

Srd5a2 Cas9-CKO Strategy

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Reviewer:

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Design Date:

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Project Overview

Project Name

Srd5a2

Project type

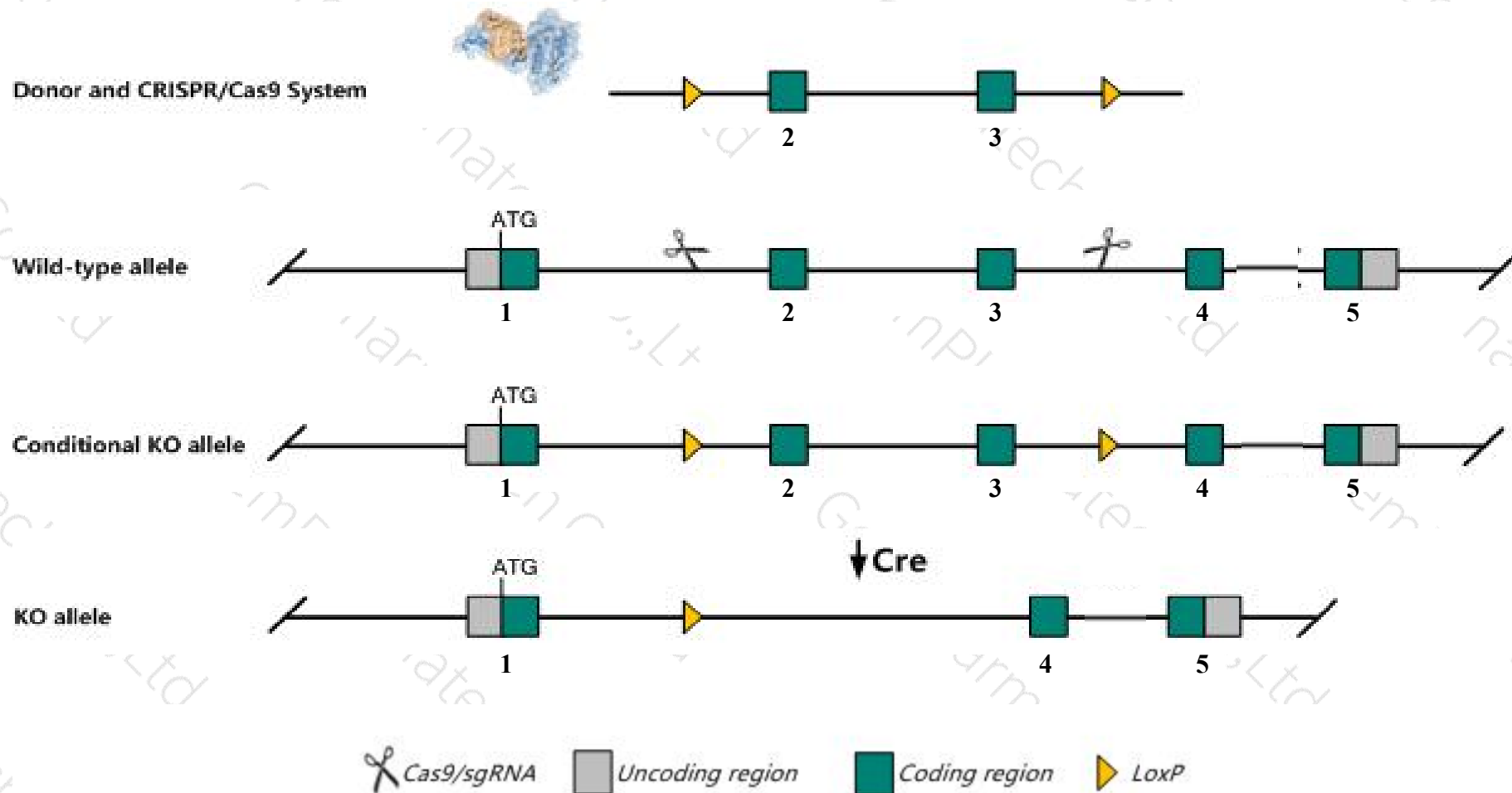
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Srd5a2* gene. The schematic diagram is as follows:



- The *Srd5a2* gene has 1 transcript. According to the structure of *Srd5a2* gene, exon2-exon3 of *Srd5a2*-201 (ENSMUST00000043458.8) transcript is recommended as the knockout region. The region contains 266bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Srd5a2* 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.

- According to the existing MGI data, Homozygous null mutant males are fertile, but have small prostates and seminal vesicles, elevated testosterone in reproductive tissue and decreased androgen-dependent gene expression. Mutant females exhibit no detectable abnormality.
- The flox region contain the Gm49880 gene, which may delete it after Cre.
- The *Srd5a2* gene is located on the Chr17. 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)

Srd5a2 steroid 5 alpha-reductase 2 [*Mus musculus* (house mouse)]

Gene ID: 94224, updated on 13-Mar-2020

Summary

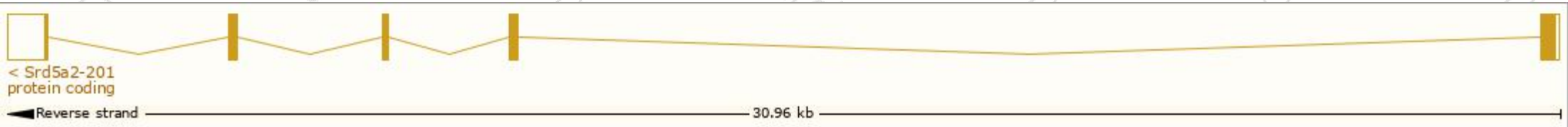
Official Symbol	Srd5a2 provided by MGI
Official Full Name	steroid 5 alpha-reductase 2 provided by MGI
Primary source	MGI:MGI:2150380
See related	Ensembl:ENSMUSG00000038541
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	5ART2; S5AR 2
Expression	Restricted expression toward adrenal adult (RPKM 248.6) See more
Orthologs	human all

Transcript information (Ensembl)

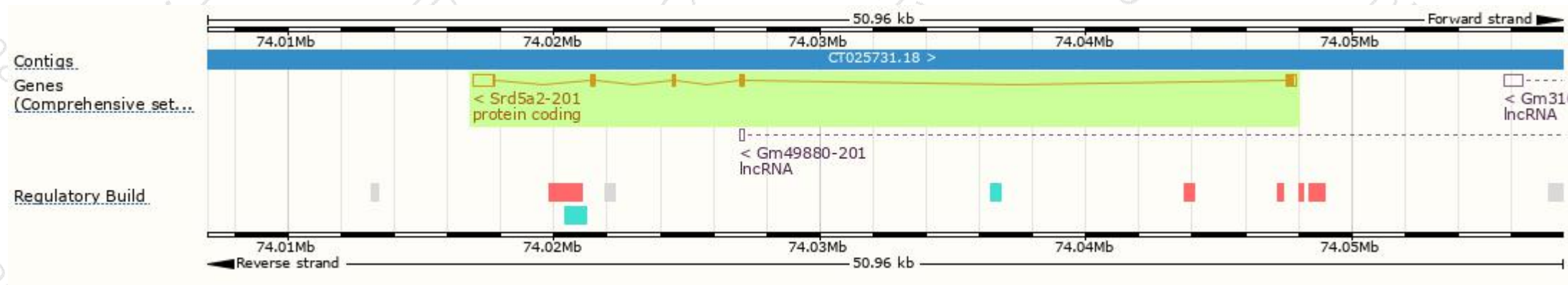
The gene has 1 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Srd5a2-201	ENSMUST00000043458.8	1605	254aa	Protein coding	CCDS28968	Q99N99	TSL:1 Gencode basic APPRIS P1

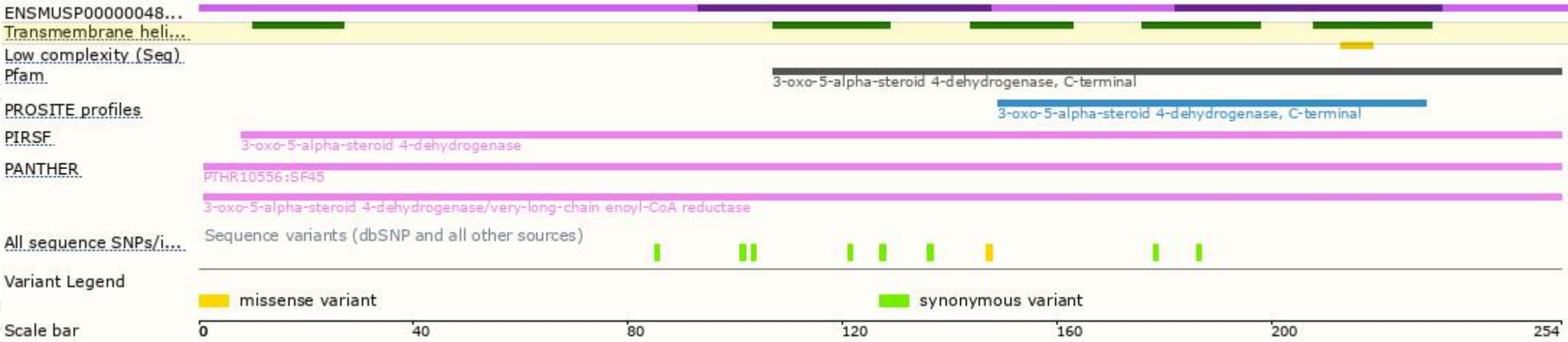
The strategy is based on the design of *Srd5a2-201* transcript,The transcription is shown below



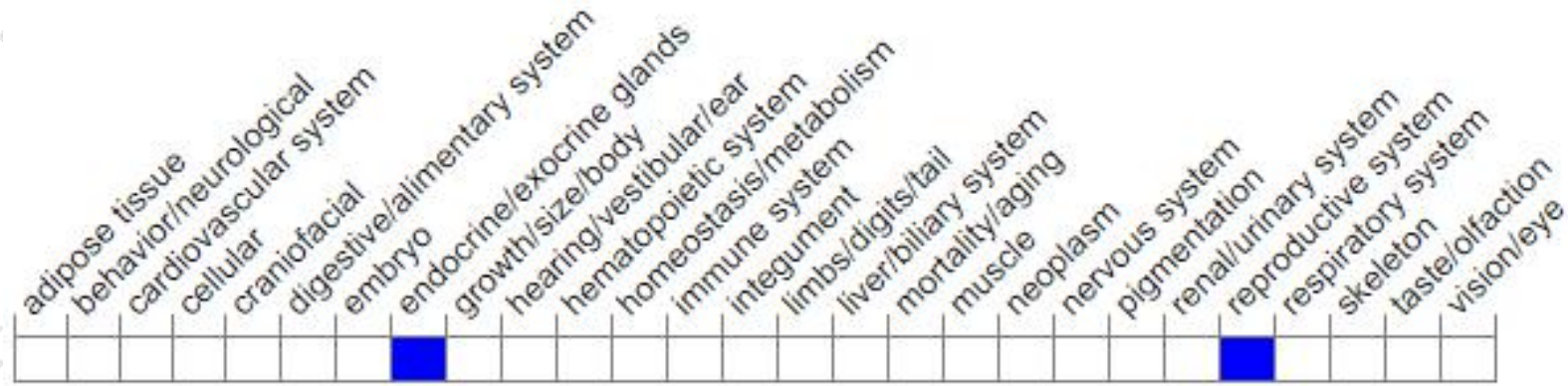
Genomic location distribution



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, Homozygous null mutant males are fertile, but have small prostates and seminal vesicles, elevated testosterone in reproductive tissue and decreased androgen-dependent gene expression. Mutant females exhibit no detectable abnormality.

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

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