

Spink2 Cas9-CKO Strategy

Designer: Xiaojing Li

Reviewer: JiaYu

Design Date: 2020-8-20

Project Overview



Project Name

Spink2

Project type

Cas9-CKO

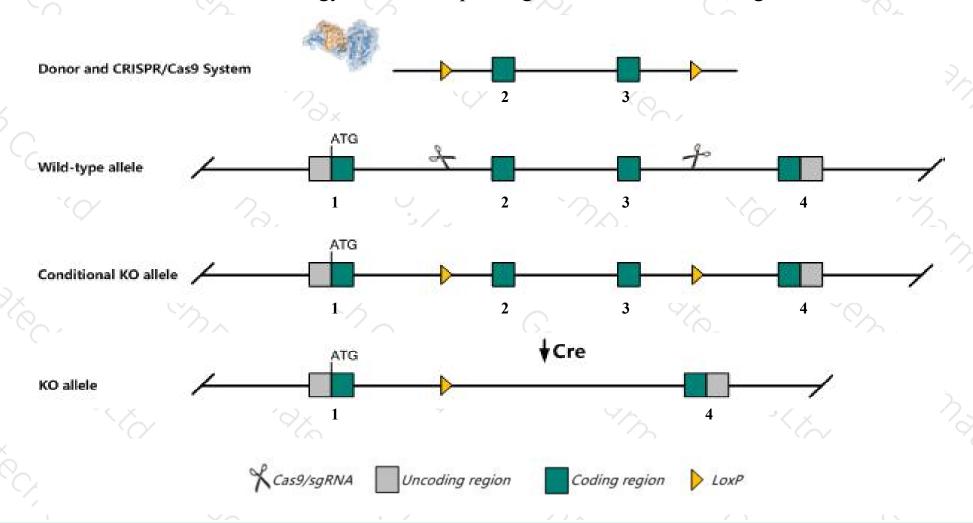
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Spink2 gene. The schematic diagram is as follows:



Technical routes



- > The *Spink2* gene has 3 transcripts. According to the structure of *Spink2* gene, exon2-exon3 of *Spink2-201*(ENSMUST00000065216.10) transcript is recommended as the knockout region. The region contains 172bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Spink2* 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/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 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, mice homozygous for a gene trap allele exhibit reduced male fertility associated with oligozoospermia, teratozoospermia, male germ cell apoptosis, and small testis.
- > The Spink2 gene is located on the Chr5. 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)



Spink2 serine peptidase inhibitor, Kazal type 2 [Mus musculus (house mouse)]

Gene ID: 69982, updated on 13-Mar-2020

Summary

☆ ?

Official Symbol Spink2 provided by MGI

Official Full Name serine peptidase inhibitor, Kazal type 2 provided byMGI

Primary source MGI:MGI:1917232

See related Ensembl:ENSMUSG00000053030

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 1700007F22Rik, AV038945, HUSI-II

Expression Restricted expression toward testis adult (RPKM 405.6)See more

Orthologs <u>human all</u>

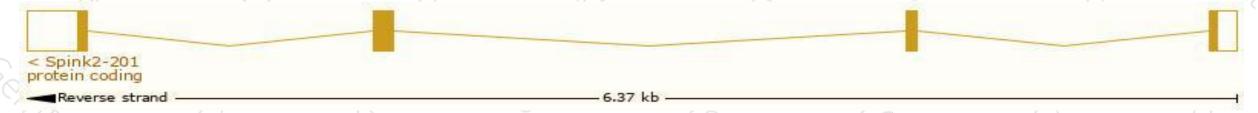
Transcript information (Ensembl)



The gene has 3 transcripts, all transcripts are shown below:

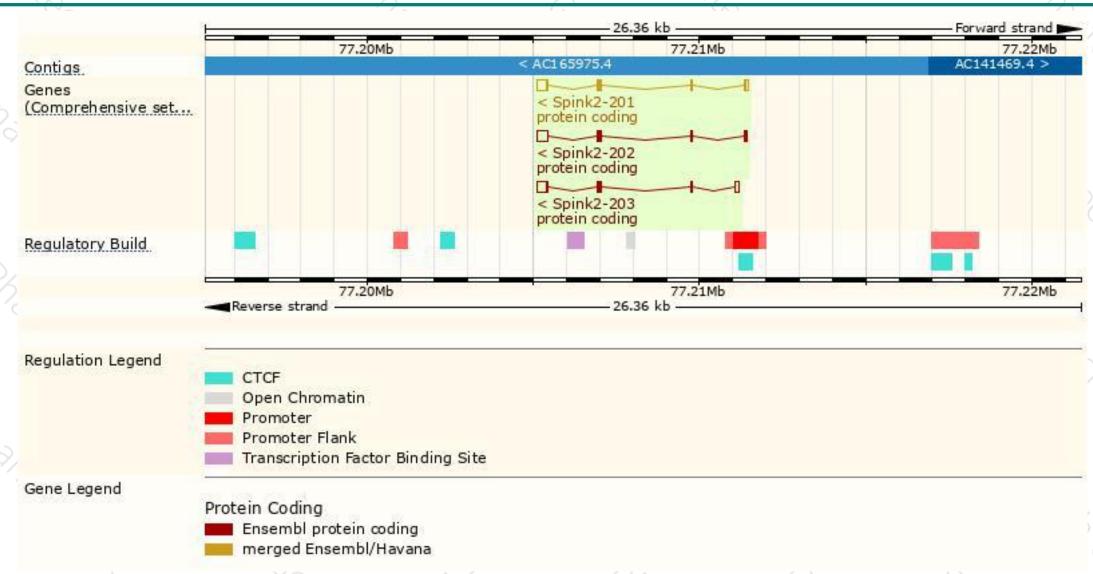
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Spink2-203	ENSMUST00000121825.1	643	<u>59aa</u>	Protein coding	CCDS80306	<u>D3Z3X9</u>	TSL:3 GENCODE basic APPRIS ALT2
Spink2-201	ENSMUST00000065216.10	631	86aa	Protein coding	CCDS39121	Q8BMY7	TSL:1 GENCODE basic APPRIS P3
Spink2-202	ENSMUST00000120429.7	594	<u>56aa</u>	Protein coding	CCDS80307	D3YTY0	TSL:1 GENCODE basic

The strategy is based on the design of *Spink2-201* transcript, the transcription is shown below:



Genomic location distribution





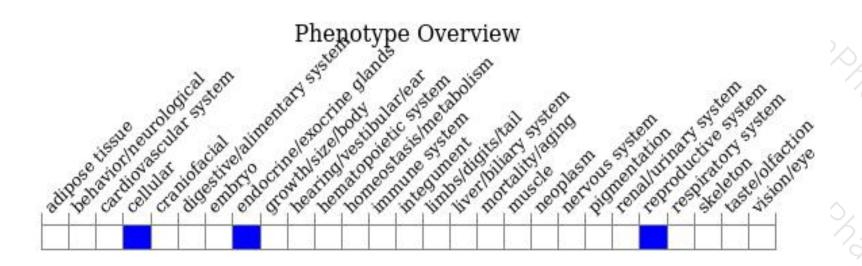
Protein domain





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 a gene trap allele exhibit reduced male fertility associated with oligozoospermia, teratozoospermia, male germ cell apoptosis, and small testis.



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

Tel: 025-5864 1534





