# *Neurog1* Cas9-KO Strategy

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# Project Overview

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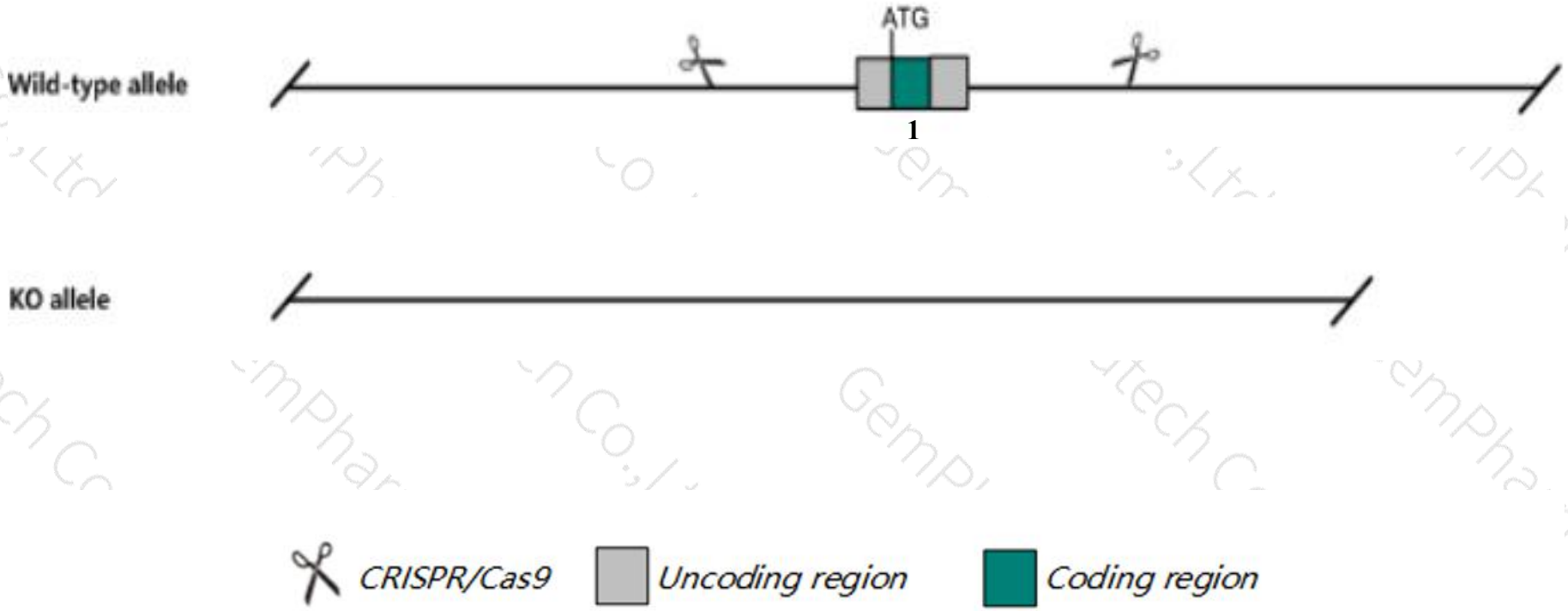
| **Project Name** | ***Neurog1*** |
| **Project type** | **Cas9-KO** |
| **Strain background** | **C57BL/6JGpt** |

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Neurog1* gene. The schematic diagram is as follows:



# Technical routes

➢ The *Neurog1* gene has 1 transcript. According to the structure of *Neurog1* gene, exon1 of *Neurog1-201*(ENSMUST00000058475.5) 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 *Neurog1* 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.

# Notice

➢ According to the existing MGI data,homozygotes for a targeted null mutation exhibit defects in midbrain, dorsal root sensory ganglia and a subset of cranial ganglia. Mutants are born alive, but fail to nurse, and die within 12 hours.

➢ The *Neurog1* gene is located on the Chr13. 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）

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## Neurog1 neurogenin 1 [Mus musculus (house mouse)]

Gene ID: 18014, updated on 13-Mar-2020

### Summary

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|---|---|
| **Official Symbol** | Neurog1 provided by MGI |
| **Official Full Name** | neurogenin 1 provided by MGI |
| **Primary source** | MGI:MGI:107754 |
| **See related** | Ensembl:ENSMUSG00000048904 |
| **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** | AKA, Math4C, Neurod3, bHLHa6, ngn1 |
| **Orthologs** | human all |

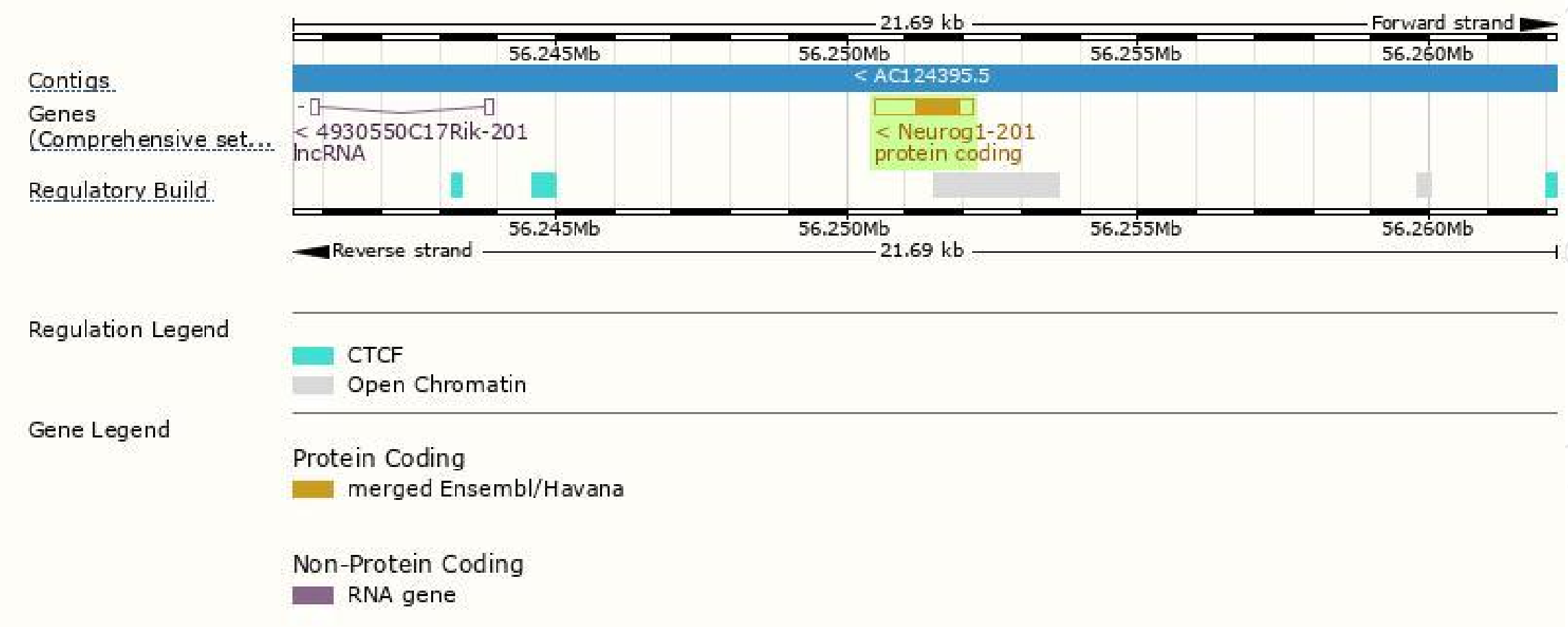
# Transcript information（Ensembl）

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

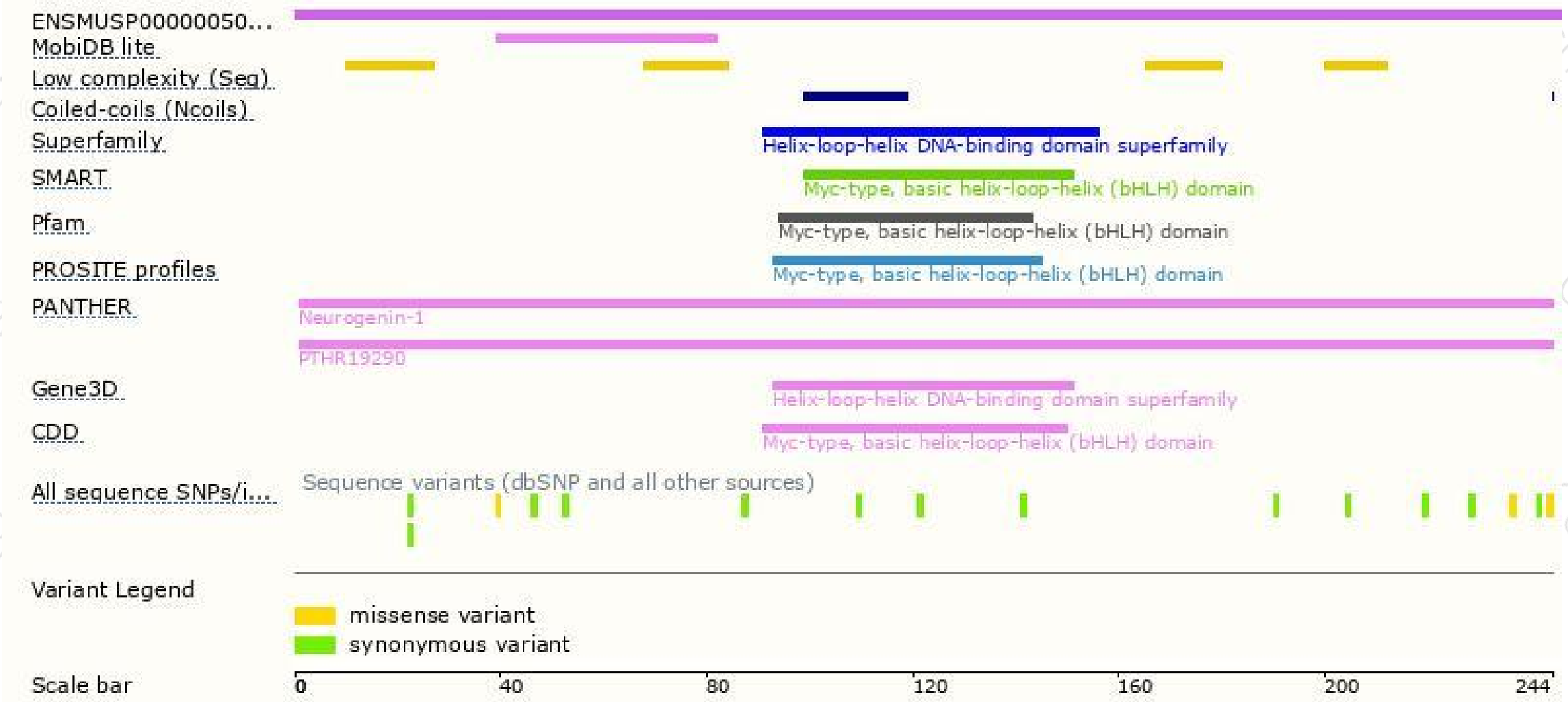
| Name | Transcript ID | bp | Protein | Biotype | CCDS | UniProt | Flags |
|---|---|---|---|---|---|---|---|
| Neurog1-201 | ENSMUST00000058475.5 | 1686 | 244aa | Protein coding | CCDS26559 | P70660 | TSL:NA GENCODE basic APPRIS is a system to annotate alternatively spliced transcripts based on a range of computational methods to identify the most functionally important transcript(s) of a gene. APPRIS P1 |

The strategy is based on the design of *Neurog1-201* transcript,the transcription is shown below:

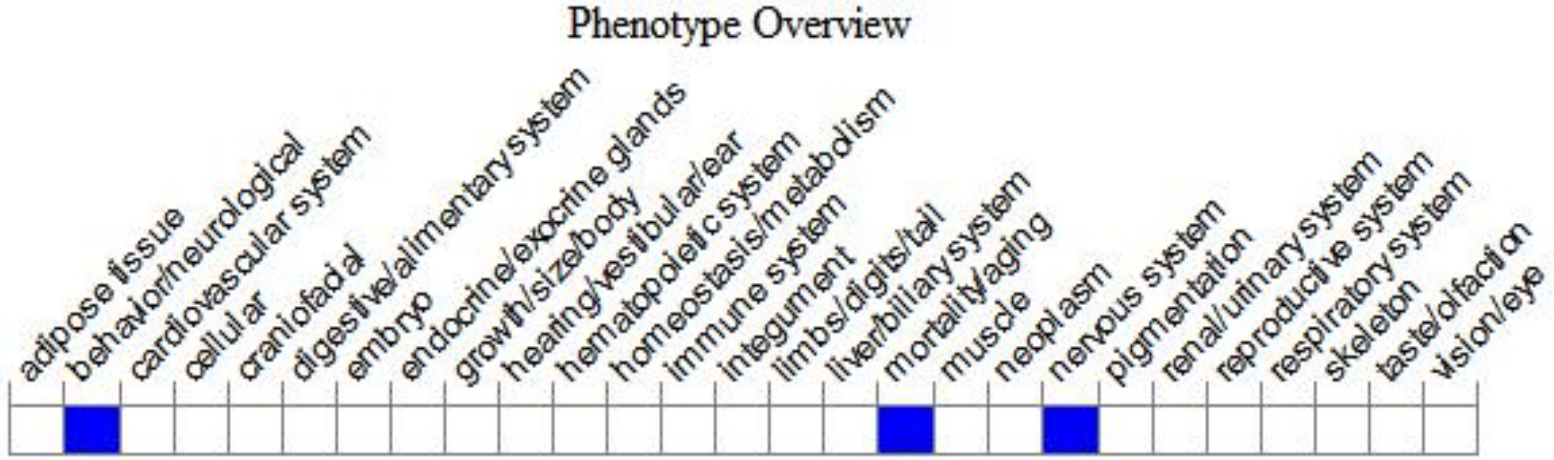
# Genomic location distribution



# 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,homozygotes for a targeted null mutation exhibit defects in midbrain, dorsal root sensory ganglia and a subset of cranial ganglia. Mutants are born alive, but fail to nurse, and die within 12 hours.

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

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