

# ***Dync1li1 Cas9-KO Strategy***

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**Reviewer :**

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# Project Overview

**Project Name**

***Dync1li1***

**Project type**

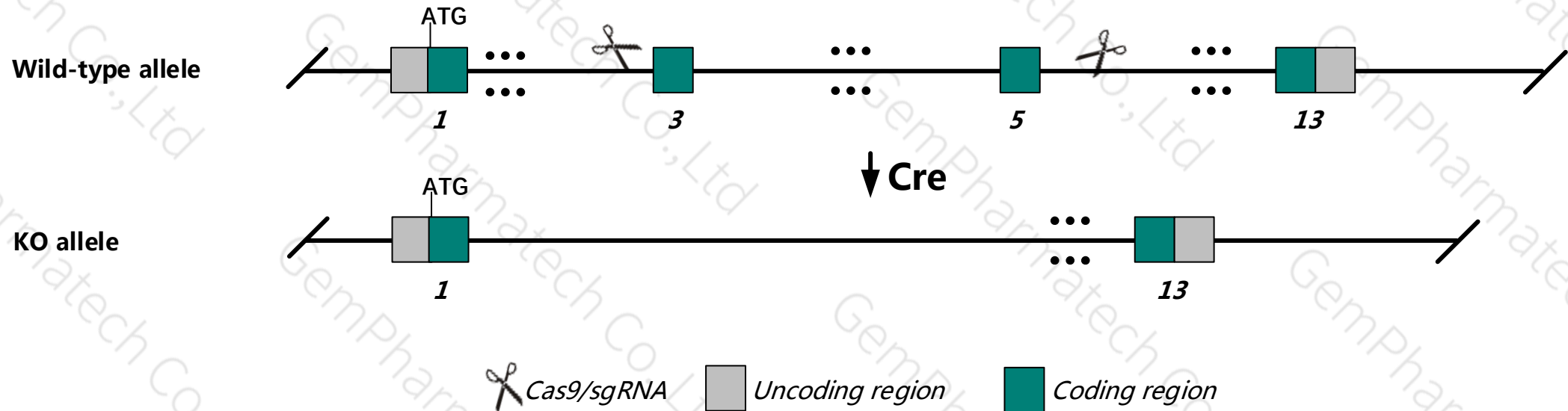
**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Dync1li1* gene. The schematic diagram is as follows:



# Technical routes

- The *Dync1li1* gene has 2 transcripts. According to the structure of *Dync1li1* gene, exon3-exon5 of *Dync1li1*-201 ( ENSMUST00000047404.6) transcript is recommended as the knockout region. The region contains 518bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Dync1li1* gene. The brief process is as follows: sgRNA was transcribed in vitro. Cas9, sgRNA 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.

- According to the existing MGI data , Mice homozygous for an ENU-induced allele exhibit increased anxiety-related response, increased dendrite length, increased neuron migration, and decreased lysosome trafficking.
- The *Dync1li1* gene is located on the Chr9. 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 )

## Dync1li1 dynein cytoplasmic 1 light intermediate chain 1 [ *Mus musculus* (house mouse) ]

Gene ID: 235661, updated on 13-Mar-2020

### Summary

Official Symbol	Dync1li1 provided by <a href="#">MGI</a>
Official Full Name	dynein cytoplasmic 1 light intermediate chain 1 provided by <a href="#">MGI</a>
Primary source	<a href="#">MGI:MGI:2135610</a>
See related	<a href="#">Ensembl:ENSMUSG00000032435</a>
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	<a href="#">Mus musculus</a>
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	LIC-1; Dncl1; 1110053F02Rik
Expression	Ubiquitous expression in CNS E18 (RPKM 27.9), CNS E14 (RPKM 23.6) and 28 other tissues <a href="#">See more</a>
Orthologs	<a href="#">human</a> <a href="#">all</a>

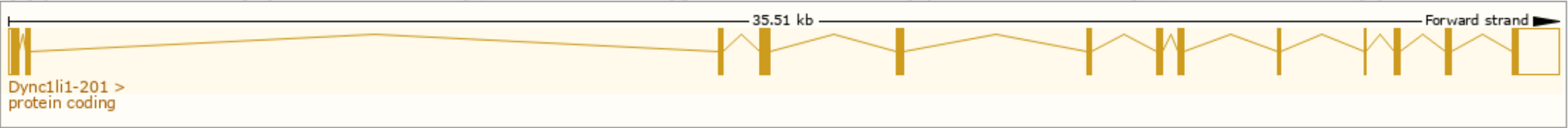


# Transcript information ( Ensembl )

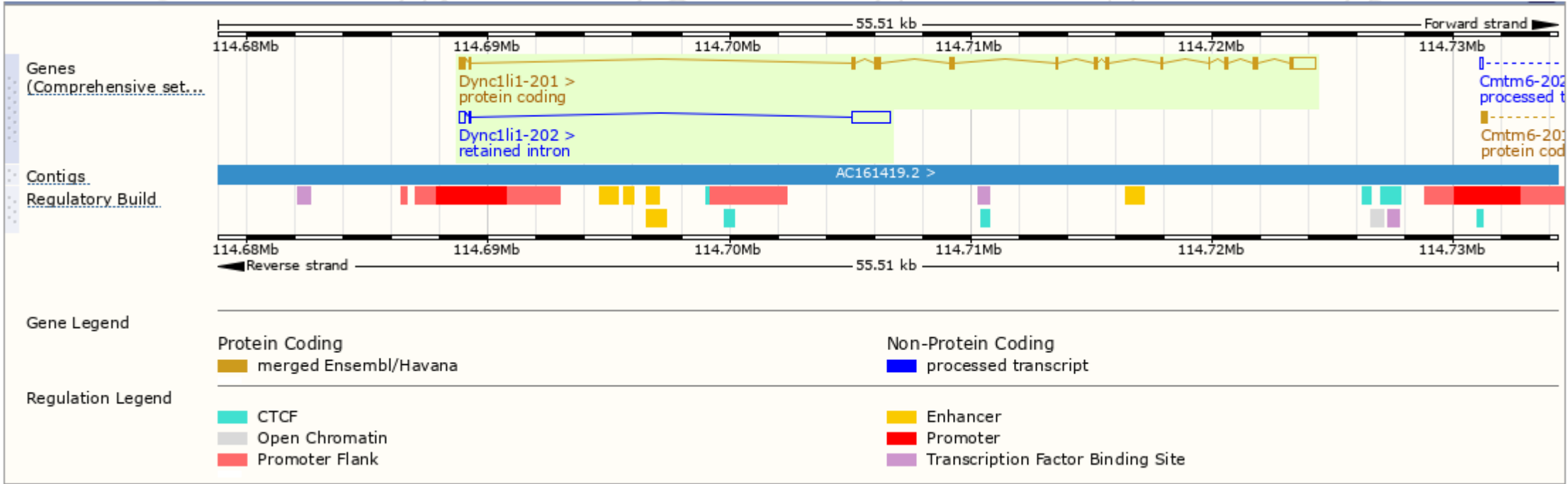
The gene has 2 transcripts, and all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Dync1li1-201	<a href="#">ENSMUST00000047404.6</a>	2601	<a href="#">523aa</a>	Protein coding	<a href="#">CCDS23595</a>	<a href="#">Q3TWG5</a> <a href="#">Q8R1Q8</a>	TSL:1 Gencode basic APPRIS P1
Dync1li1-202	<a href="#">ENSMUST00000215345.1</a>	1930	No protein	Retained intron	-	-	TSL:1

The strategy is based on the design of *Dync1li1*-201 transcript, The transcription is shown below

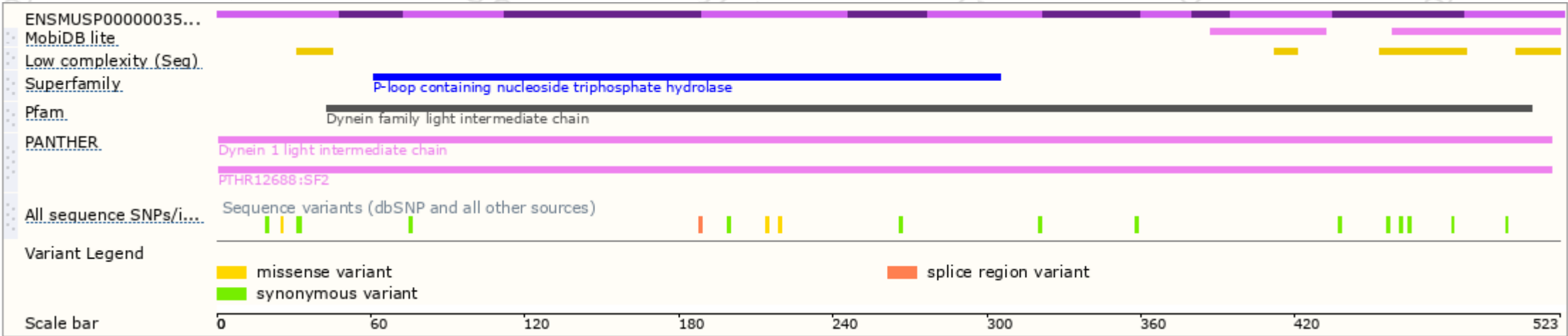


# Genomic location distribution

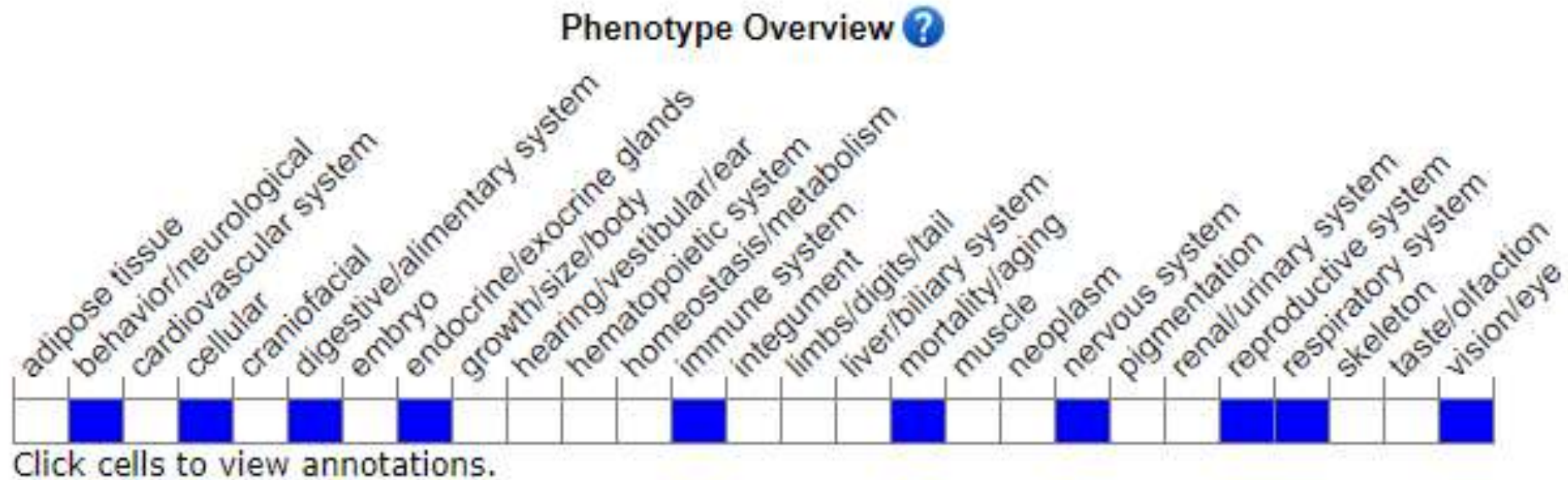




# Protein domain



# 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 an ENU-induced allele exhibit increased anxiety-related response, increased dendrite length, increased neuron migration, and decreased lysosome trafficking.

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