

Agr3 Cas9-CKO Strategy

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Project Overview

Project Name

Agr3

Project type

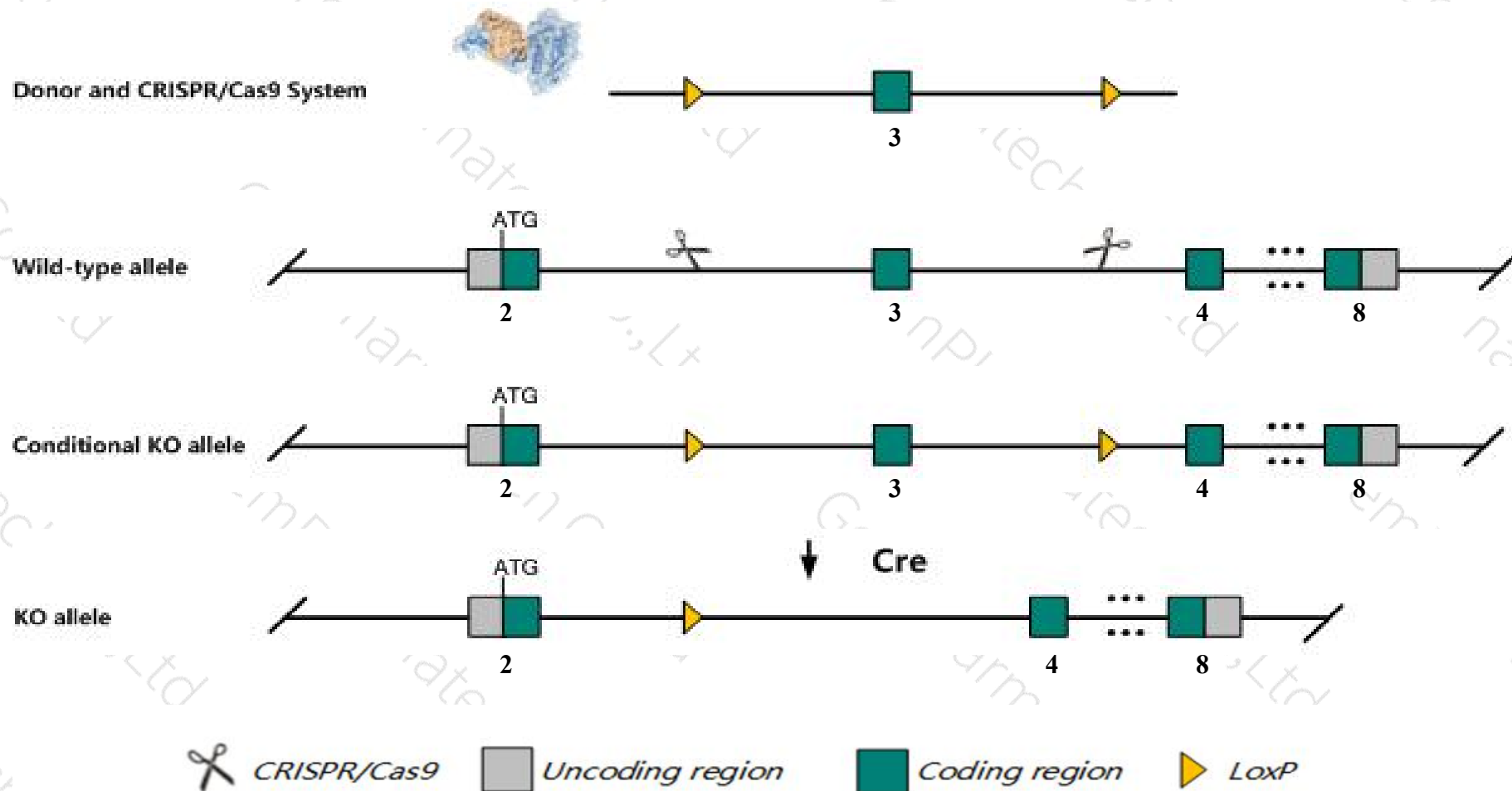
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Agr3* gene. The schematic diagram is as follows:



- The *Agr3* gene has 2 transcripts. According to the structure of *Agr3* gene, exon3 of *Agr3-201* (ENSMUST00000042101.4) transcript is recommended as the knockout region. The region contains 64bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Agr3* 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

- The effect on transcript *Agr3*-202 is unknown.
- The *Agr3* gene is located on the Chr12. 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)

Agr3 anterior gradient 3 [*Mus musculus* (house mouse)]

Gene ID: 403205, updated on 12-Aug-2019

Summary

Official Symbol Agr3 provided by [MGI](#)
Official Full Name anterior gradient 3 provided by [MGI](#)
Primary source [MGI:MGI:2685734](#)
See related [Ensembl:ENSMUSG00000036231](#)
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 Gm888; BCMP11; E030025L21Rik
Expression Low expression observed in reference dataset [See more](#)
Orthologs [human](#) [all](#)

Genomic context

Location: 12; 12 A3

See Agr3 in [Genome Data Viewer](#)

Exon count: 10

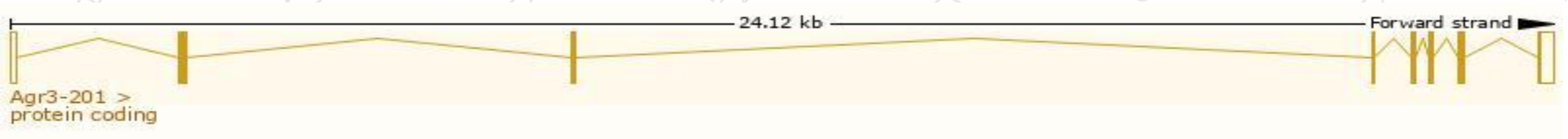
Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	12	NC_000078.6 (35925621..35949737)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	12	NC_000078.5 (36652208..36676317)

Transcript information (Ensembl)

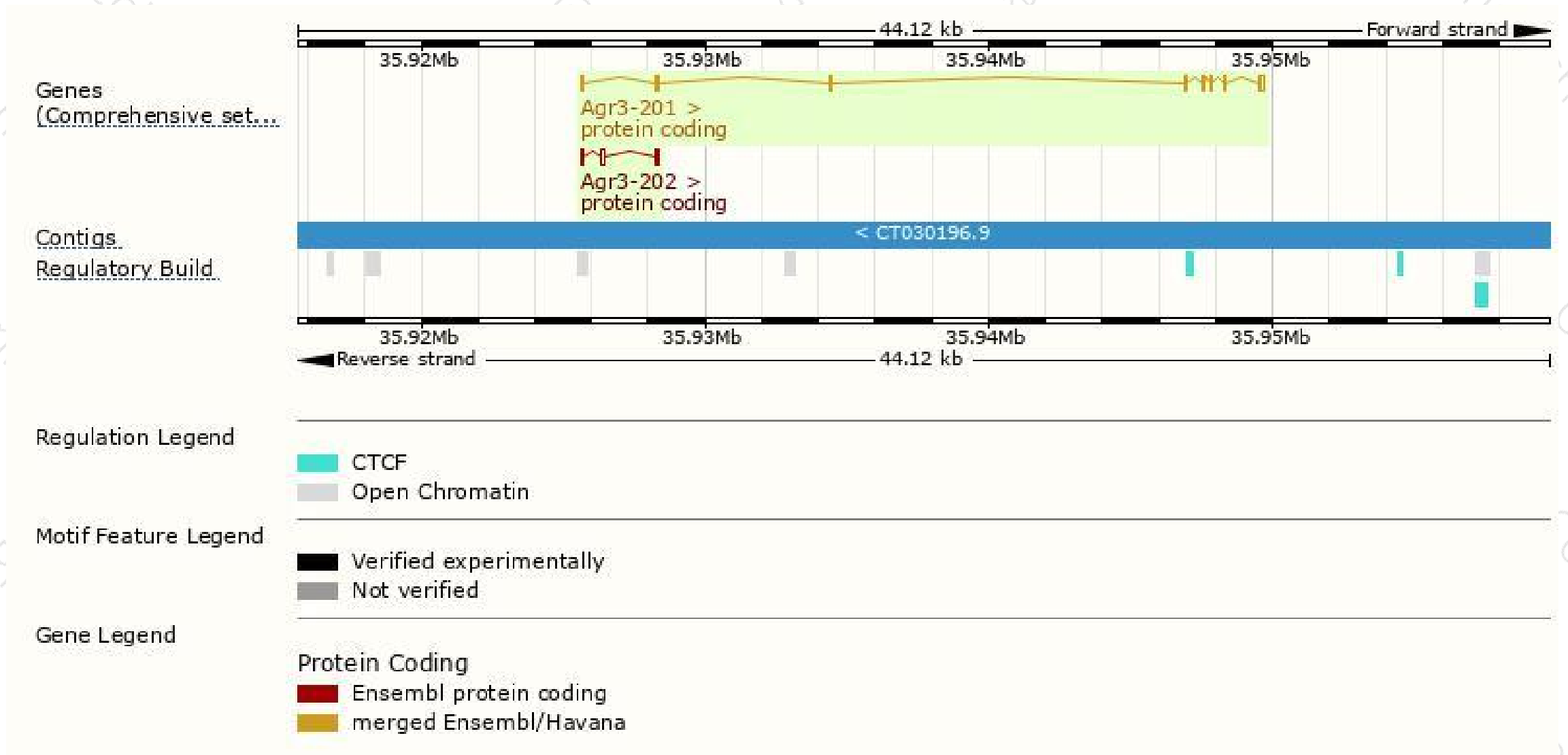
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Agr3-201	ENSMUST00000042101.4	826	165aa	Protein coding	CCDS25880	Q8R3W7	TSL:1 GENCODE basic APPRIS P1
Agr3-202	ENSMUST00000154042.1	388	35aa	Protein coding	-	A7NSI2	CDS 3' incomplete TSL:2

The strategy is based on the design of *Agr3-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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