

Scgb3a2 Cas9-CKO Strategy

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Design Date:

2019-9-17

Project Overview

Project Name

Scgb3a2

Project type

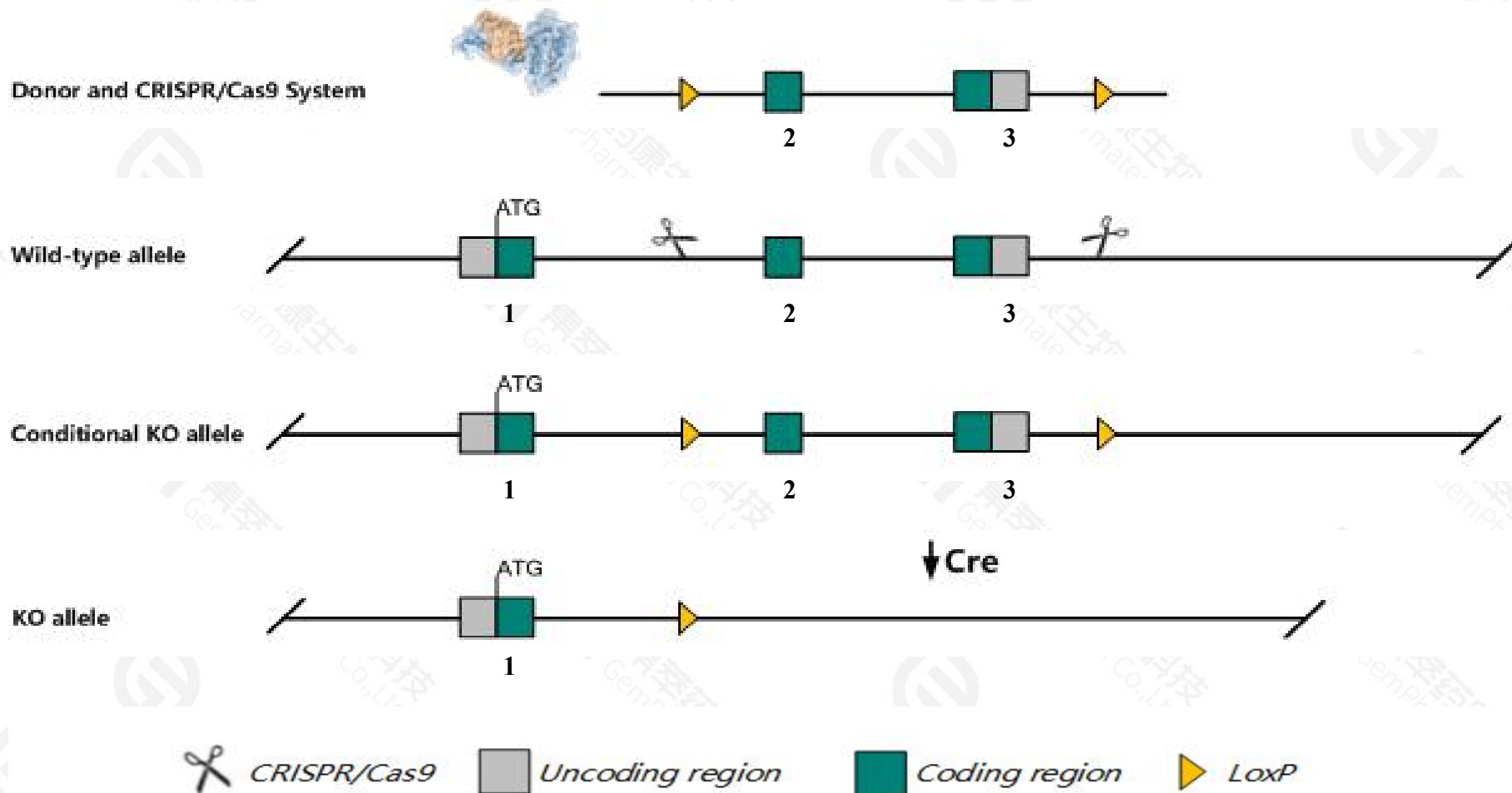
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Scgb3a2* gene. The schematic diagram is as follows:



- The *Scgb3a2* gene has 3 transcripts. According to the structure of *Scgb3a2* gene, exon2-exon3 of *Scgb3a2*-203(ENSMUST00000189750.1) transcript is recommended as the knockout region. The region contains 221bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Scgb3a2* 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, mice homozygous for a knock-out allele exhibit background sensitive lung inflammatory response to ovalbumin exposure.
- The *Scgb3a2* 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)

Scgb3a2 secretoglobin, family 3A, member 2 [Mus musculus (house mouse)]

Gene ID: 117158, updated on 17-Feb-2021

Summary



Official Symbol Scgb3a2 provided by [MGI](#)

Official Full Name secretoglobin, family 3A, member 2 provided by [MGI](#)

Primary source [MGI:MGI:2153470](#)

See related [Ensembl:ENSMUSG00000038791](#)

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 LuLe, LuLeu1, Pnsp1, UGR, UGRP1, Utgrp1

Expression Restricted expression toward lung adult (RPKM 559.3)[See more](#)

Orthologs [human](#) [all](#)

Transcript information (Ensembl)

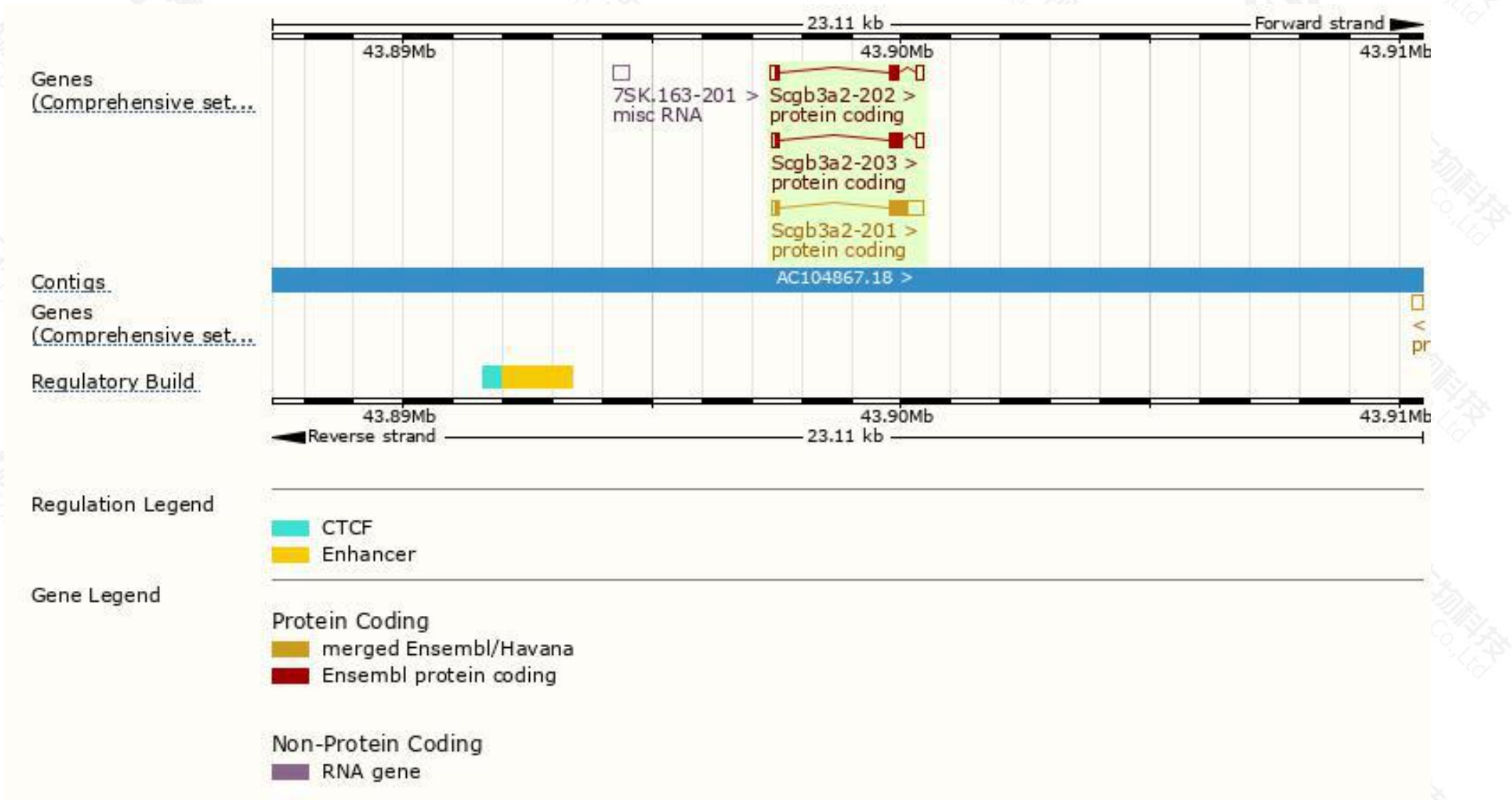
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Scgb3a2-203	ENSMUST00000189750.2	578	113aa	Protein coding	CCDS79641		TSL:1 , GENCODE basic ,
Scgb3a2-202	ENSMUST00000187157.7	550	91aa	Protein coding	CCDS79640		TSL:1 , GENCODE basic , APPRIS P1 ,
Scgb3a2-201	ENSMUST00000043803.13	842	139aa	Protein coding	-		TSL:1 , GENCODE basic ,

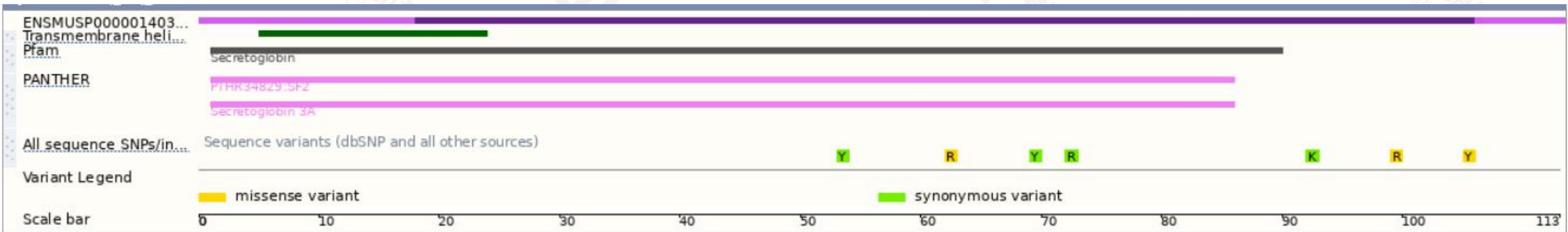
The strategy is based on the design of *Scgb3a2-203* transcript,the transcription is shown below:



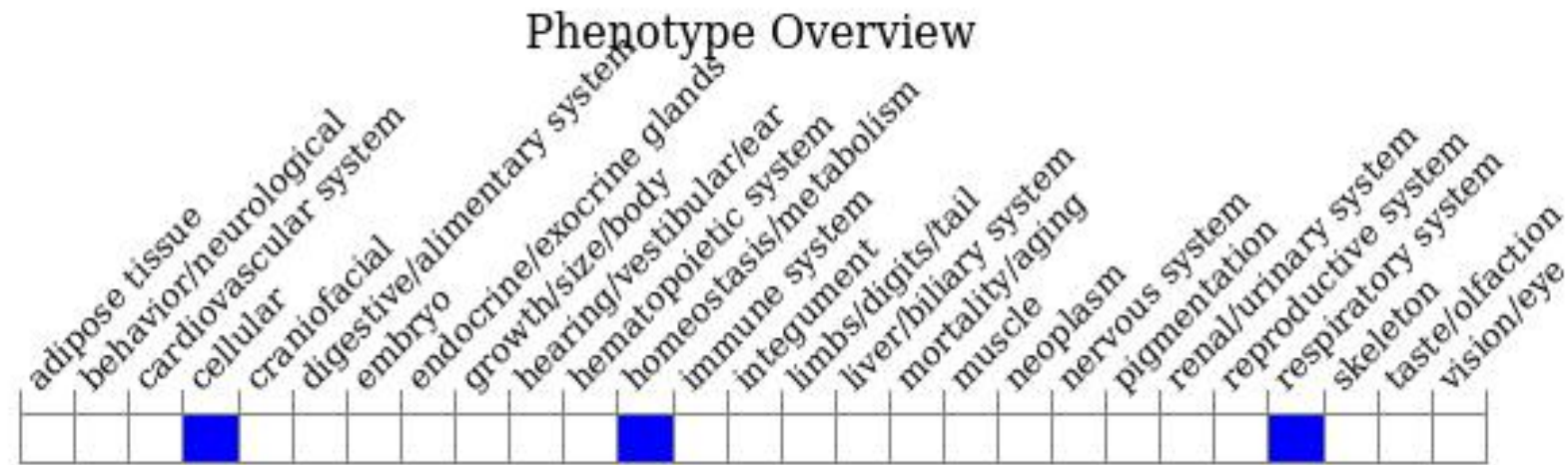
Genomic location distribution



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 knock-out allele exhibit background sensitive lung inflammatory response to ovalbumin exposure.

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

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