

John Skech Co. 1/4 Pth1r Cas9-CKO Strategy Ronnohamakech Co. / Ky

Designer: JiaYu

Project Overview



Project Name Pth1r

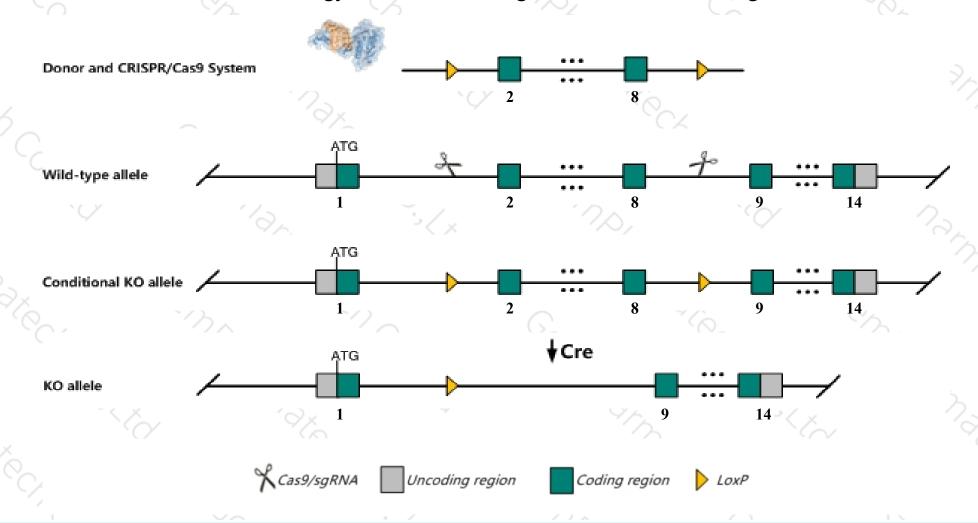
Project type Cas9-CKO

Strain background C57BL/6J

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Pth1r* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Pth1r* gene has 6 transcripts. According to the structure of *Pth1r* gene, exon2-exon8 of *Pth1r-201*(ENSMUST0000006005.11) transcript is recommended as the knockout region. The region contains 913bp coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Pth1r* 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/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.
- > 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.

Notice



- > According to the existing MGI data, Homozygous mutant mice die in mid-gestation or shortly after birth depending on genetic background, are small in size, have short limbs, and accelerated differentiation of chondrocytes resulting in accelerated bone mineralization.
- > The *Pth1r* gene is located on the Chr9. 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)



Pth1r parathyroid hormone 1 receptor [Mus musculus (house mouse)]

Gene ID: 19228, updated on 19-Mar-2019

Summary

☆ ?

Official Symbol Pth1r provided by MGI

Official Full Name parathyroid hormone 1 receptor provided by MGI

Primary source MGI:MGI:97801

See related Ensembl:ENSMUSG00000032492

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 PPR, Pthr, Pthr1

Expression Biased expression in kidney adult (RPKM 207.3), limb E14.5 (RPKM 43.7) and 4 other tissuesSee more

Orthologs <u>human</u> <u>all</u>

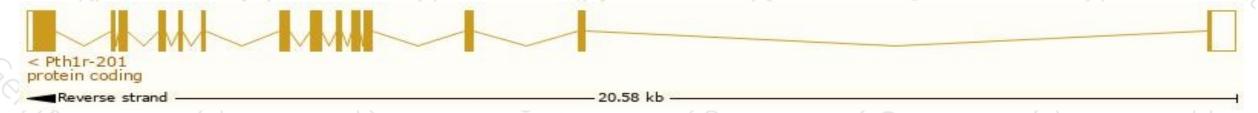
Transcript information (Ensembl)



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

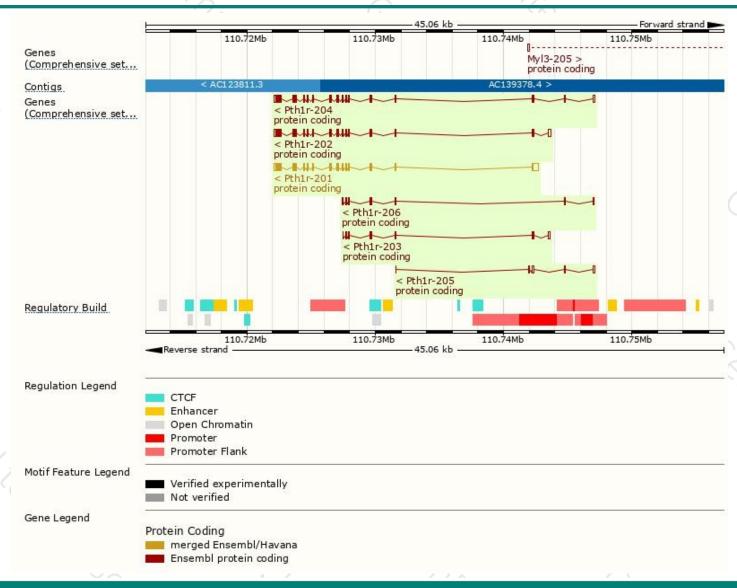
Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
ENSMUST00000006005.11	2307	<u>591aa</u>	Protein coding	CCDS52940	P41593	TSL:1 GENCODE basic APPRIS P1
ENSMUST00000166716.7	2219	<u>591aa</u>	Protein coding	CCDS52940	P41593	TSL:1 GENCODE basic APPRIS P1
ENSMUST00000198865.4	2201	<u>591aa</u>	Protein coding	CCDS52940	<u>P41593</u>	TSL:1 GENCODE basic APPRIS P1
ENSMUST00000196057.1	830	<u>197aa</u>	Protein coding	-	A0A0G2JG89	CDS 3' incomplete TSL:5
ENSMUST00000199862.4	682	<u>198aa</u>	Protein coding	-	A0A0G2JE85	CDS 3' incomplete TSL:5
ENSMUST00000199791.1	508	<u>54aa</u>	Protein coding	-	A0A0G2JEZ1	CDS 3' incomplete TSL:5
	ENSMUST000000166716.7 ENSMUST00000198865.4 ENSMUST00000196057.1 ENSMUST00000199862.4	ENSMUST00000166716.7 2219 ENSMUST00000198865.4 2201 ENSMUST00000196057.1 830 ENSMUST00000199862.4 682	ENSMUST00000166716.7 2219 591aa ENSMUST00000198865.4 2201 591aa ENSMUST00000196057.1 830 197aa ENSMUST00000199862.4 682 198aa	ENSMUST00000006005.11 2307 591aa Protein coding ENSMUST00000166716.7 2219 591aa Protein coding ENSMUST00000198865.4 2201 591aa Protein coding ENSMUST00000196057.1 830 197aa Protein coding ENSMUST00000199862.4 682 198aa Protein coding	ENSMUST00000006005.11 2307 591aa Protein coding CCDS52940 ENSMUST00000166716.7 2219 591aa Protein coding CCDS52940 ENSMUST00000198865.4 2201 591aa Protein coding CCDS52940 ENSMUST00000196057.1 830 197aa Protein coding - ENSMUST00000199862.4 682 198aa Protein coding -	ENSMUST00000006005.11 2307 591aa Protein coding CCDS52940 P41593 ENSMUST00000166716.7 2219 591aa Protein coding CCDS52940 P41593 ENSMUST00000198865.4 2201 591aa Protein coding CCDS52940 P41593 ENSMUST00000196057.1 830 197aa Protein coding - A0A0G2JG89 ENSMUST00000199862.4 682 198aa Protein coding - A0A0G2JE85

The strategy is based on the design of *Pth1r-201* transcript, The transcription is shown below



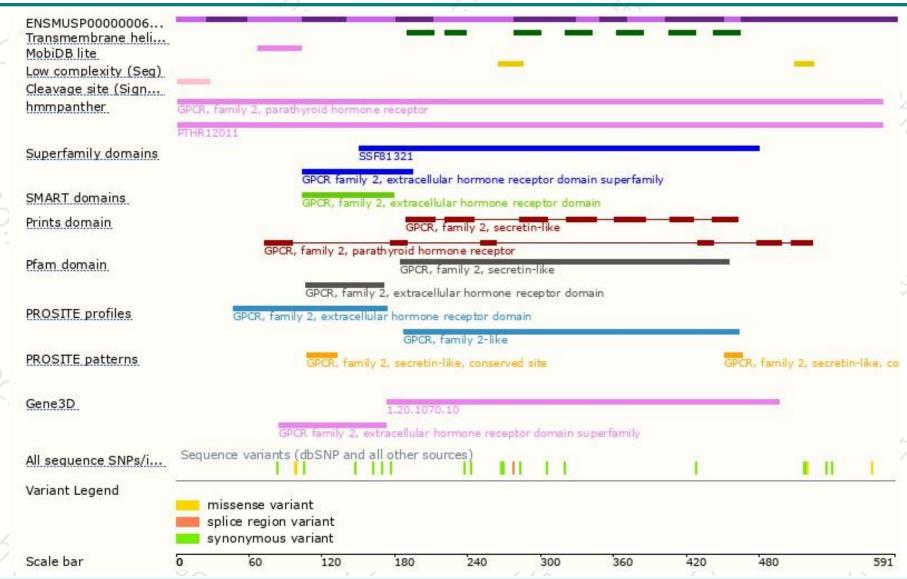
Genomic location distribution





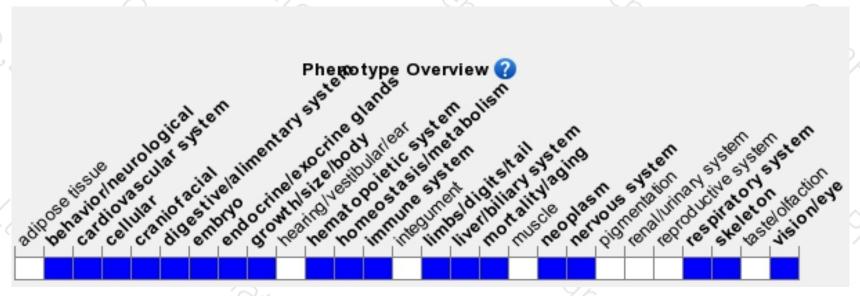
Protein domain





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, Homozygous mutant mice die in mid-gestation or shortly after birth depending on genetic background, are small in size, have short limbs, and accelerated differentiation of chondrocytes resulting in accelerated bone mineralization.



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

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