

Med26 Cas9-CKO Strategy

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Project Overview

Project Name

Med26

Project type

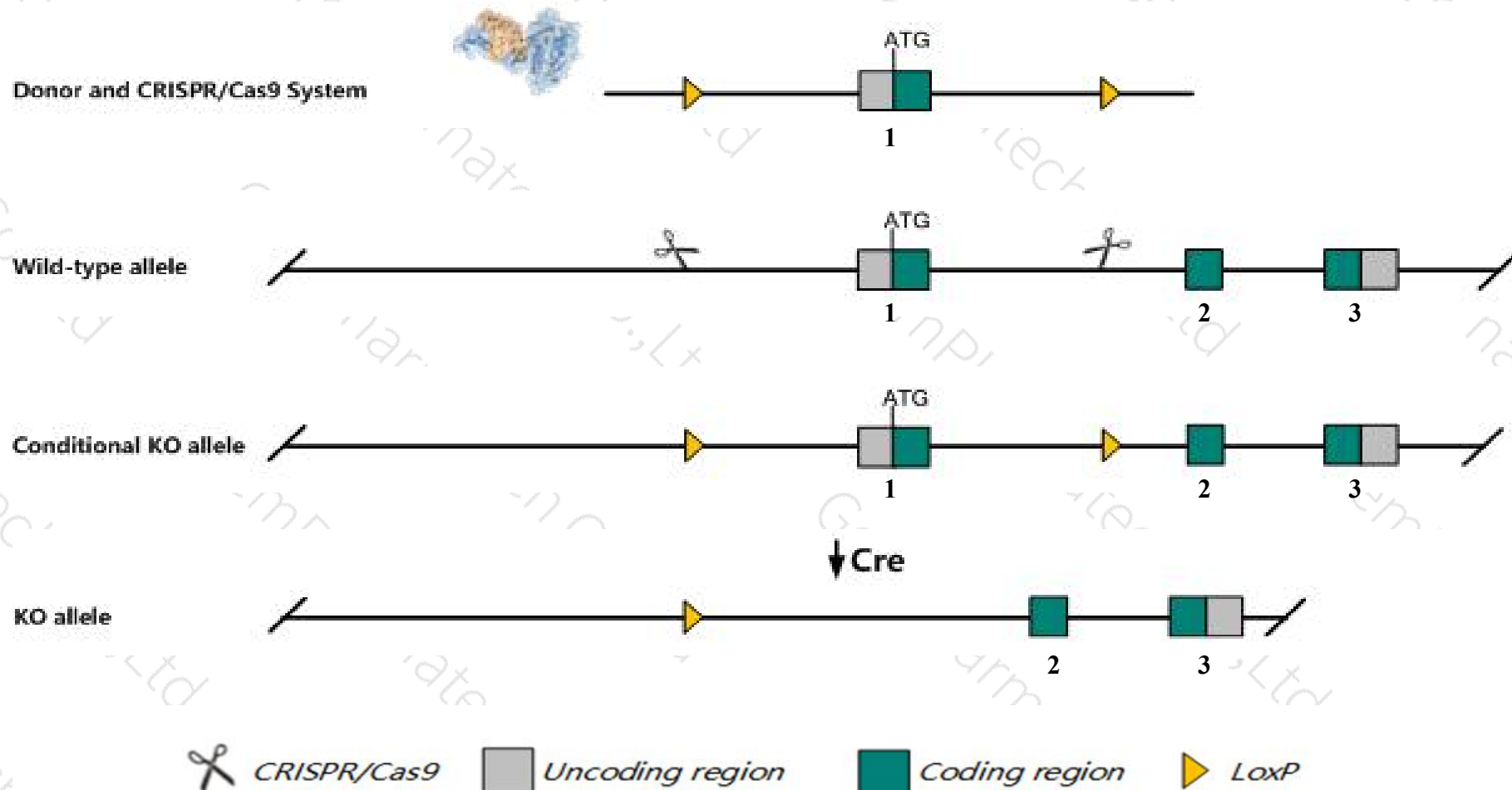
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Med26* gene. The schematic diagram is as follows:



Technical routes

- The *Med26* gene has 1 transcript. According to the structure of *Med26* gene, exon1 of *Med26-201* (ENSMUST00000058534.6) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Med26* 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.

- A new translation initiation site ATG, may be recognized after knocking out ATG, and there is a risk of forming an unknown protein.
- The *Med26* gene is located on the Chr8. 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)

Med26 mediator complex subunit 26 [*Mus musculus* (house mouse)]

Gene ID: 70625, updated on 12-Aug-2019

Summary

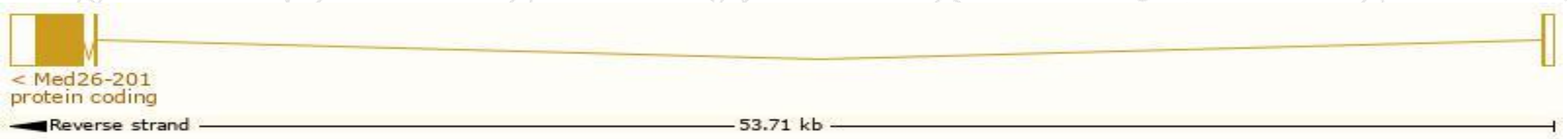
Official Symbol	Med26 provided by MGI
Official Full Name	mediator complex subunit 26 provided by MGI
Primary source	MGI:MGI:1917875
See related	Ensembl:ENSMUSG000000045248
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Crsp7; AI414941; AW495270; 5730493L18Rik
Expression	Ubiquitous expression in testis adult (RPKM 35.9), ovary adult (RPKM 12.2) and 28 other tissues 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
Med26-201	ENSMUST00000058534.6	2997	588aa	Protein coding	CCDS22416	Q7TN02	TSL:1 GENCODE basic APPRIS P1

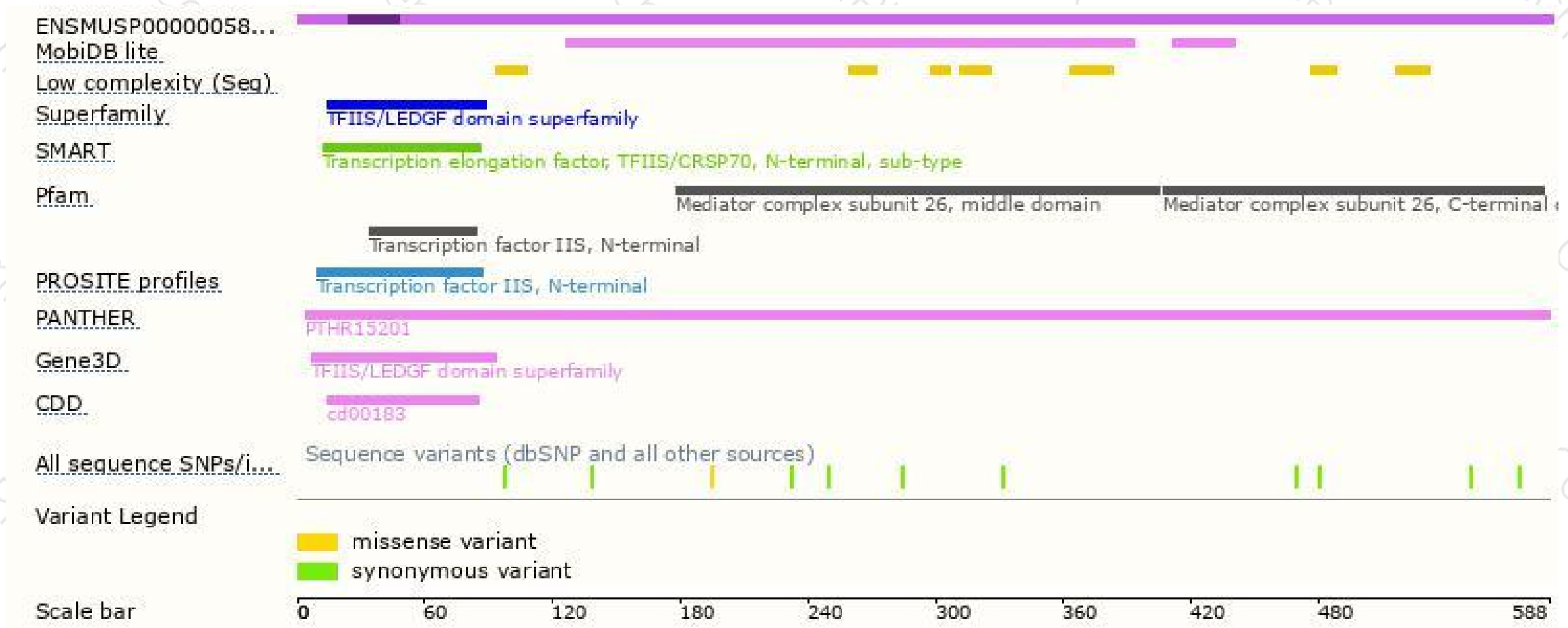
The strategy is based on the design of *Med26-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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