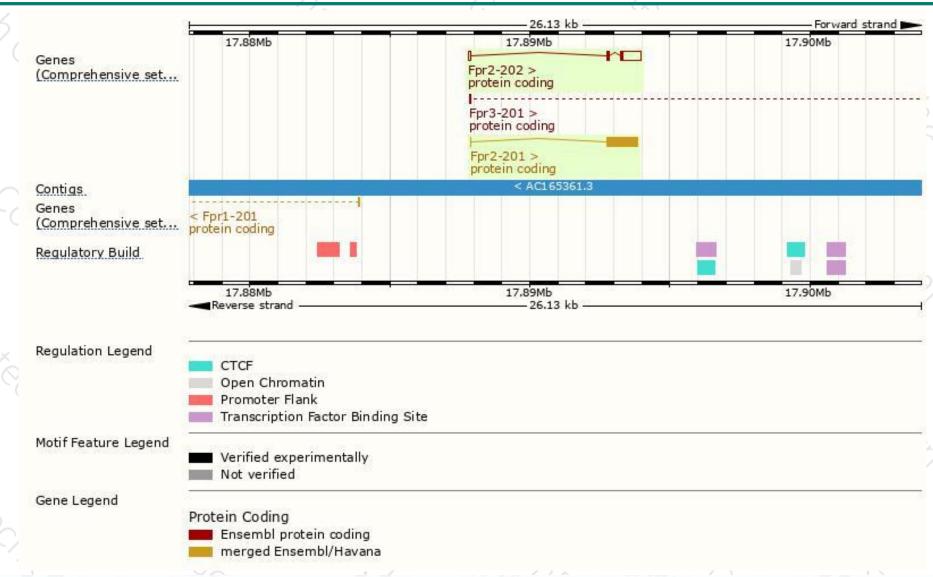
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Fpr2-201	ENSMUST00000064068.4	1133	351aa	Protein coding	CCDS28419	<u>088536</u>	TSL:1 GENCODE basic APPRIS P1
Fpr2-202	ENSMUST00000149944.1	883	<u>48aa</u>	Protein coding	-	A0A2I3BPU6	TSL:1 GENCODE basic

The strategy is based on the design of *Fpr2-201* transcript, The transcription is shown below

Fpr2-201 > protein coding

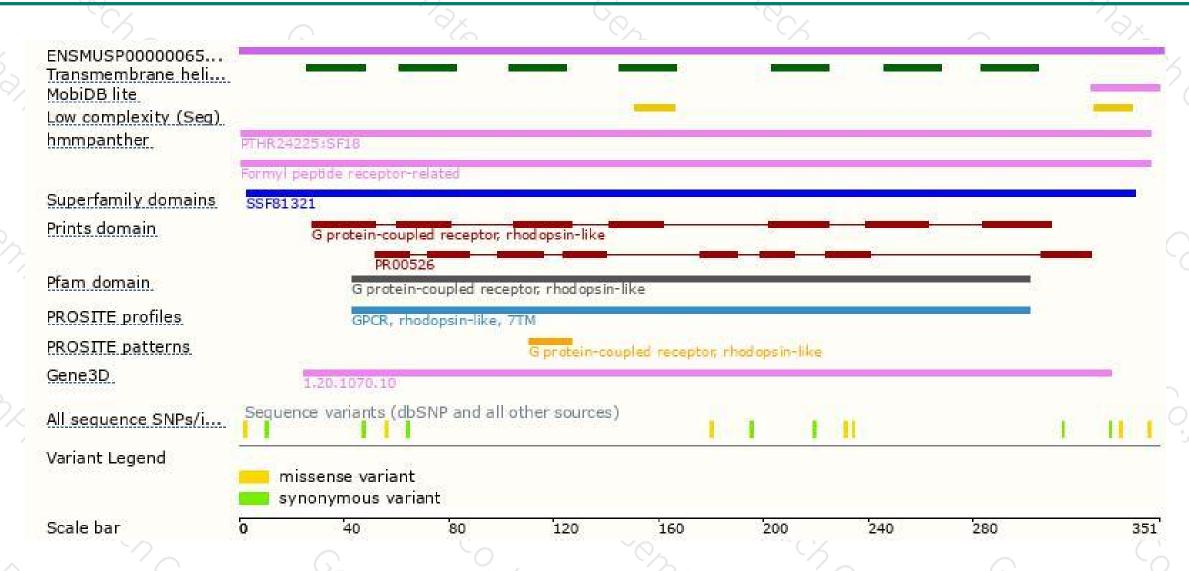
Genomic location distribution





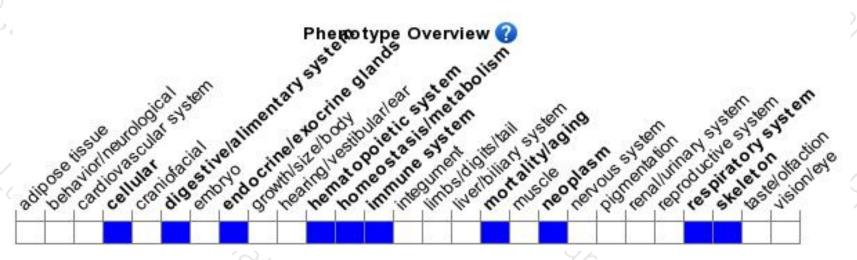
Protein domain





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 a targeted reporter allele exhibit altered leukocyte responses and experimentally induced inflammation.



If you have any questions, you are welcome to inquire. Tel: 400-9660890





