

***Med11* Cas9-KO Strategy**

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Project Overview

Project Name

Med11

Project type

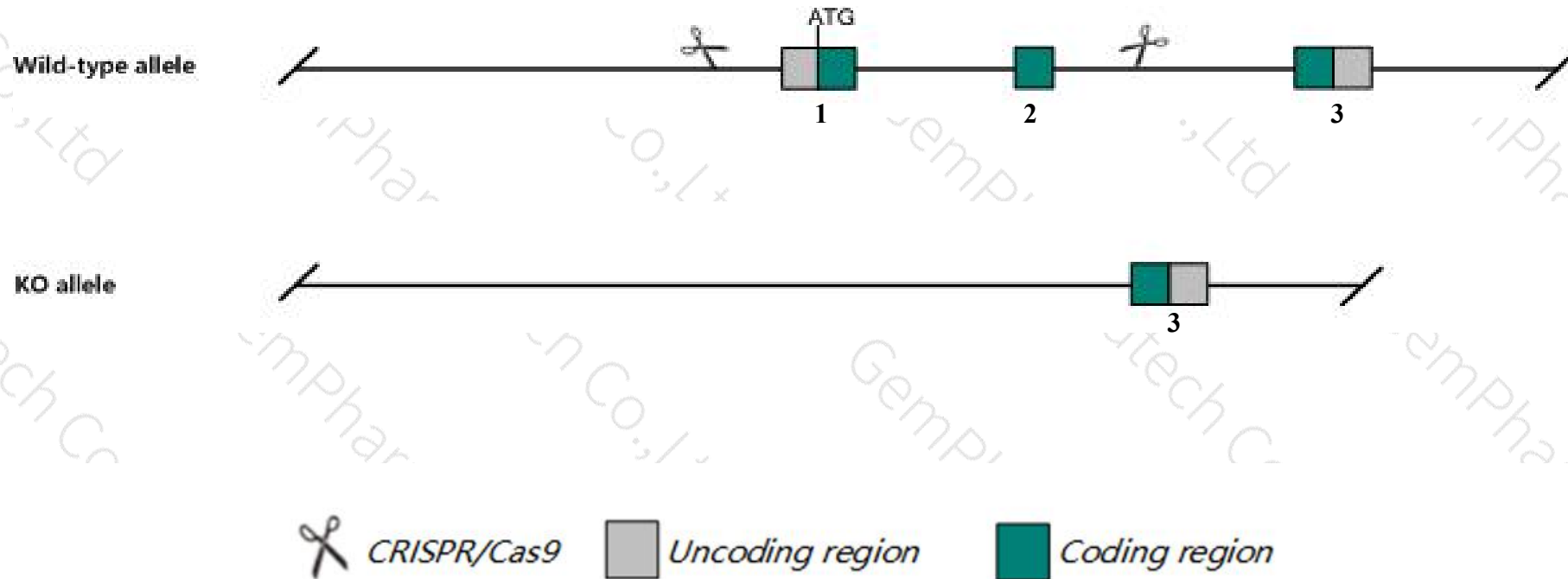
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Med11* gene. The schematic diagram is as follows:



- The *Med11* gene has 3 transcripts. According to the structure of *Med11* gene, exon1-exon2 of *Med11-201* (ENSMUST00000019067.7) 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 *Med11* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Med11* gene is located on the Chr11. 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)

Med11 mediator complex subunit 11 [Mus musculus (house mouse)]

Gene ID: 66172, updated on 31-Jan-2019

Summary



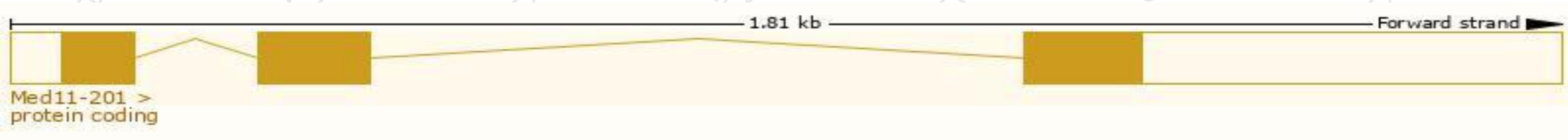
Official Symbol	Med11 provided by MGI
Official Full Name	mediator complex subunit 11 provided by MGI
Primary source	MGI:MGI:1913422
See related	Ensembl:ENSMUSG00000018923
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	1110030J09Rik, AI465144, AW545069
Expression	Ubiquitous expression in CNS E11.5 (RPKM 12.5), placenta adult (RPKM 11.9) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

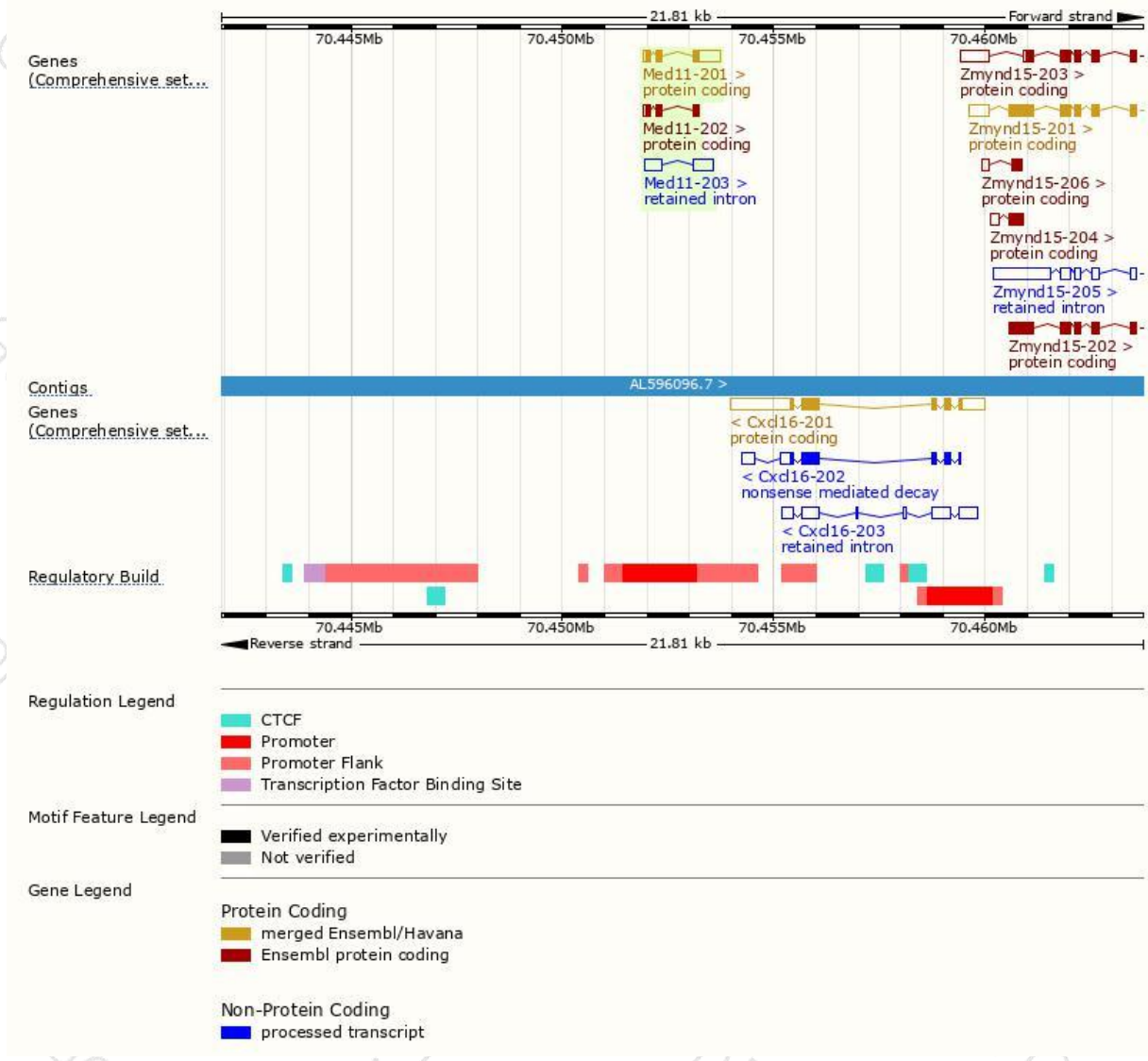
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Med11-201	ENSMUST00000019067.7	902	117aa	Protein coding	CCDS24947	Q9D8C6	TSL:1 GENCODE basic APPRIS P1
Med11-202	ENSMUST00000151013.7	445	127aa	Protein coding	-	G3UZ31	TSL:2 GENCODE basic
Med11-203	ENSMUST00000152348.1	870	No protein	Retained intron	-	-	TSL:2

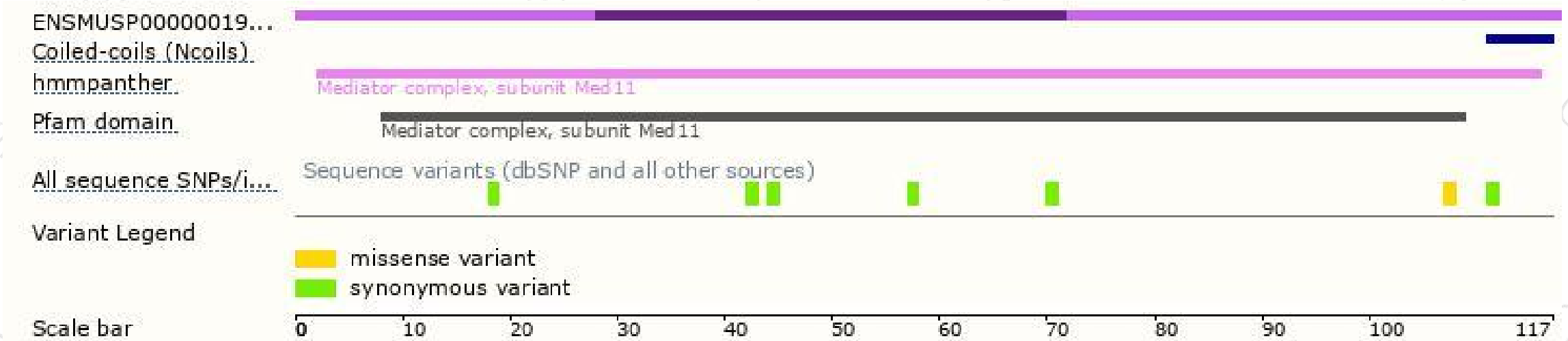
The strategy is based on the design of *Med11-201* transcript,The transcription is shown below



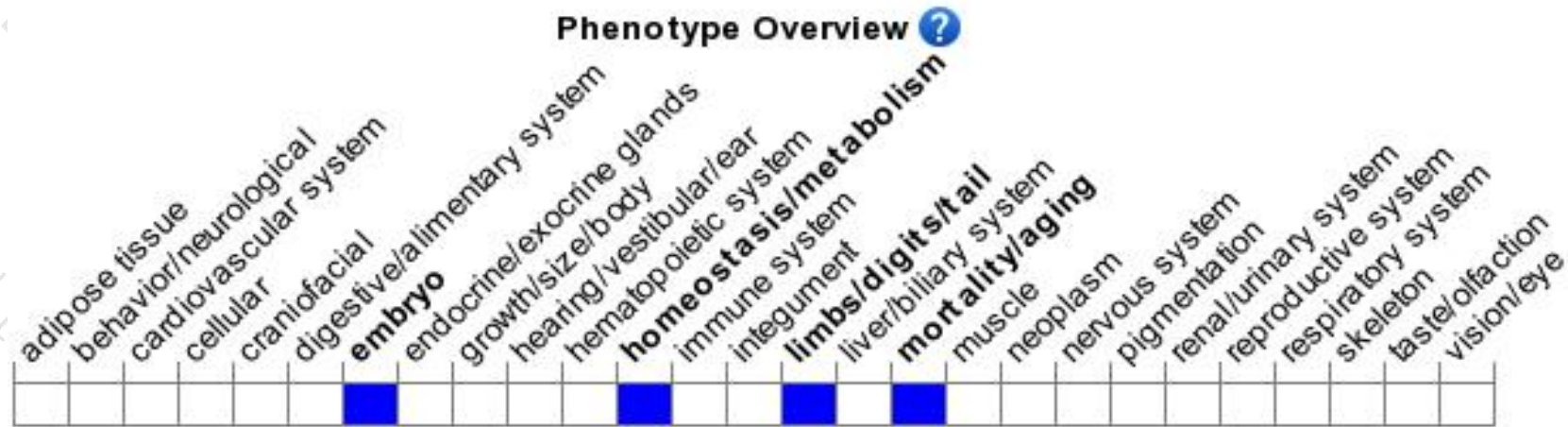
Genomic location distribution



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/>).

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

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