

Neurog2 Cas9-KO Strategy

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Project Overview

Project Name

Neurog2

Project type

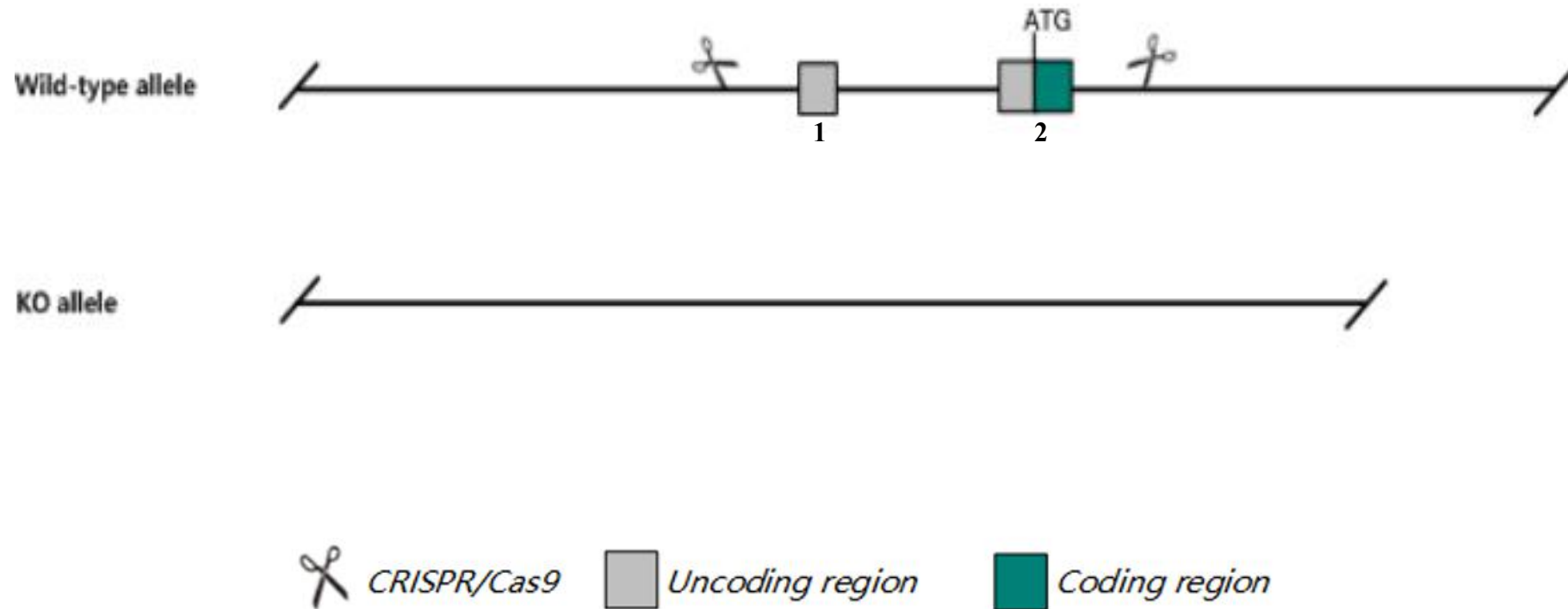
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Neurog2* gene. The schematic diagram is as follows:



The *Neurog2* gene has 1 transcript. According to the structure of *Neurog2* gene, exon1-exon2 of *Neurog2-201*(ENSMUST00000029587.9) transcript is recommended as the knockout region. The region contains all of the coding sequence. Knock out the region will result in disruption of protein function.

In this project we use CRISPR/Cas9 technology to modify *Neurog2* gene. The brief process is as follows: CRISPR/Cas9 system 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, homozygous null mice die after birth and have neuronal differentiation defects, affecting retinal development, spinal cord interneuron development and behavior.

The *Neurog2* gene is located on the Chr3. 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.

Neurog2 neurogenin 2 [Mus musculus (house mouse)]

Gene ID: 11924, updated on 23-Feb-2021

Summary



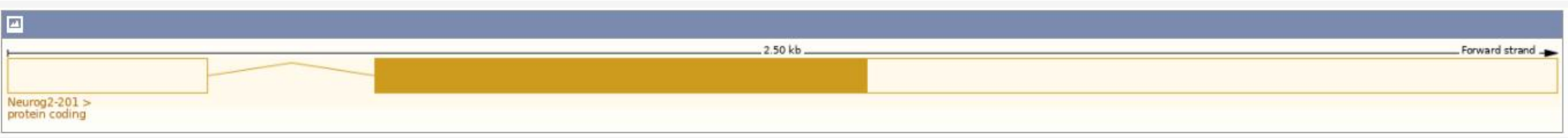
Official Symbol	Neurog2 provided by MGI
Official Full Name	neurogenin 2 provided by MGI
Primary source	MGI:MGI:109619
See related	Ensembl:ENSMUSG00000027967
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Ato, Atoh4, Math, Math4A, bHLH, bHLHa8, ng, ngn-2, ngn2
Expression	Biased expression in CNS E14 (RPKM 28.5), whole brain E14.5 (RPKM 22.8) and 2 other tissues See more
Orthologs	human all

Transcript information Ensembl

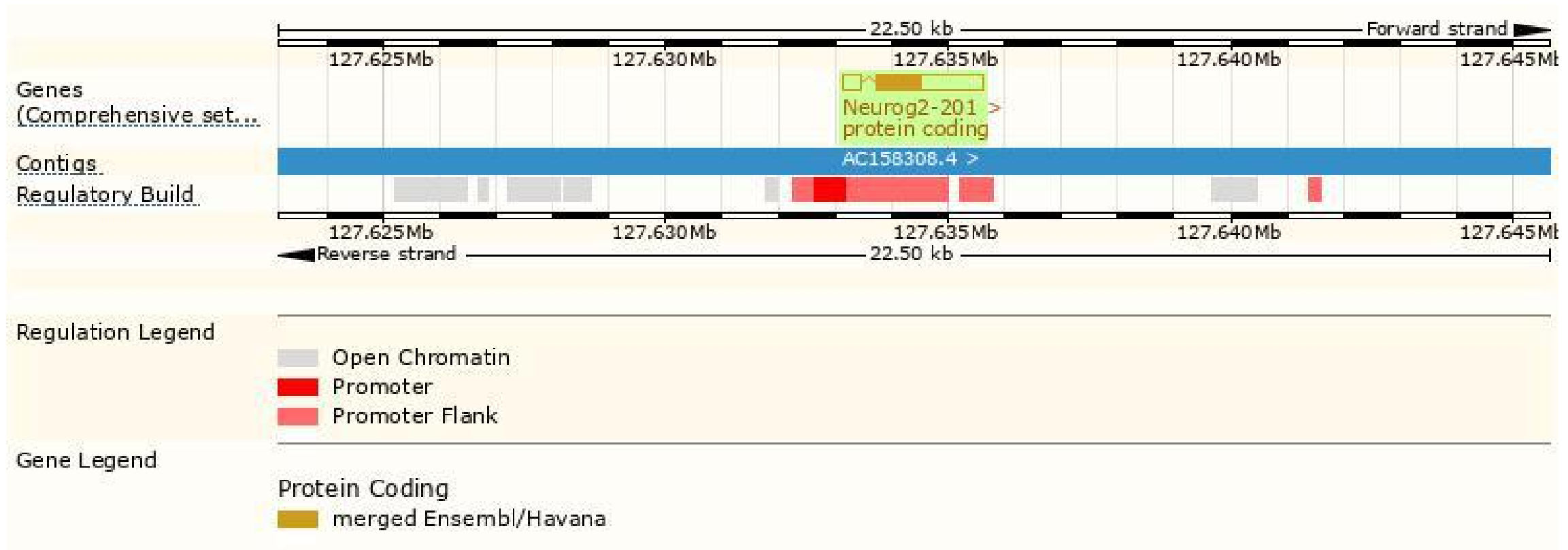
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Neurog2-201	ENSMUST00000029587.9	2227	263aa	Protein coding	CCDS17825		TSL:1 , GENCODE basic , APPRIS P1 ,

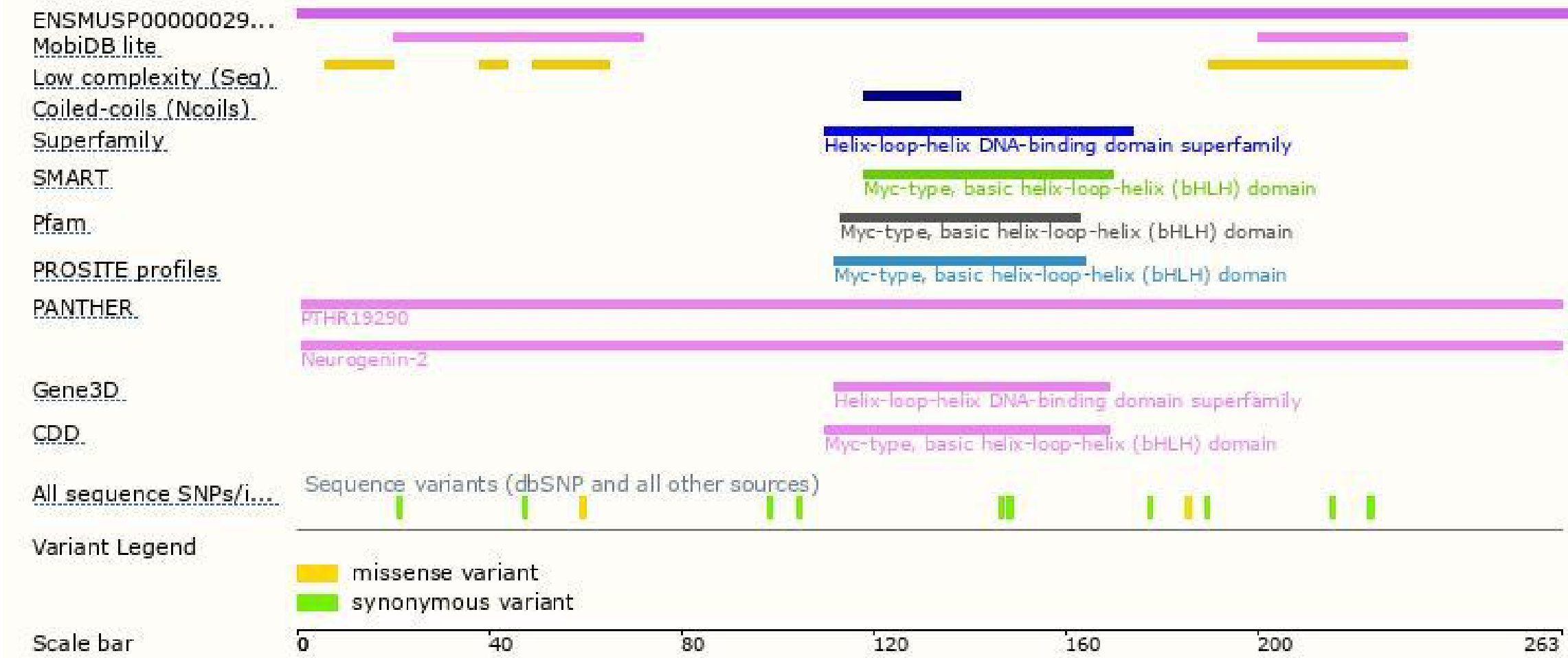
The strategy is based on the design of *Neurog2-201* transcript, the transcription is shown below:



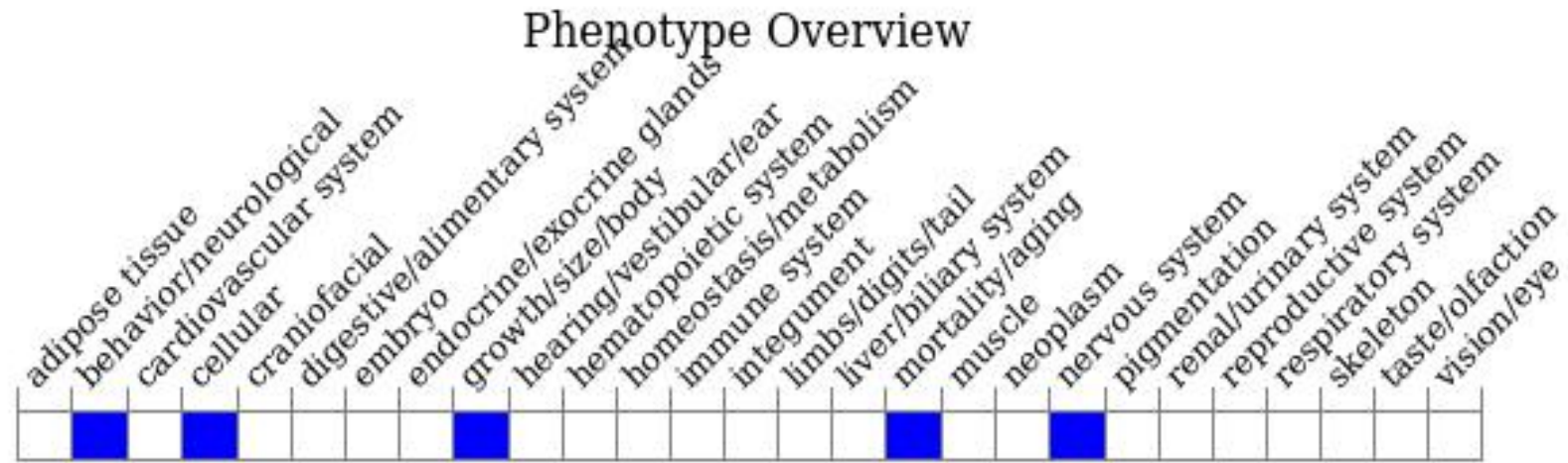
Genomic location distribution



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, homozygous null mice die after birth and have neuronal differentiation defects, affecting retinal development, spinal cord interneuron development and behavior.

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