

Jak1 Cas9-CKO Strategy

Designer:

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Reviewer:

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Design Date:

Project Overview



Project Name

Project type

Strain background

C57BL/6JGpt

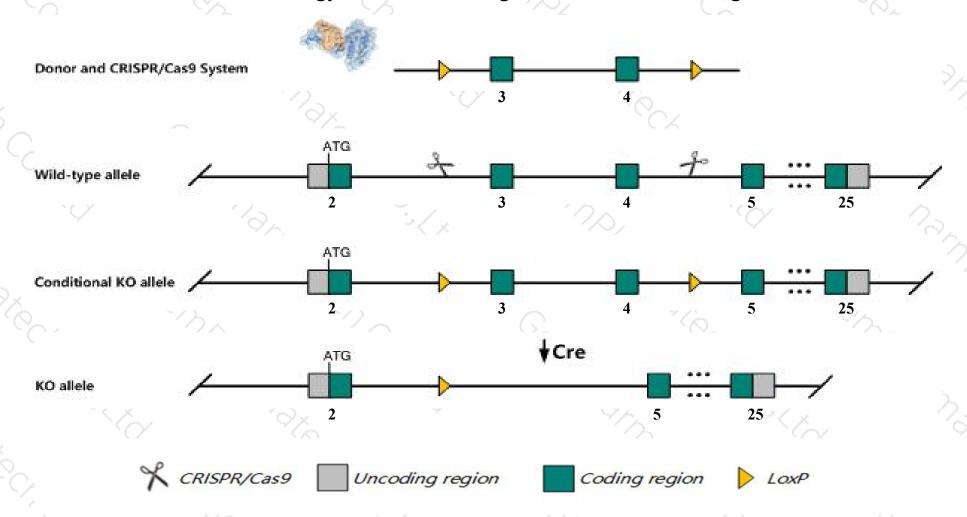
Cas9-CKO

Jak1

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Jak1 gene. The schematic diagram is as follows:



Technical routes



- The *Jak1* gene has 8 transcripts. According to the structure of *Jak1* gene, exon3-exon4 of *Jak1-201*(ENSMUST00000102781.9) transcript is recommended as the knockout region. The region contains 323bp coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Jak1* 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



- ➤ According to the existing MGI data, mice homozygous for disruption of this gene die within the first 24 hours after birth.
- The distance between gene *Gm24468* and exon4 of *Jak1-201* is about 2kb, this strategy may affect the regulatory function of *Gm24468* gene.
- The *Jak1* gene is located on the Chr4. 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)



Jak1 Janus kinase 1 [Mus musculus (house mouse)]

Gene ID: 16451, updated on 14-Dec-2019



☆ ?

Official Symbol Jak1 provided by MGI

Official Full Name Janus kinase 1 provided by MGI

Primary source MGI:MGI:96628

See related Ensembl:ENSMUSG00000028530

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 BAP004; AA960307; C130039L05Rik

Expression Ubiquitous expression in thymus adult (RPKM 28.5), spleen adult (RPKM 26.3) and 28 other tissues See more

Orthologs human all

Transcript information (Ensembl)



The gene has 8 transcripts, all transcripts are shown below:

3	f 1		f 2	"//)		1	
Name A	Transcript ID	bp 🌲	Protein	Biotype 🍦	CCDS 🍦	UniProt 🌲	Flags
Jak1-201	ENSMUST00000102781.9	4906	<u>1153aa</u>	Protein coding	CCDS18393 ₽	B1ASP2 ₺	TSL:1 GENCODE basic APPRIS P1
Jak1-202	ENSMUST00000125968.1	374	No protein	IncRNA	(14)	(H)	TSL:3
Jak1-203	ENSMUST00000132292.1	778	No protein	IncRNA	8578	1. Table 1.	TSL:5
Jak1-204	ENSMUST00000136167.7	589	No protein	IncRNA	(S 4)	(H)	TSL:3
Jak1-205	ENSMUST00000147211.1	2039	No protein	IncRNA	8578	12 5 3	TSL:1
Jak1-206	ENSMUST00000149297.1	421	<u>18aa</u>	Protein coding	641	B1ASP3 ₺	CDS 3' incomplete TSL:3
Jak1-207	ENSMUST00000151235.1	504	No protein	IncRNA	8578	\ -	TSL:1
Jak1-208	ENSMUST00000155328.7	671	No protein	IncRNA	(S#)	(H)	TSL:5

The strategy is based on the design of Jak1-201 transcript, the transcription is shown below:

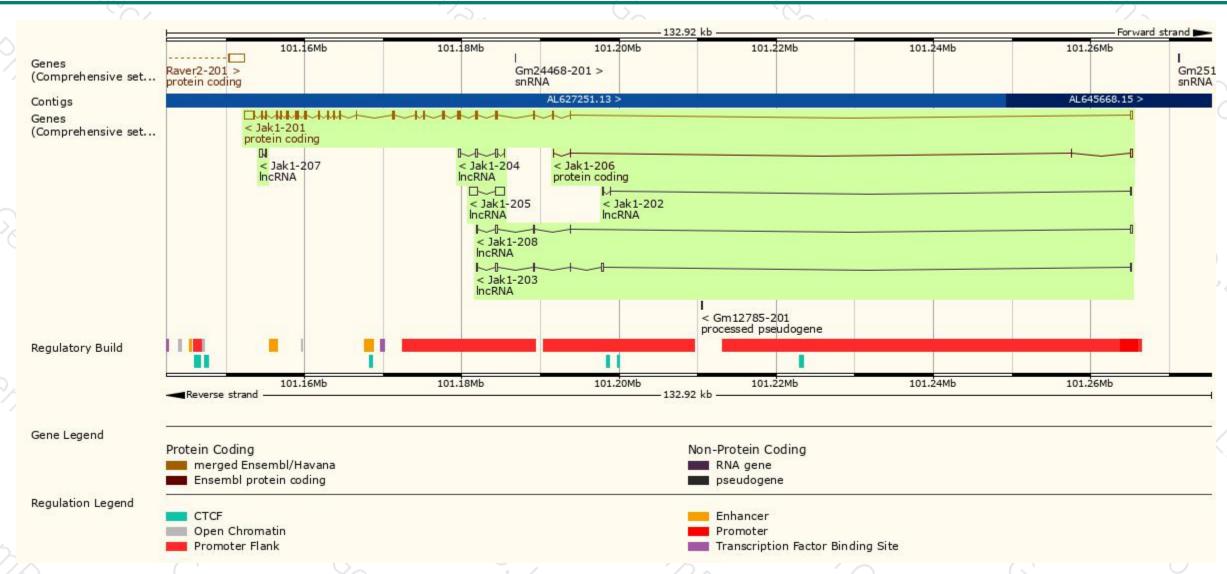


Reverse strand —

112.92 kb

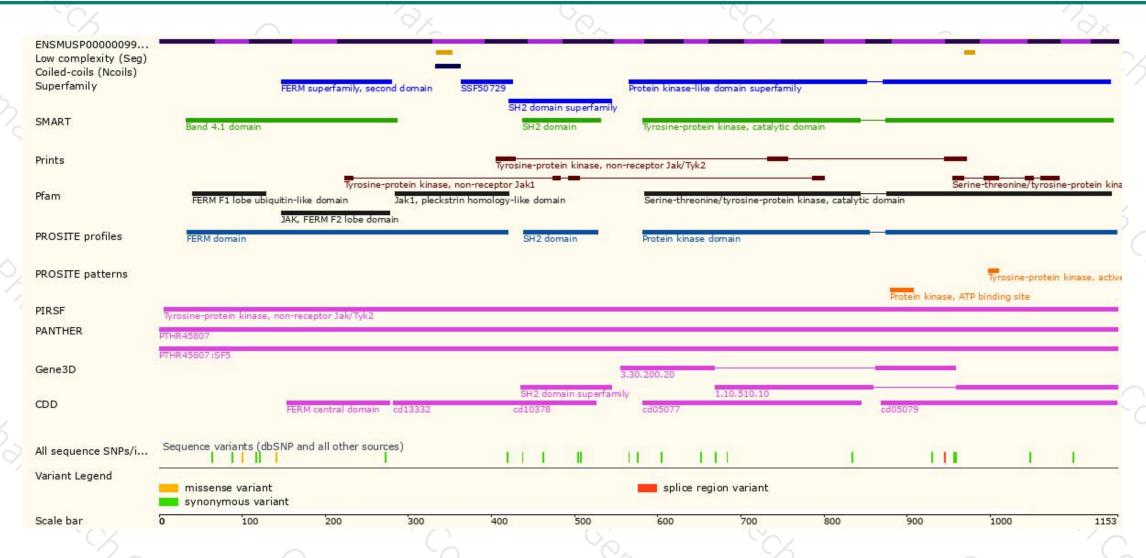
Genomic location distribution





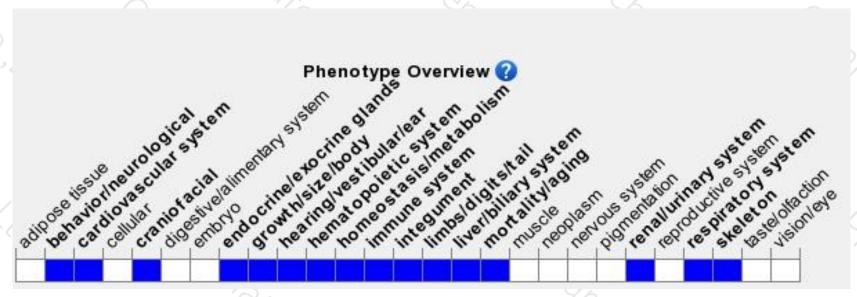
Protein domain





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 disruption of this gene die within the first 24 hours after birth.



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





