Duox1 Cas9-KO Strategy

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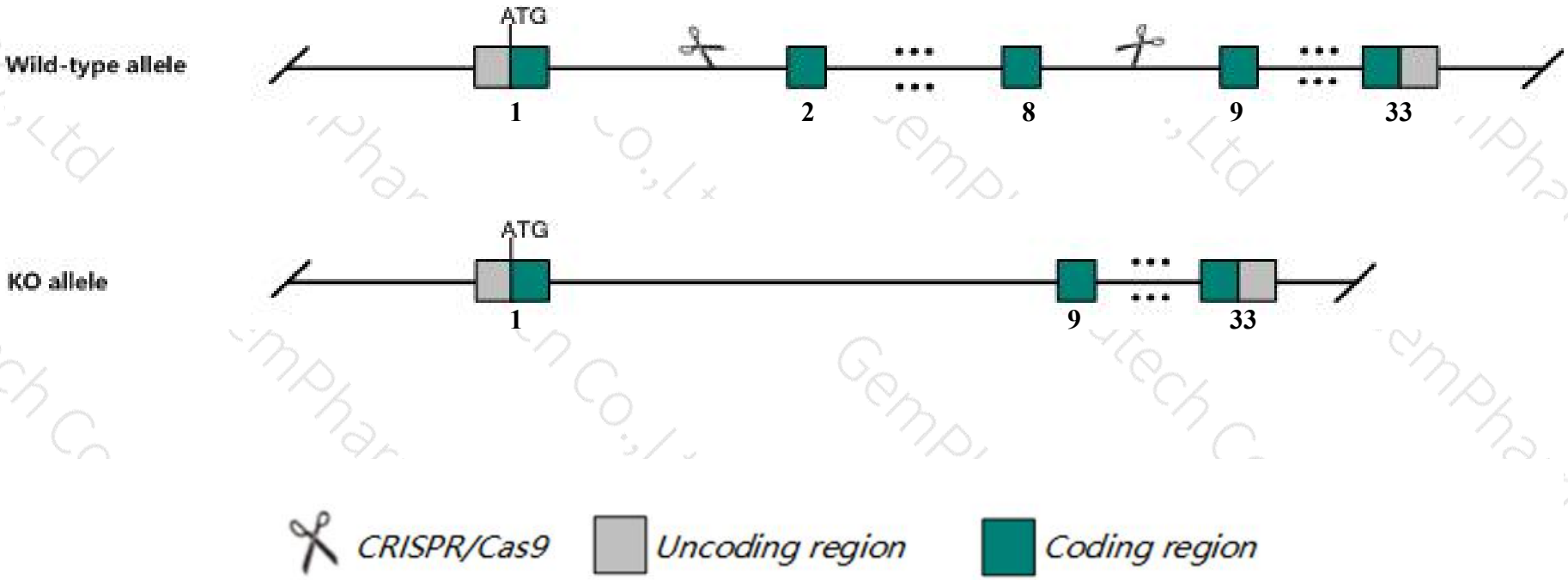
Project Overview

Project Name	***Duox1***
Project type	**Cas9-KO**
Strain background	**C57BL/6JGpt**

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Duox1* gene. The schematic diagram is as follows:



Technical routes

➢ The *Duox1* gene has 1 transcript. According to the structure of *Duox1* gene, exon2-exon8 of *Duox1-201* (ENSMUST00000099461.3) transcript is recommended as the knockout region.The region contains 964bp coding sequence. Knock out the region will result in disruption of protein function.

➢ In this project we use CRISPR/Cas9 technology to modify *Duox1* gene. The brief process is as follows: CRISPR/Cas9 system

Notice

➢The knockout region is near to the N-terminal of *Duoxa1* gene,this strategy may influence the regulatory function of the N-terminal of *Duoxa1* gene.

➢ The *Duox1* gene is located on the Chr2. 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information（NCBI）

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Duox1 dual oxidase 1 [*Mus musculus* (house mouse)]

Gene ID: 99439, updated on 5-Nov-2019

Summary

Official Symbol	Duox1 provided by MGI
Official Full Name	dual oxidase 1 provided by MGI
Primary source	MGI:MGI:2139422
See related	Ensembl:ENSMUSG00000033268
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	Duox2; LNOX1; LNOX2; THOX1; THOX2; NOXEF1; NOXEF2; AW987690; P138-TOX; 9930101G15Rik
Expression	Biased expression in stomach adult (RPKM 3.3), bladder adult (RPKM 2.4) and 9 other tissues See more
Orthologs	human all

Genomic context

Location: 2; 2 E5

See Duox1 in Genome Data Viewer

Exon count: 34

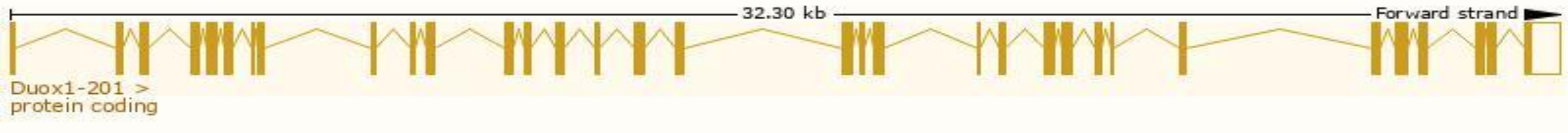
Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	2	NC_000068.7 (122313010..122347972)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	2	NC_000068.6 (122141408..122173708)

Transcript information（Ensembl）

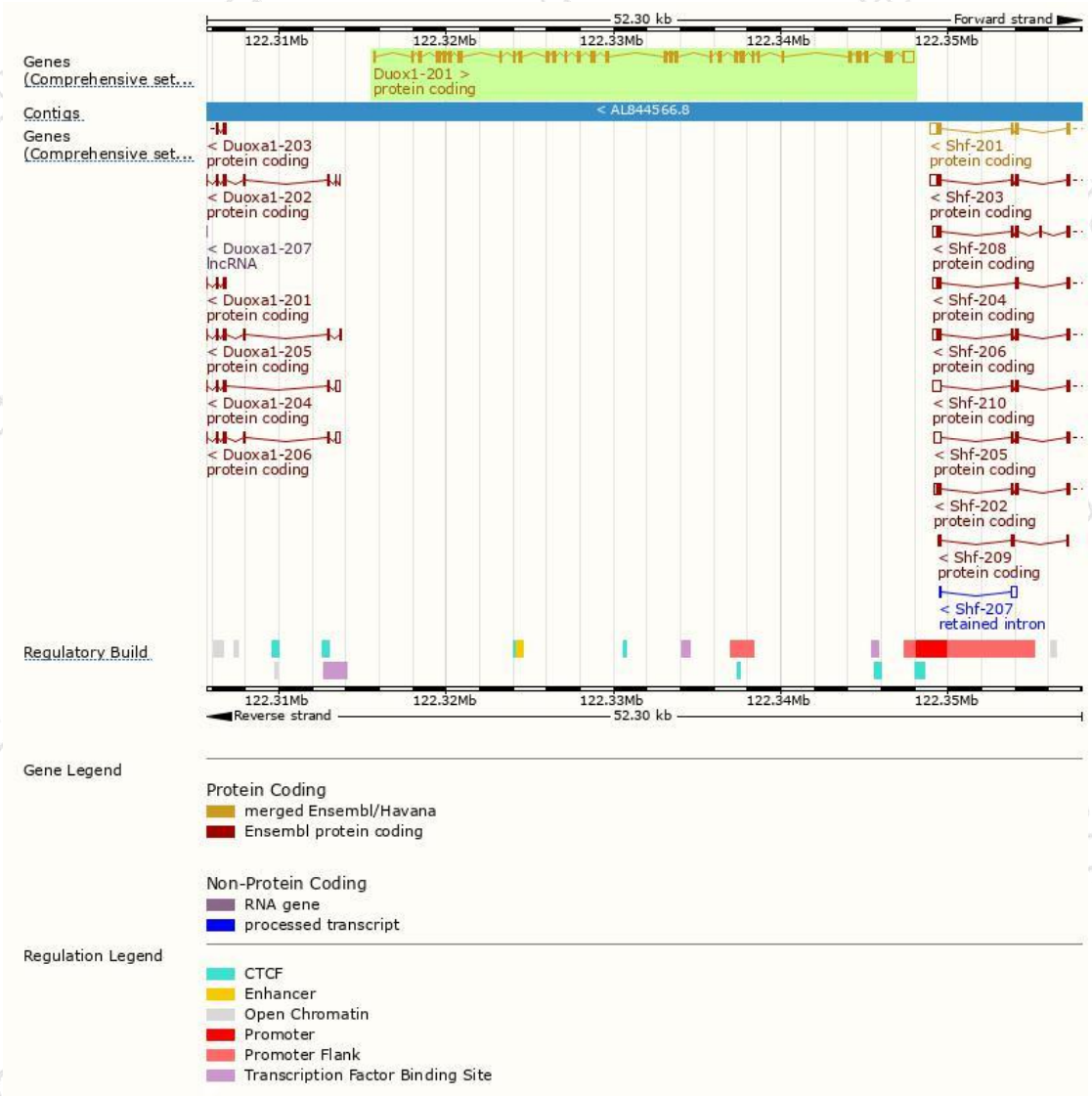
The gene has 1 transcript,and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Duox1-201	ENSMUST00000099461.3	5267	1551aa	Protein coding	CCDS38222	A2AQ92	TSL:2 GENCODE basic APPRIS P1

The strategy is based on the design of *Duox1-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain

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

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