

***Cand2* Cas9-KO Strategy**

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Project Overview

Project Name

Cand2

Project type

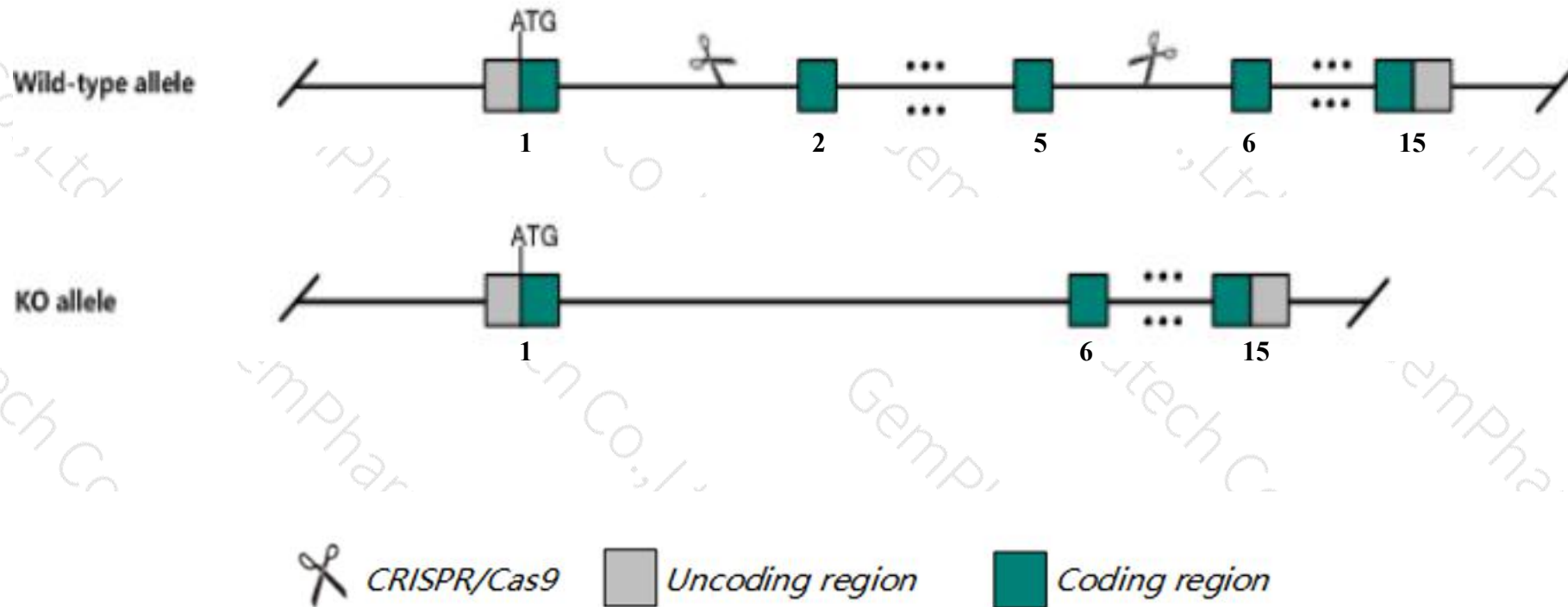
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Cand2* gene. The schematic diagram is as follows:



- The *Cand2* gene has 1 transcript. According to the structure of *Cand2* gene, exon2-exon5 of *Cand2-201* (ENSMUST00000075995.6) transcript is recommended as the knockout region. The region contains 689bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Cand2* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Cand2* gene is located on the Chr6. 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)

Cand2 cullin-associated and neddylation-dissociated 2 (putative) [Mus musculus (house mouse)]

Gene ID: 67088, updated on 13-Mar-2020

Summary



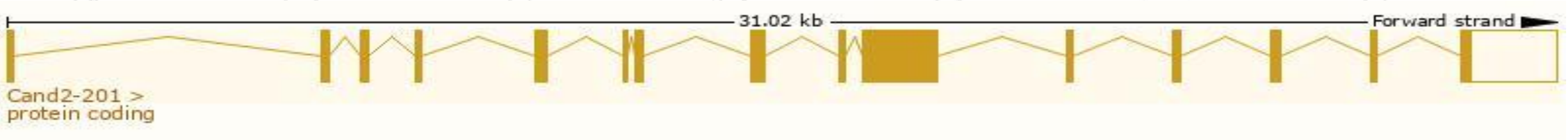
Official Symbol	Cand2 provided by MGI
Official Full Name	cullin-associated and neddylation-dissociated 2 (putative) provided by MGI
Primary source	MGI:MGI:1914338
See related	Ensembl:ENSMUSG00000030319
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	2210404G23Rik, Tp120b, mKIAA0667
Expression	Broad expression in limb E14.5 (RPKM 24.2), heart adult (RPKM 22.0) and 26 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

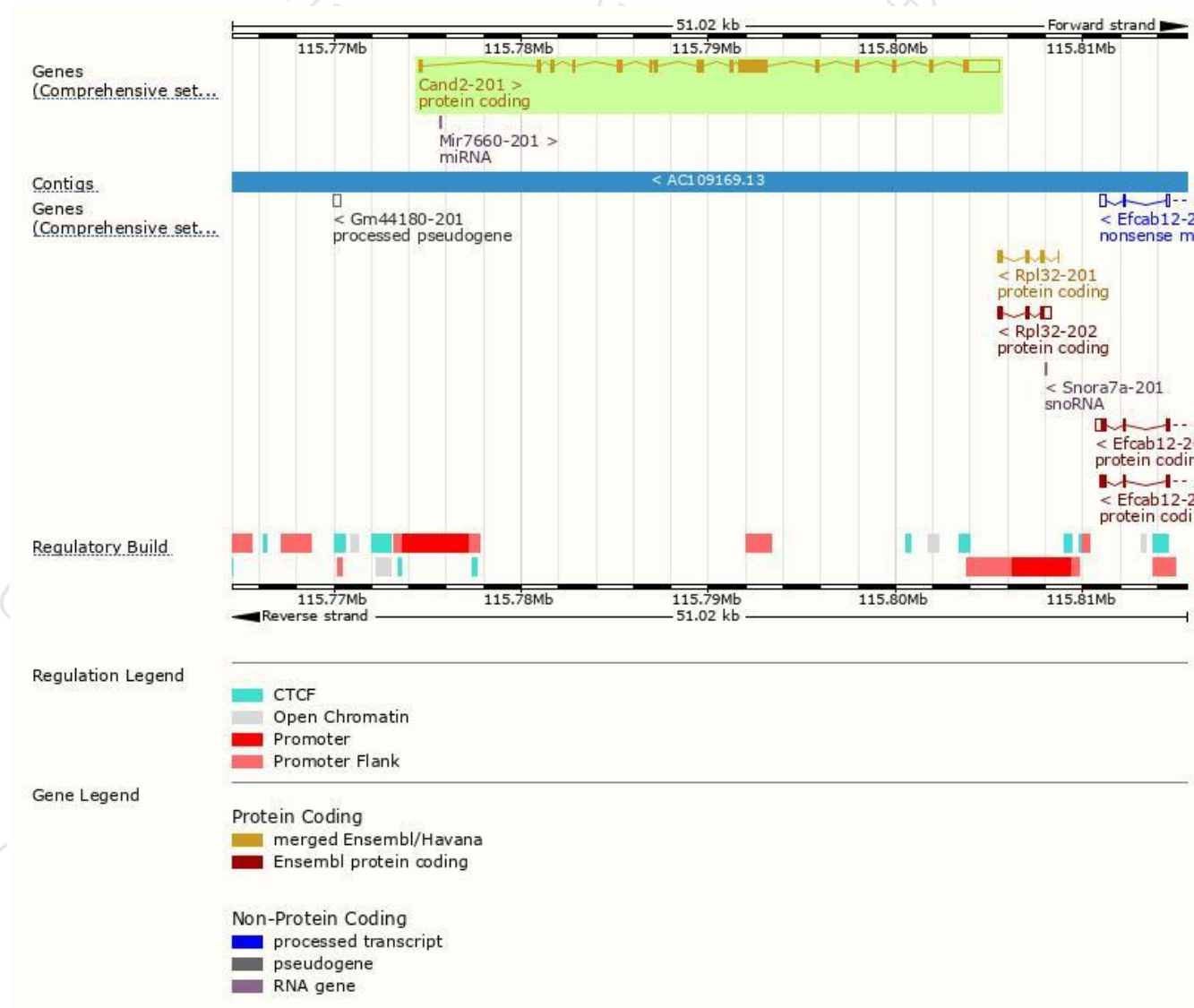
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Cand2-201	ENSMUST00000075995.6	5474	1235aa	Protein coding	CCDS20442	Q6ZQ73	TSL:1 GENCODE basic APPRIS P1

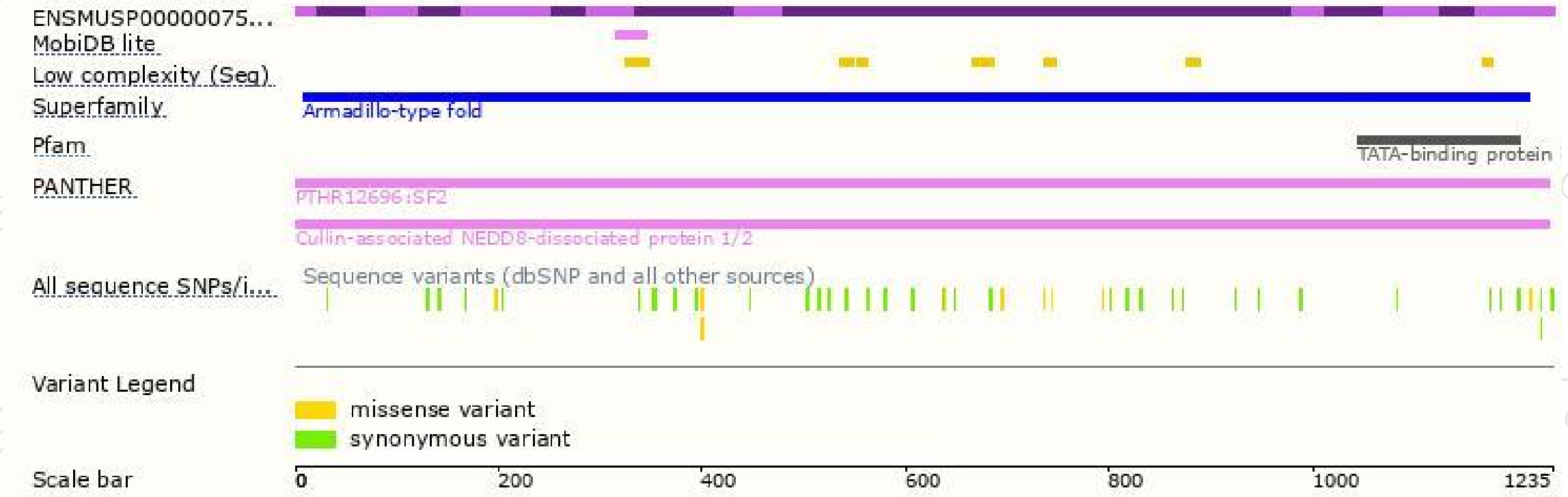
The strategy is based on the design of *Cand2-201* transcript, the transcription is shown below:



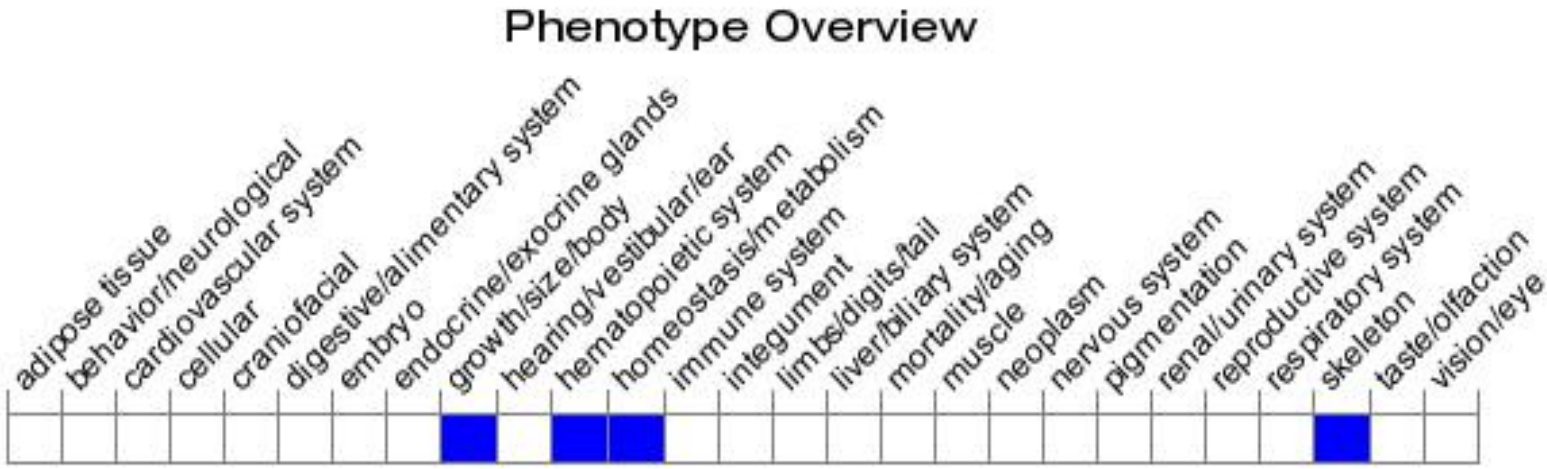
Genomic location distribution



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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