

Ascc1 Cas9-CKO Strategy

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Project Overview

Project Name

Ascc1

Project type

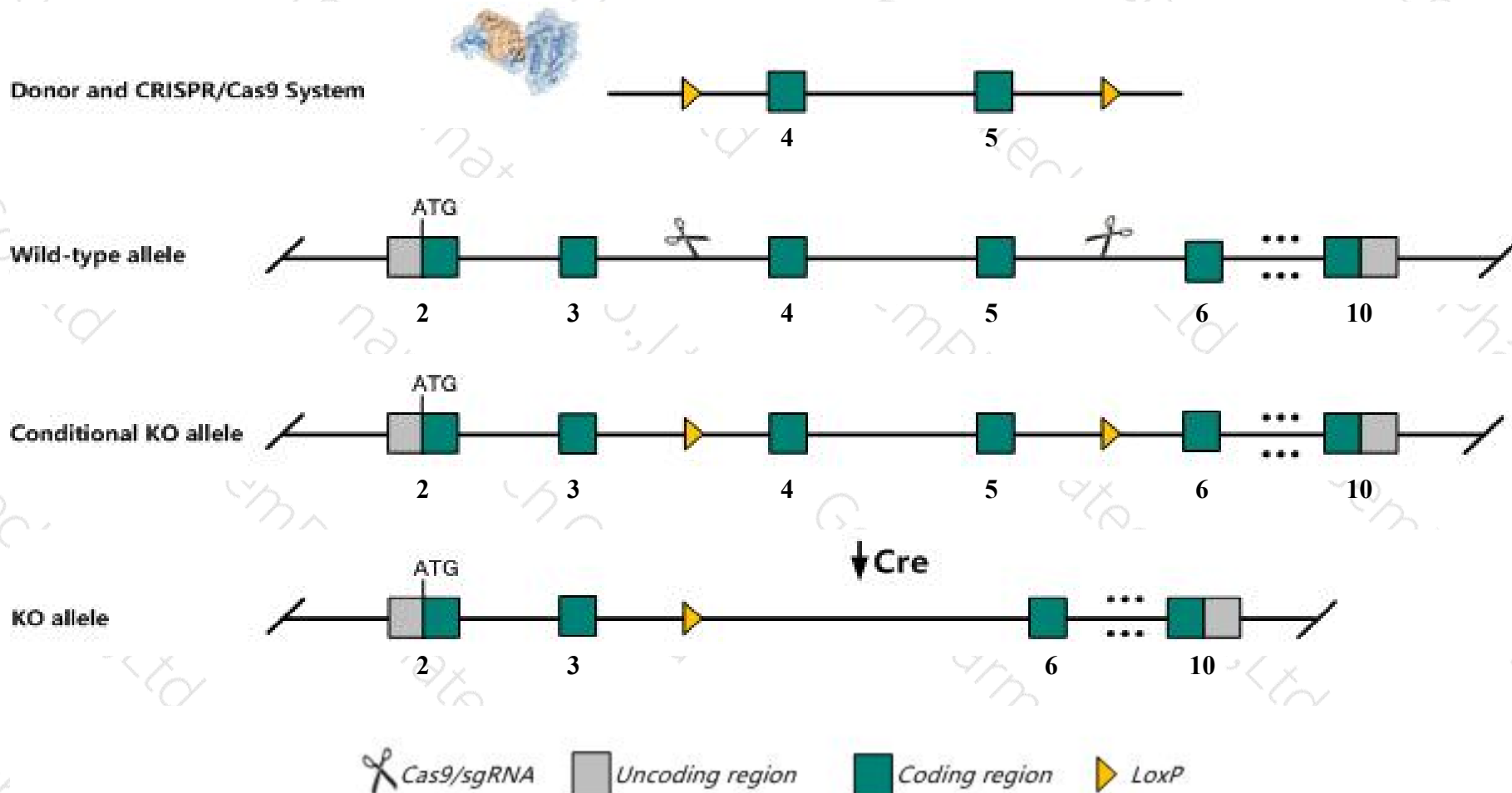
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Ascc1* gene. The schematic diagram is as follows:



- The *Ascc1* gene has 3 transcripts. According to the structure of *Ascc1* gene, exon4-exon5 of *Ascc1*-202(ENSMUST00000164083.3) transcript is recommended as the knockout region. The region contains 277bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Ascc1* 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.

- The *Ascc1* gene is located on the Chr10. 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)

Ascc1 activating signal cointegrator 1 complex subunit 1 [*Mus musculus* (house mouse)]

Gene ID: 69090, updated on 26-Jun-2020

Summary

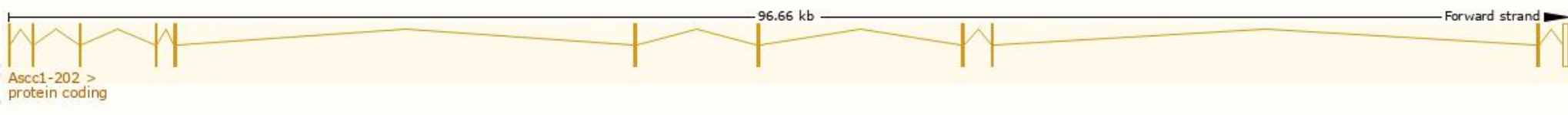
Official Symbol	Ascc1 provided by MGI
Official Full Name	activating signal cointegrator 1 complex subunit 1 provided by MGI
Primary source	MGI:MGI:1916340
See related	Ensembl:ENSMUSG00000044475
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	CGI-18; ASC1p50; AI550520; 1810015P09Rik
Expression	Ubiquitous expression in testis adult (RPKM 16.8), kidney adult (RPKM 16.1) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

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

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ascc1-202	ENSMUST00000164083.3	1473	356aa	Protein coding	CCDS23870	Q9D8Z1	TSL:1 GENCODE basic APPRIS P1
Ascc1-201	ENSMUST00000050516.13	1450	356aa	Protein coding	CCDS23870	Q9D8Z1	TSL:1 GENCODE basic APPRIS P1
Ascc1-203	ENSMUST00000219778.1	1230	No protein	Processed transcript	-	-	TSL:1

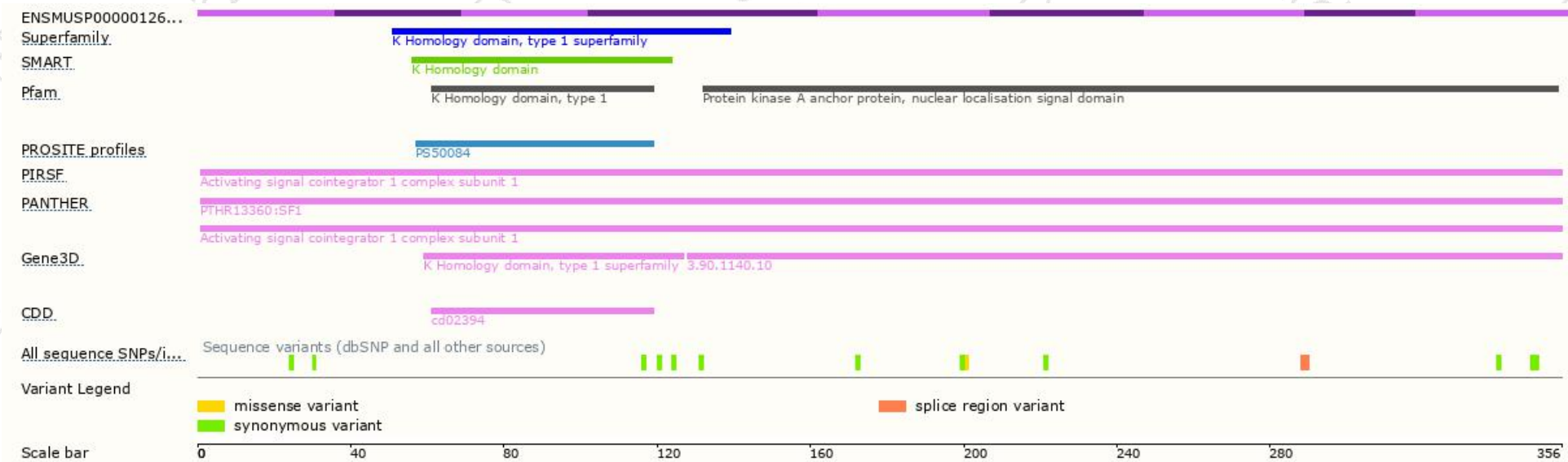
The strategy is based on the design of *Ascc1-202* transcript,the transcription is shown below:



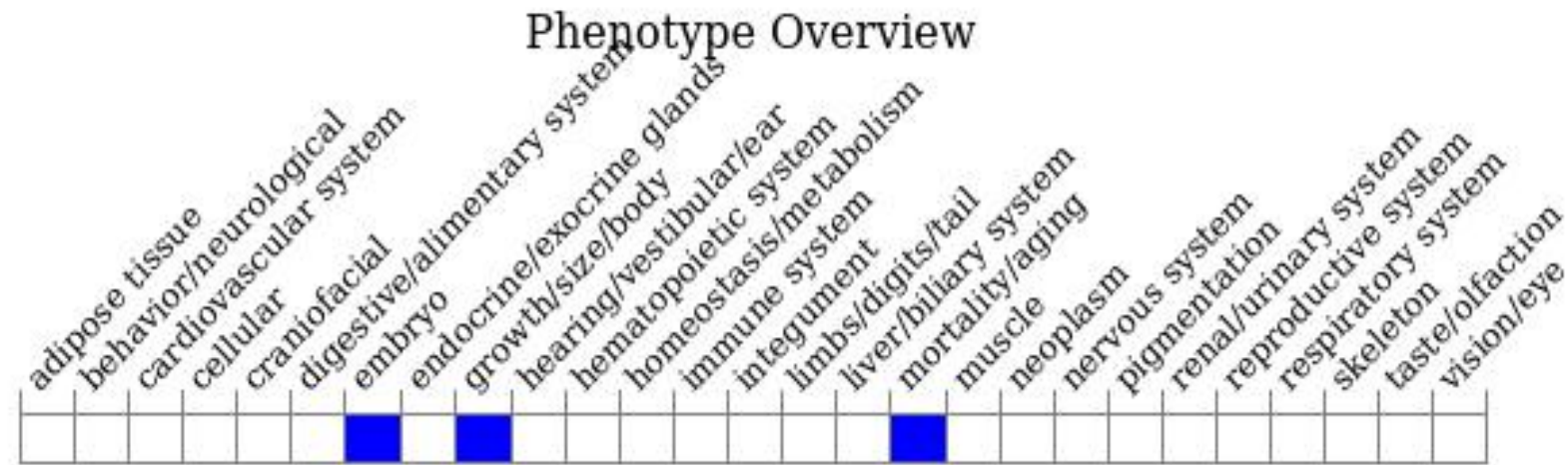
Genomic location distribution



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/>).

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

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