

Defb26 Cas9-KO Strategy

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Project Overview

Project Name

Defb26

Project type

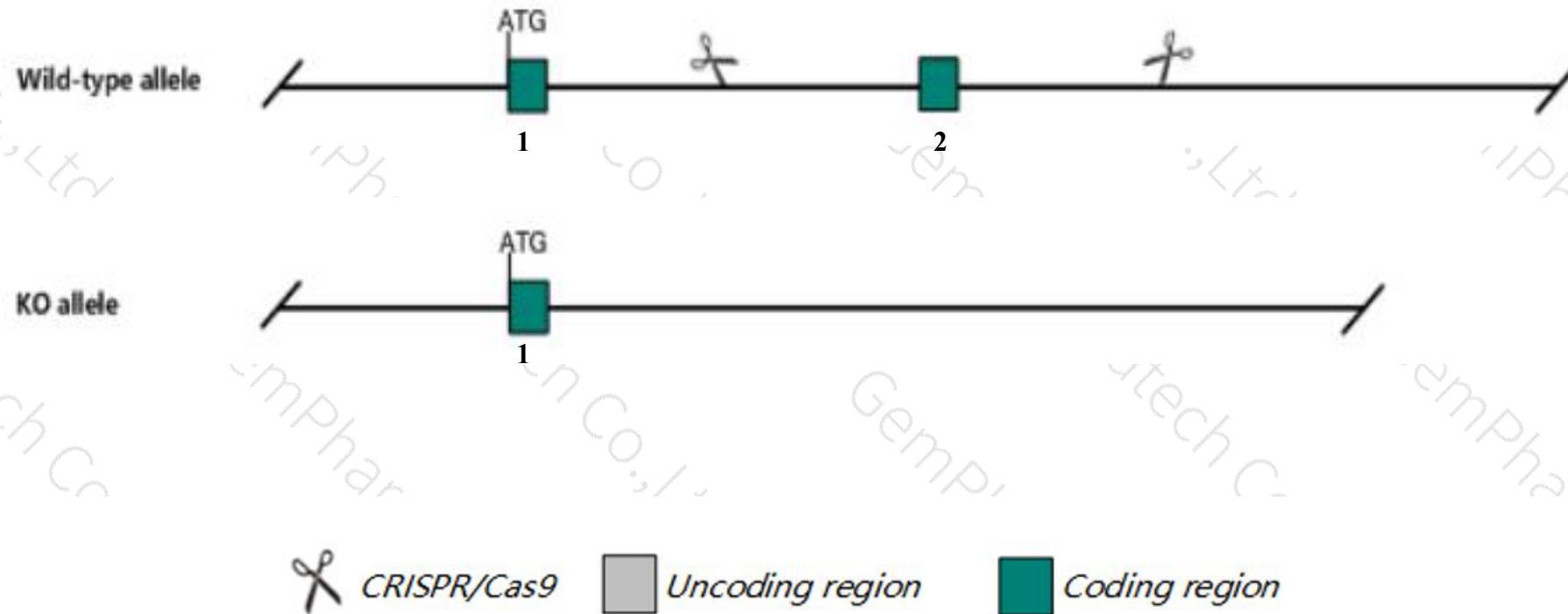
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Defb26* gene. The schematic diagram is as follows:



- The *Defb26* gene has 1 transcript. According to the structure of *Defb26* gene, exon2 of *Defb26-201*(ENSMUST00000099205.1) transcript is recommended as the knockout region. The region contains 503bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Defb26* gene. The brief process is as follows: CRISPR/Cas9 system 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 *Defb26* 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)

Defb26 defensin beta 26 [Mus musculus (house mouse)]

Gene ID: 654457, updated on 13-Mar-2020

Summary



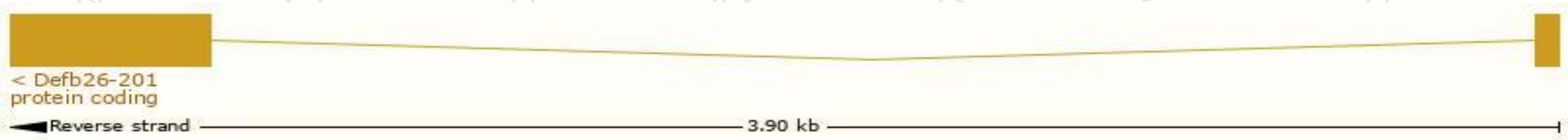
Official Symbol	Defb26 provided by MGI
Official Full Name	defensin beta 26 provided by MGI
Primary source	MGI:MGI:3643488
See related	Ensembl:ENSMUSG00000074680
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	EG654457
Expression	Restricted expression toward genital fat pad adult (RPKM 53.5) See more
Orthologs	human all

Transcript information (Ensembl)

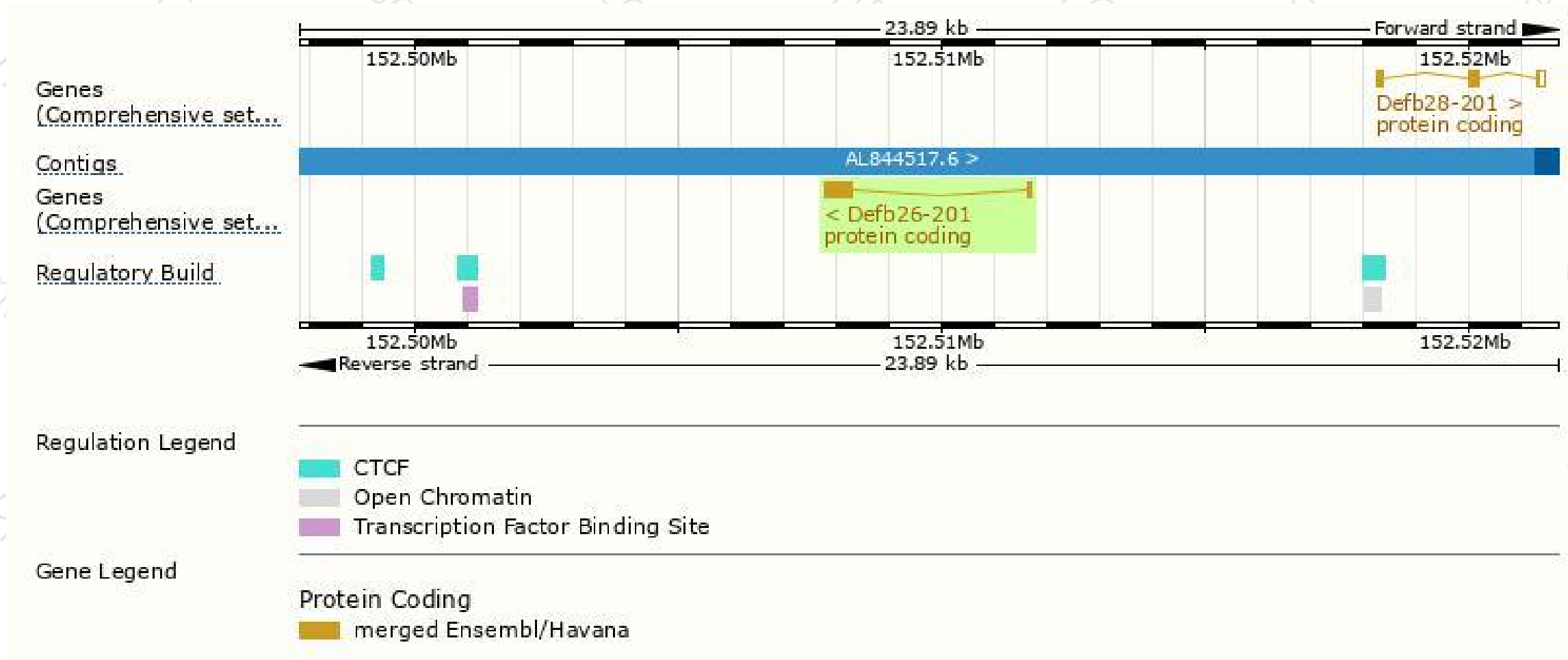
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Defb26-201	ENSMUST00000099205.1	561	186aa	Protein coding	CCDS16888	Q30KN7	TSL:1 GENCODE basic APPRIS P1

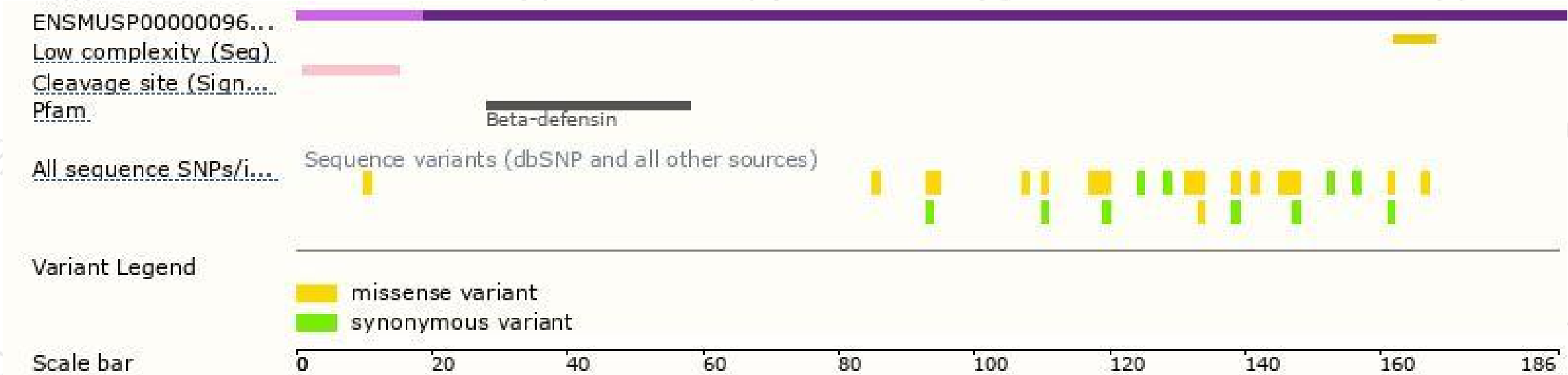
The strategy is based on the design of *Defb26-201* transcript, the transcription is shown below:



Genomic location distribution



Protein domain



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

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