

# *Daglb* Cas9-KO Strategy

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**Reviewer:**

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# Project Overview

**Project Name**

*Daglb*

**Project type**

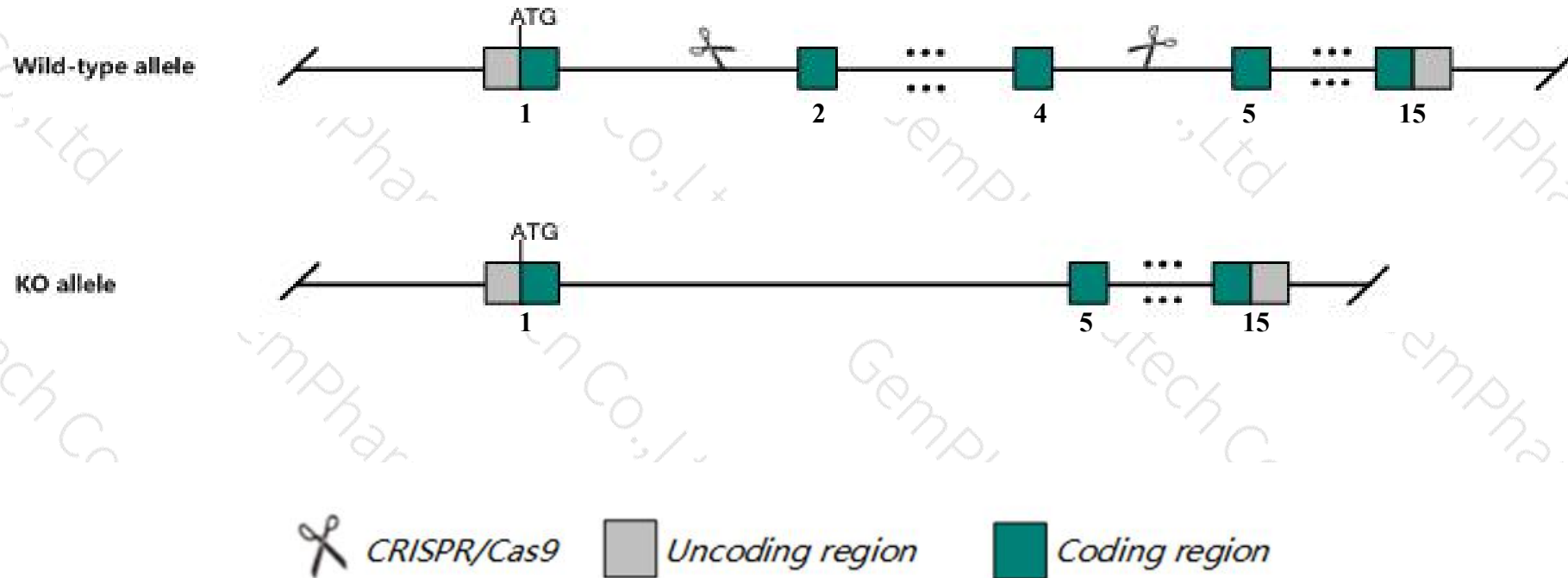
**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

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



- The *Daglb* gene has 4 transcripts. According to the structure of *Daglb* gene, exon2-exon4 of *Daglb-201* (ENSMUST00000045593.11) transcript is recommended as the knockout region. The region contains 583bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Daglb* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice homozygous for null mutations have a reduction in endocannabinoids in the brain and a decrease in adult neuronal proliferation in the hippocampus.
- The *Daglb* gene is located on the Chr5. 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)

## Daglb diacylglycerol lipase, beta [Mus musculus (house mouse)]

Gene ID: 231871, updated on 31-Jan-2019

### Summary



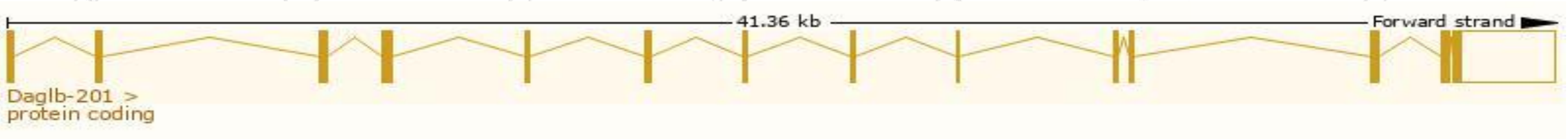
<b>Official Symbol</b>	Daglb provided by <a href="#">MGI</a>
<b>Official Full Name</b>	diacylglycerol lipase, beta provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:2442032</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000039206</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	VALIDATED
<b>Organism</b>	<a href="#">Mus musculus</a>
<b>Lineage</b>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
<b>Also known as</b>	E330036I19Rik
<b>Expression</b>	Ubiquitous expression in adrenal adult (RPKM 20.1), mammary gland adult (RPKM 16.6) and 27 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

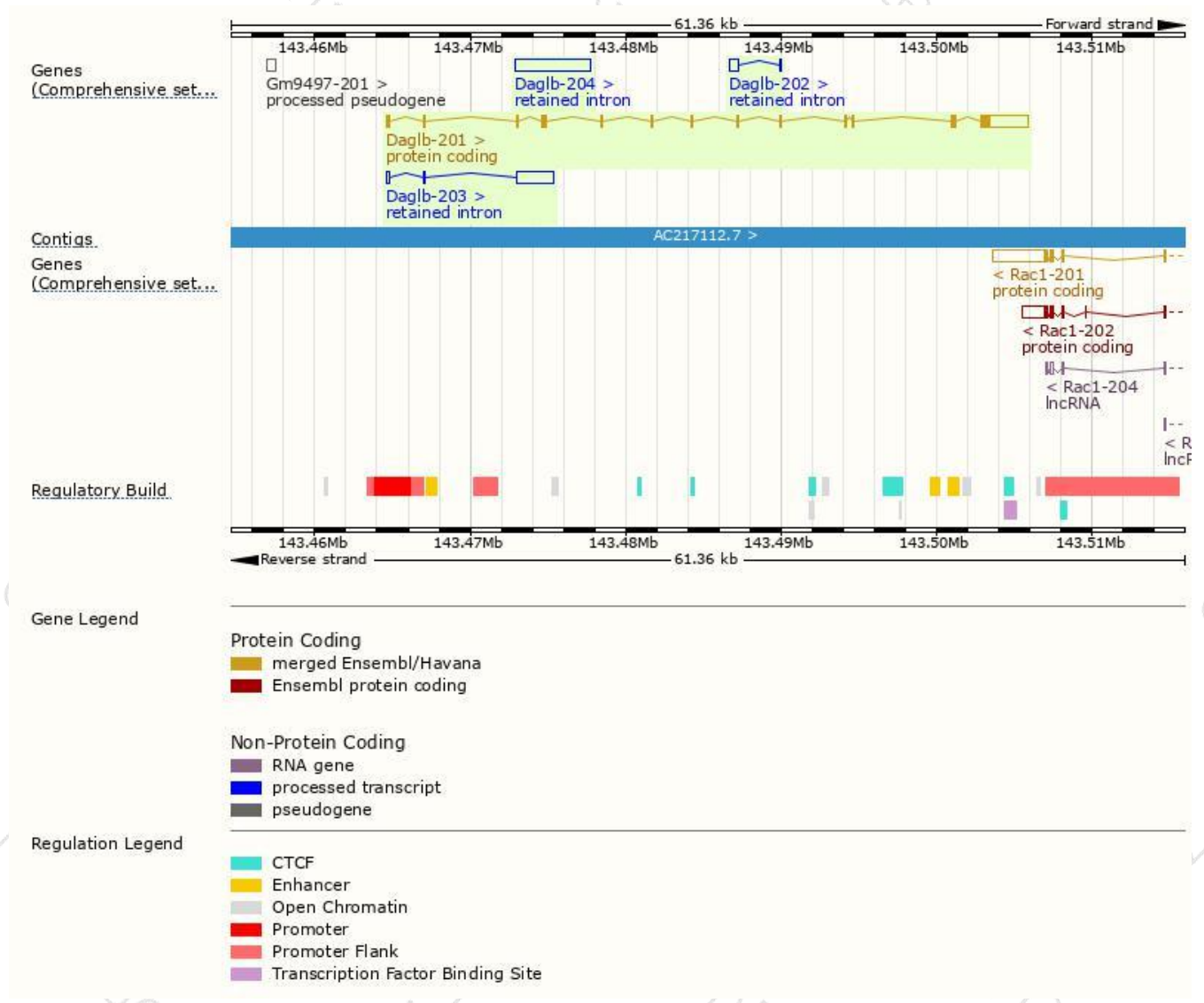
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Daglb-201	<a href="#">ENSMUST00000045593.11</a>	4651	<a href="#">669aa</a>	Protein coding	<a href="#">CCDS39370</a>	<a href="#">Q91WC9</a>	TSL:1 GENCODE basic APPRIS P1
Daglb-204	<a href="#">ENSMUST00000199577.1</a>	4877	No protein	Retained intron	-	-	TSL:NA
Daglb-203	<a href="#">ENSMUST00000174273.2</a>	2712	No protein	Retained intron	-	-	TSL:1
Daglb-202	<a href="#">ENSMUST00000155383.1</a>	622	No protein	Retained intron	-	-	TSL:2

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

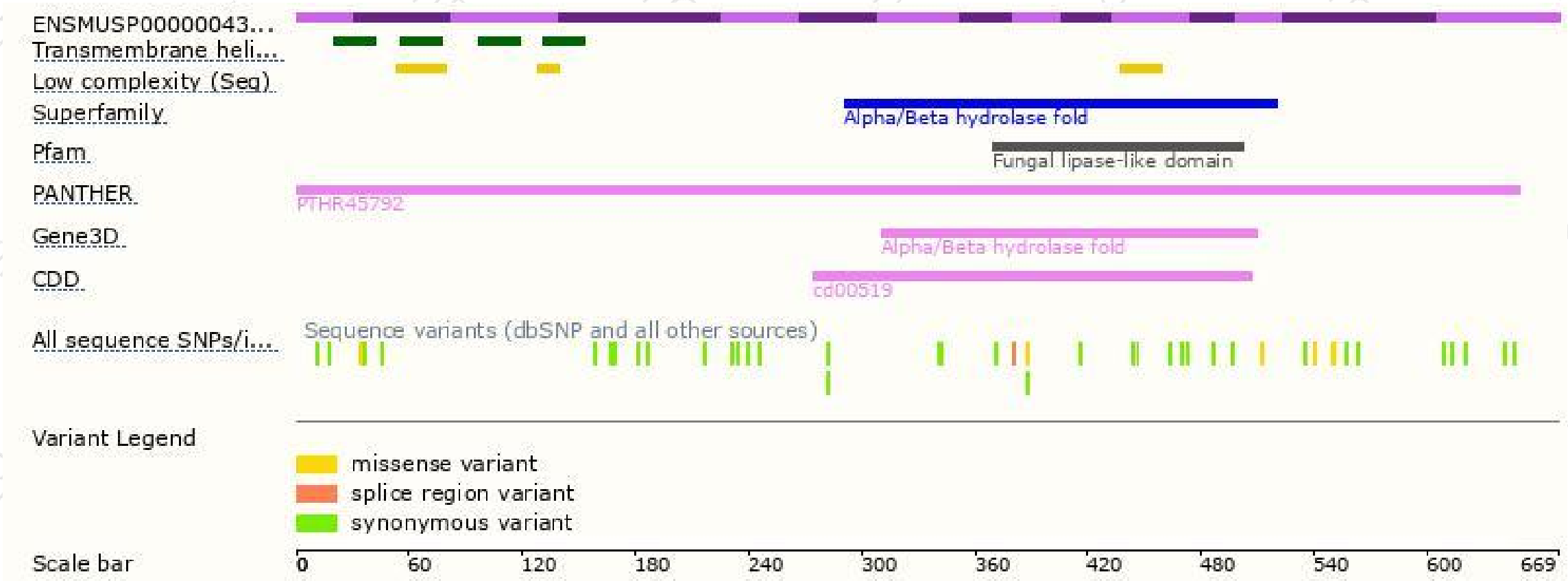


# Genomic location distribution

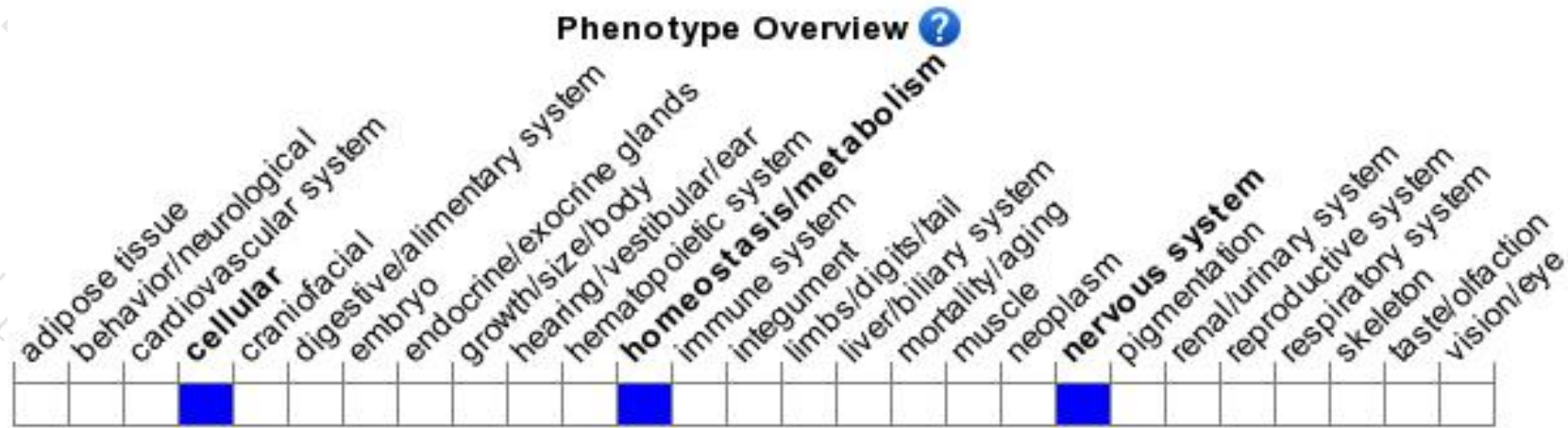




# Protein domain



# Mouse phenotype description(MGI )



*Phenotypes affected by the gene are marked in blue. Data quoted from MGI database(<http://www.informatics.jax.org/>).*

According to the existing MGI data, Mice homozygous for null mutations have a reduction in endocannabinoids in the brain and a decrease in adult neuronal proliferation in the hippocampus.

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

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