

Rock1 Cas9-CKO Strategy

Designer: Daohua Xu

Reviewer: Huimin Su

Design Date: 2019-9-28

Project Overview



Project Name

Rock1

Project type

Cas9-CKO

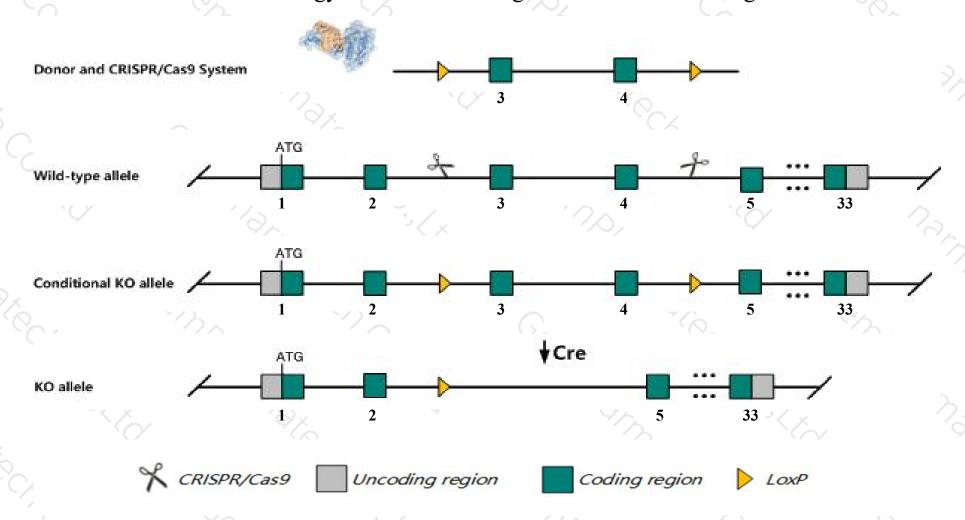
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Rock1* gene. The schematic diagram is as follows:



Technical routes



- The *Rock1* gene has 8 transcripts. According to the structure of *Rock1* gene, exon3-exon4 of *Rock1-201* (ENSMUST00000067947.6) transcript is recommended as the knockout region. The region contains 239bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Rock1* 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, Homozygous null mice have open eyes at birth, omphalocele and most die soon after birth as a result of cannibalization by the mom. Survivors develop inflammation of the eyelid.

 Another homozygous mutant shows partial lethality around implantation and reduced cardiac fibrosis after pressure overload.
- The *Rock1* gene is located on the Chr18. 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)



Rock1 Rho-associated coiled-coil containing protein kinase 1 [Mus musculus (house mouse)]

Gene ID: 19877, updated on 9-Mar-2019

Summary

☆ ?

Official Symbol Rock1 provided by MGI

Official Full Name Rho-associated coiled-coil containing protein kinase 1 provided by MGI

Primary source MGI:MGI:107927

See related Ensembl:ENSMUSG00000024290

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 1110055K06Rik, Rock-I

Expression Ubiquitous expression in bladder adult (RPKM 7.8), placenta adult (RPKM 6.0) and 28 other tissuesSee more

Orthologs human all

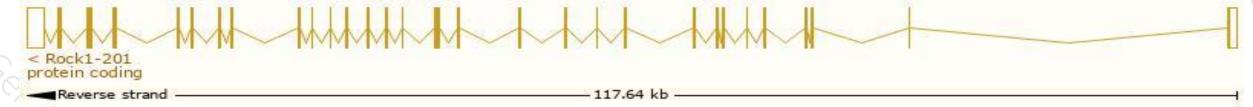
Transcript information (Ensembl)



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

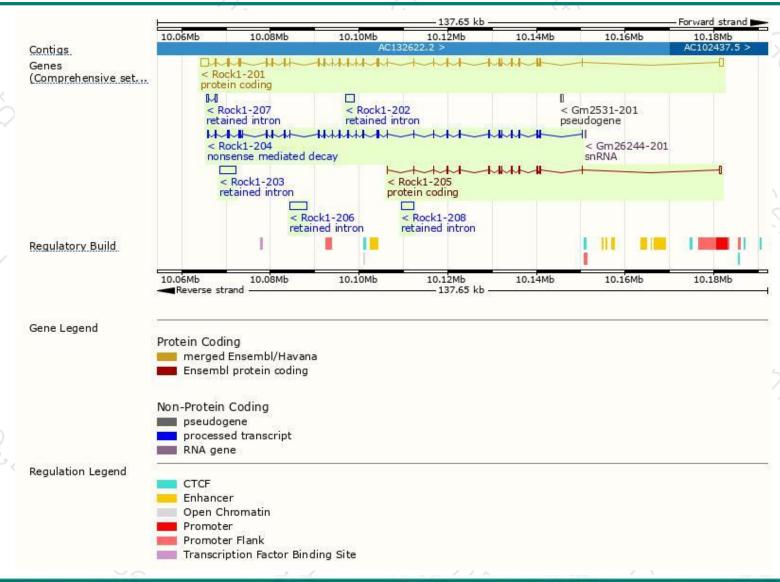
Name	Transcript ID	bn	Protein	Pietyne	ccps	UniProt	Flage
Name	Transcript ib	bp	Frotein	Biotype	CCDS	UIIIFIUL	Flags
Rock1-201	ENSMUST00000067947.6	6440	<u>1354aa</u>	Protein coding	CCDS29053	P70335	TSL:5 GENCODE basic APPRIS P1
Rock1-205	ENSMUST00000234734.1	1989	504aa	Protein coding	-	-8	CDS 3' incomplete
Rock1-204	ENSMUST00000234610.1	4320	<u>1089aa</u>	Nonsense mediated decay		-	
Rock1-206	ENSMUST00000234926.1	3840	No protein	Retained intron	72	20	
Rock1-203	ENSMUST00000234463.1	3494	No protein	Retained intron	-	-	
Rock1-208	ENSMUST00000235075.1	2735	No protein	Retained intron	-	*	
Rock1-202	ENSMUST00000234050.1	1877	No protein	Retained intron		-	
Rock1-207	ENSMUST00000235067.1	813	No protein	Retained intron	72	20	

The strategy is based on the design of *Rock1-201* transcript, The transcription is shown below



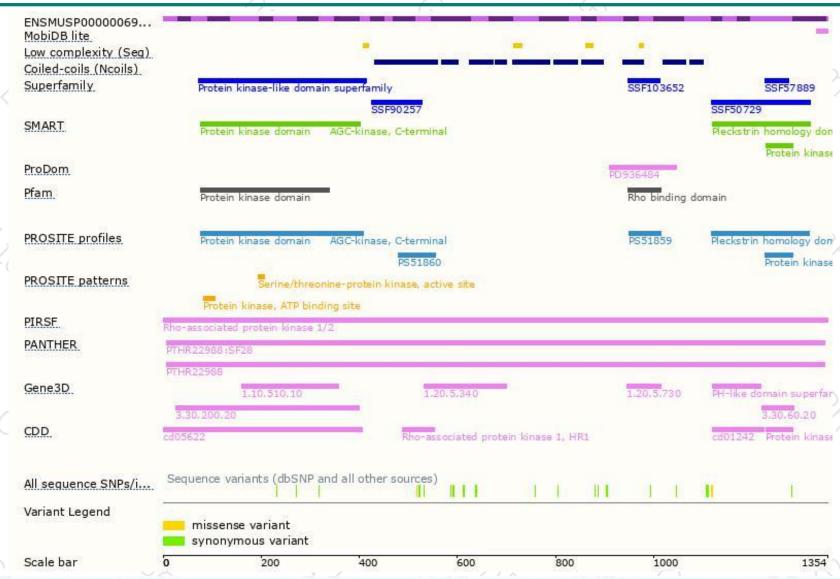
Genomic location distribution





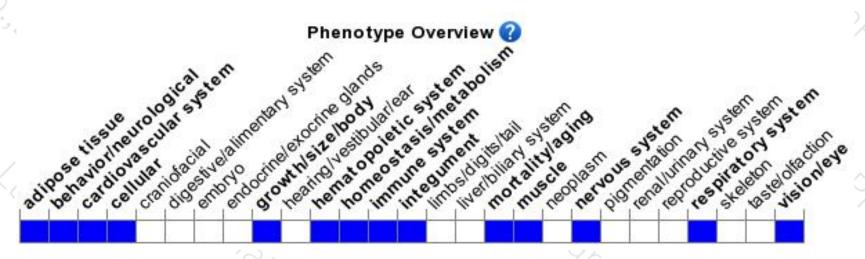
Protein domain





Mouse phenotype description(MGI)





Phenotypes affected by the gene are marked in blue.Data quoted from MGI database(http://www.informatics.jax.org/).

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





