

Yipf6 Cas9-KO Strategy

Designer:Qiong Zhou

Project Overview



Project Name

Yipf6

Project type

Cas9-KO

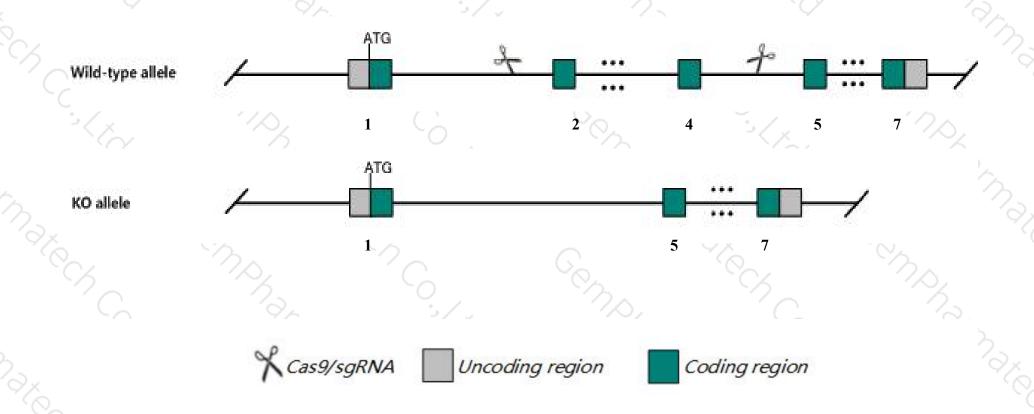
Strain background

C57BL/6J

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Yipf6 gene. The schematic diagram is as follows:



Technical routes



- ➤ The Yipf6 gene has 6 transcripts. According to the structure of Yipf6 gene, exon2-exon4 of Yipf6-201 (ENSMUST0000054697.6) transcript is recommended as the knockout region. The region contains 251bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Yipf6* 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, Mice homozygous or hemizygous for an ENU-induced allele exhibit colitis and increased susceptibility to induced colitis with decreased Paneth and goblet cells.
- ➤ The *Yipf6* gene is located on the ChrX. 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)



Yipf6 Yip1 domain family, member 6 [Mus musculus (house mouse)]

Gene ID: 77929, updated on 19-Feb-2019

Summary

☆ ?

Official Symbol Yipf6 provided by MGI

Official Full Name Yip1 domain family, member 6 provided by MGI

Primary source MGI:MGI:1925179

See related Ensembl:ENSMUSG00000047694

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 A430107J06Rik

Expression Ubiquitous expression in bladder adult (RPKM 7.2), placenta adult (RPKM 5.4) and 28 other tissuesSee more

Orthologs <u>human</u> all

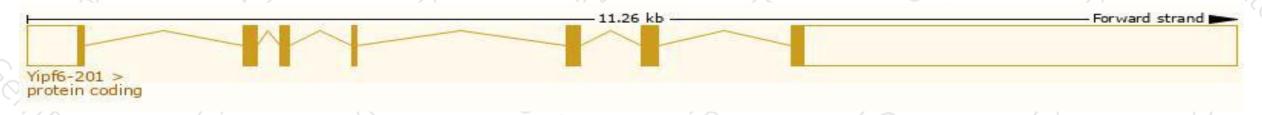
Transcript information (Ensembl)



The gene has 6 transcripts, all transcripts are shown below:

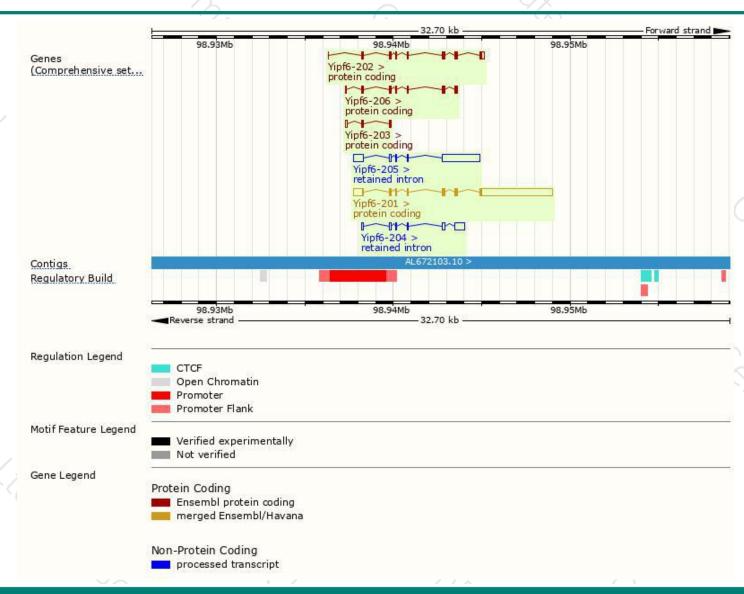
1 1/2	William II		f agent				
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Yipf6-201	ENSMUST00000054697.6	5211	236aa	Protein coding	CCDS30296	Q8BR70	TSL:1 GENCODE basic APPRIS P1
Yipf6-202	ENSMUST00000113811.7	930	<u>236aa</u>	Protein coding	CCDS30296	Q8BR70	TSL:1 GENCODE basic APPRIS P1
Yipf6-206	ENSMUST00000151353.7	690	<u>198aa</u>	Protein coding	828	B1AV66	CDS 3' incomplete TSL:5
Yipf6-203	ENSMUST00000124010.1	284	<u>55aa</u>	Protein coding	3525	B1AV65	CDS 3' incomplete TSL:3
Yipf6-205	ENSMUST00000149082.7	2908	No protein	Retained intron	187	1.00	TSL:1
Yipf6-204	ENSMUST00000140479.1	1020	No protein	Retained intron	14.		TSL:1

The strategy is based on the design of Yipf6-201 transcript, The transcription is shown below



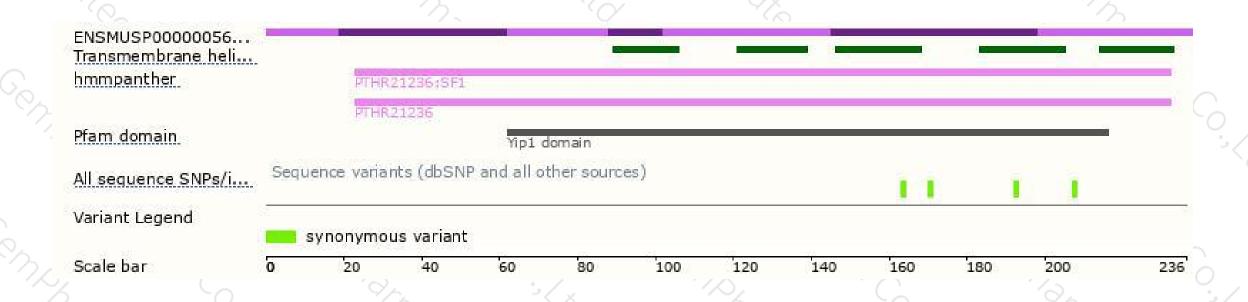
Genomic location distribution





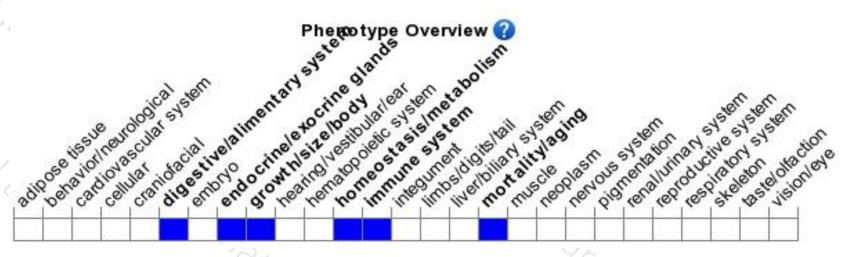
Protein domain





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 or hemizygous for an ENU-induced allele exhibit colitis and increased susceptibility to induced colitis with decreased Paneth and goblet cells.



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

Tel: 025-5864 1534





