

Smc1a Cas9-CKO Strategy

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Project Overview



Project Name

Smc1a

Project type

Cas9-CKO

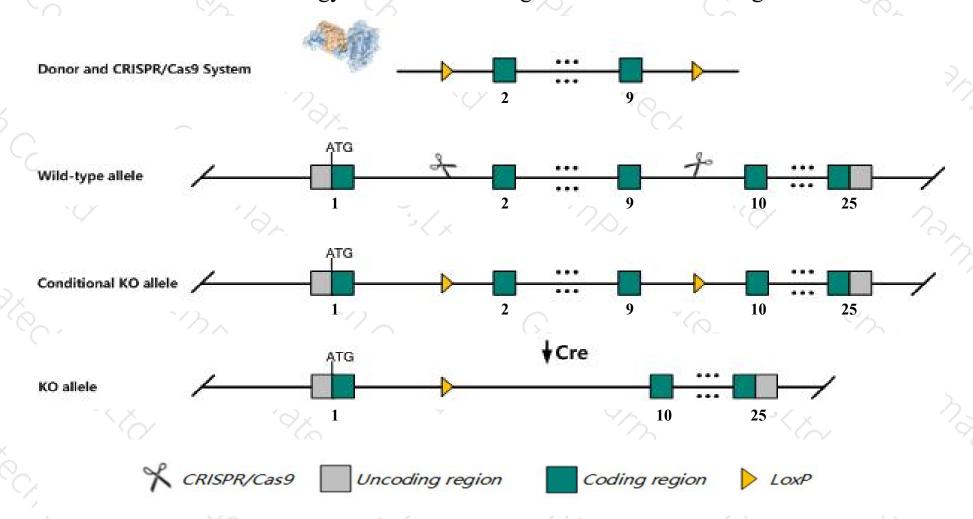
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Smc1a* gene. The schematic diagram is as follows:



Technical routes



- The *Smc1a* gene has 6 transcripts. According to the structure of *Smc1a* gene, exon2-exon9 of *Smc1a-201* (ENSMUST00000045312.5) transcript is recommended as the knockout region. The region contains 1436bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Smc1a* 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 a disruption in this gene display increased chromosomal instability, decreased cell survival, and defective S-phase checkpoint after ionizing radiation exposure.
- The *Smc1a* gene is located on the ChrX. 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)



Smc1a structural maintenance of chromosomes 1A [Mus musculus (house mouse)]

Gene ID: 24061, updated on 31-Jan-2019

Summary

☆ ?

Official Symbol Smc1a provided by MGI

Official Full Name structural maintenance of chromosomes 1A provided by MGI

Primary source MGI:MGI:1344345

See related Ensembl:ENSMUSG00000041133

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 5830426l24Rik, SMC-1A, Sb1.8, Smc1, Smc1alpha, Smc1l1, Smcb, mKlAA0178

Expression Ubiquitous expression in CNS E11.5 (RPKM 47.7), liver E14 (RPKM 30.8) and 27 other tissuesSee more

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

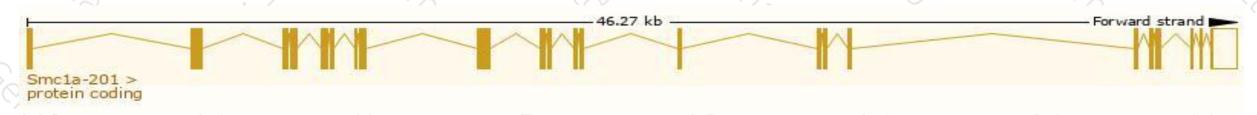
Transcript information (Ensembl)



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

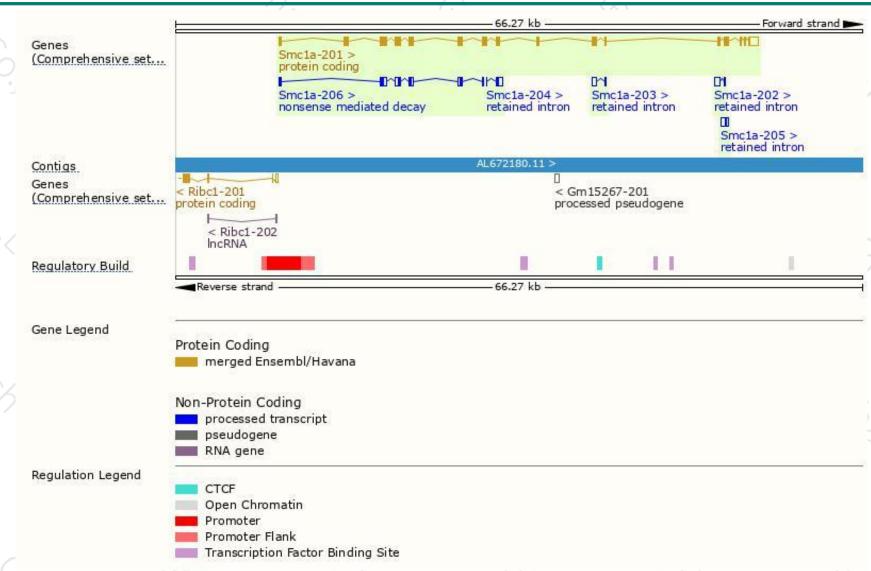
Transcript ID	-					
. rancompt to	bp	Protein	Biotype	CCDS	UniProt	Flags
ENSMUST00000045312.5	4691	<u>1233aa</u>	Protein coding	CCDS30473	Q9CU62	TSL:1 GENCODE basic APPRIS P1
ENSMUST00000145518.7	1824	<u>54aa</u>	Nonsense mediated decay	-	S4R179	TSL:1
ENSMUST00000124681.1	620	No protein	Retained intron		2	TSL:2
ENSMUST00000135172.1	617	No protein	Retained intron	92	2	TSL:3
ENSMUST00000141457.1	591	No protein	Retained intron	-		TSL:3
ENSMUST00000131395.1	499	No protein	Retained intron	-	-	TSL:3
	ENSMUST00000145518.7 ENSMUST00000124681.1 ENSMUST00000135172.1 ENSMUST00000141457.1	ENSMUST00000145518.7 1824 ENSMUST00000124681.1 620 ENSMUST00000135172.1 617 ENSMUST00000141457.1 591	ENSMUST00000145518.7 1824 54aa ENSMUST00000124681.1 620 No protein ENSMUST00000135172.1 617 No protein ENSMUST00000141457.1 591 No protein	ENSMUST00000145518.7 1824 54aa Nonsense mediated decay ENSMUST00000124681.1 620 No protein Retained intron ENSMUST00000135172.1 617 No protein Retained intron ENSMUST00000141457.1 591 No protein Retained intron	ENSMUST00000145518.7 1824 54aa Nonsense mediated decay - ENSMUST00000124681.1 620 No protein Retained intron - ENSMUST00000135172.1 617 No protein Retained intron - ENSMUST00000141457.1 591 No protein Retained intron - ENSMUST00000131395.1 499 No protein Retained intron -	ENSMUST00000145518.7 1824 54aa Nonsense mediated decay - S4R179 ENSMUST00000124681.1 620 No protein Retained intron - - ENSMUST00000135172.1 617 No protein Retained intron - - ENSMUST00000141457.1 591 No protein Retained intron - -

The strategy is based on the design of Smc1a-201 transcript, The transcription is shown below



Genomic location distribution





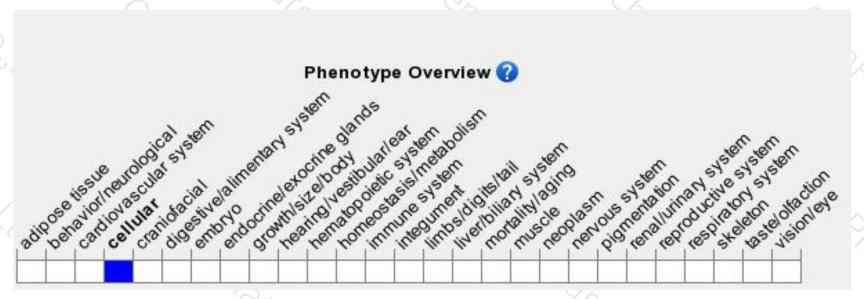
Protein domain





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 disruption in this gene display increased chromosomal instability, decreased cell survival, and defective S-phase checkpoint after ionizing radiation exposure.



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





