

Phospho2 Cas9-CKO Strategy

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Project Overview



Project Name

Phospho2

Project type

Cas9-CKO

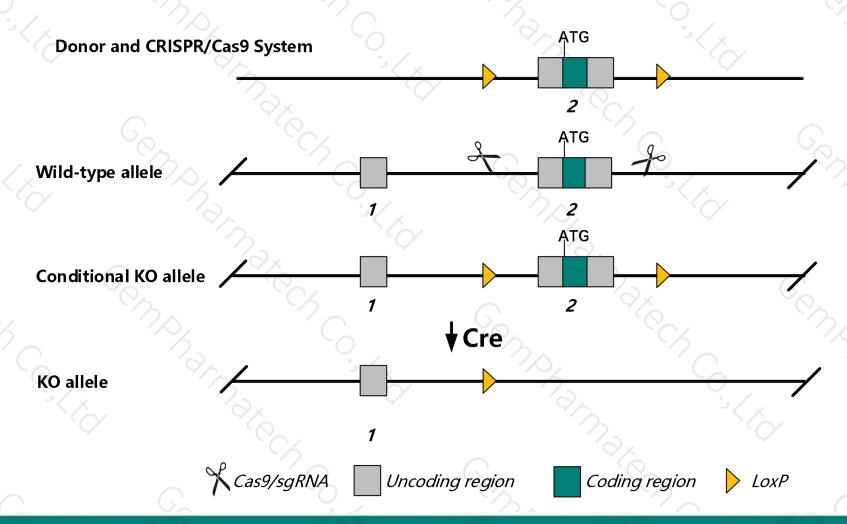
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Phospho2* gene. The schematic diagram is as follows:



Technical routes



- The *Phospho2* gene has 6 transcripts. According to the structure of *Phospho2* gene, exon2 of *Phospho2-206* (ENSMUST00000180290.1) 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 *Phospho2* 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



- Transcript 204 may not be affected. The effect of transcripts 203,205 is unknown.
- > The *Phospho2* gene is located on the Chr2. 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)



Phospho2 phosphatase, orphan 2 [Mus musculus (house mouse)]

Gene ID: 73373, updated on 13-Mar-2020

Summary

☆ ?

Official Symbol Phospho2 provided by MGI

Official Full Name phosphatase, orphan 2 provided by MGI

Primary source MGI:MGI:1920623

See related Ensembl: ENSMUSG00000027088

Gene type protein coding
RefSeq status PROVISIONAL
Organism Mus musculus

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;

Muroidea; Muridae; Murinae; Mus; Mus

Also known as 1700048E23Rik, Al661517, AU021728, AV006103, Phos2

Expression Broad expression in testis adult (RPKM 58.1), adrenal adult (RPKM 8.9) and 25 other tissuesSee more

Orthologs <u>human all</u>

Transcript information (Ensembl)



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

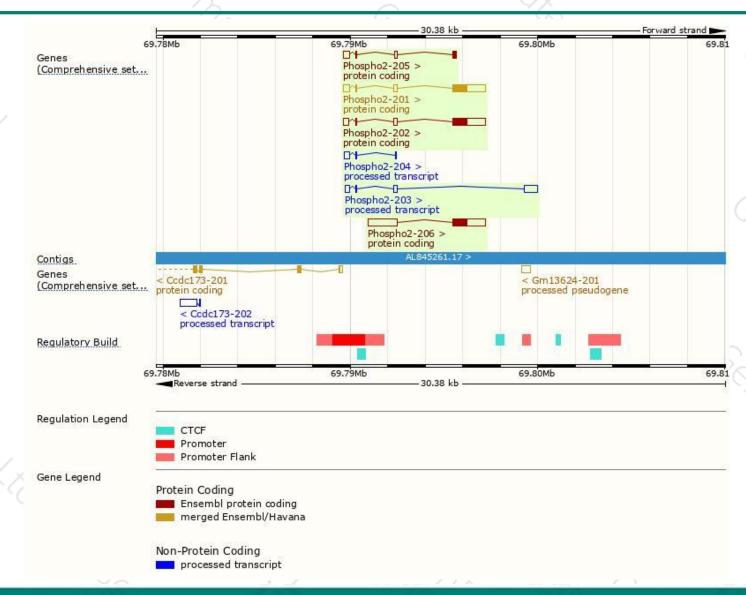
and the same	f in						
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Phospho2-206	ENSMUST00000180290.1	3267	241aa	Protein coding	CCDS16099	Q9D9M5	TSL:1 GENCODE basic APPRIS is a system to annotate alternatively spliced transcripts based on a range of computational methods to identify the most functionally important transcript(s) of a gene. APPRIS P1
Phospho2-201	ENSMUST00000028494.8	2260	241aa	Protein coding	CCDS16099	Q9D9M5	TSL:1 GENCODE basic APPRIS is a system to annotate alternatively spliced transcripts based on a range of computational methods to identify the most functionally important transcript(s) of a gene. APPRIS P1
Phospho2-202	ENSMUST00000112266.7	2194	241aa	Protein coding	CCDS16099	Q9D9M5	TSL:3 GENCODE basic APPRIS is a system to annotate alternatively spliced transcripts based on a range of computational methods to identify the most functionally important transcript(s) of a gene. APPRIS P1
Phospho2-205	ENSMUST00000151298.7	667	<u>50aa</u>	Protein coding	- 64	B0R0D2	CDS 3' incomplete TSL:2
Phospho2-203	ENSMUST00000128765.1	1133	No protein	Processed transcript	1.5	-	TSL:1
Phospho2-204	ENSMUST00000131100.7	415	No protein	Processed transcript	1-	-	TSL:2

The strategy is based on the design of *Phospho2-206* transcript, The transcription is shown below

Phospho2-206 > protein coding

Genomic location distribution





Protein domain







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





