

Nom1 Cas9-KO Strategy

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Project Overview

Project Name

Nom1

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Nom1* gene. The schematic diagram is as follows:



- The *Nom1* gene has 2 transcripts. According to the structure of *Nom1* gene, exon3 of *Nom1-201* (ENSMUST00000001611.10) transcript is recommended as the knockout region. The region contains 196bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Nom1* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Nom1* 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)

Nom1 nucleolar protein with MIF4G domain 1 [Mus musculus (house mouse)]

Gene ID: 433864, updated on 31-Jan-2019

Summary



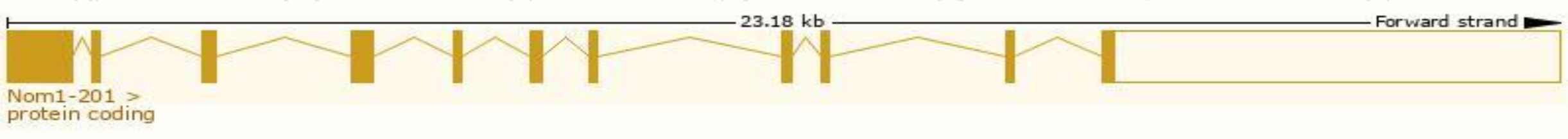
Official Symbol	Nom1 provided by MGI
Official Full Name	nucleolar protein with MIF4G domain 1 provided by MGI
Primary source	MGI:MGI:1861749
See related	Ensembl:ENSMUSG000000001569
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	D5Kng1, Gm1040
Expression	Ubiquitous expression in CNS E11.5 (RPKM 7.8), liver E14 (RPKM 6.0) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

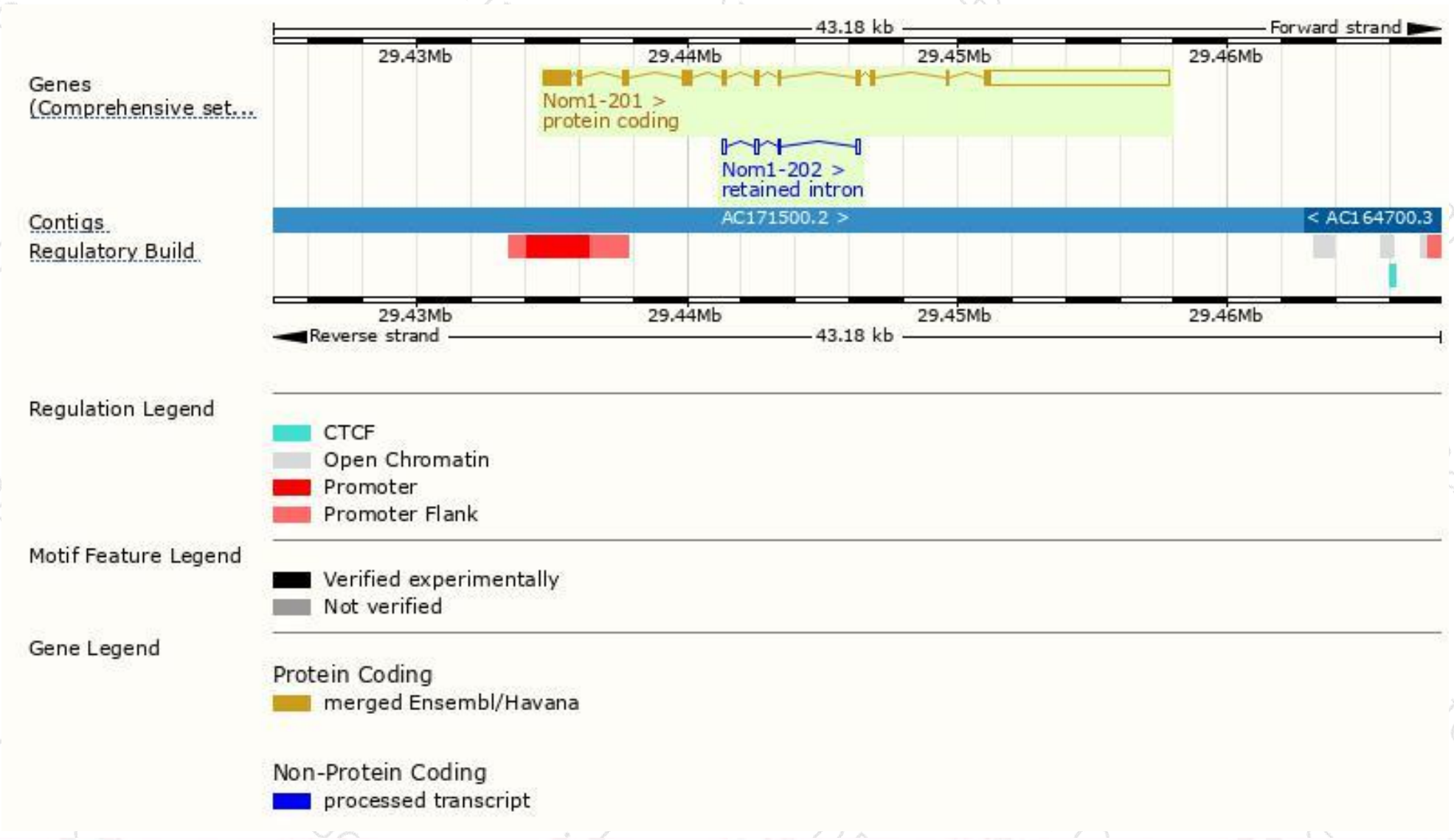
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Nom1-201	ENSMUST00000001611.10	9219	854aa	Protein coding	CCDS19149	Q3UFM5	TSL:1 GENCODE basic APPRIS P1
Nom1-202	ENSMUST00000146347.1	531	No protein	Retained intron	-	-	TSL:2

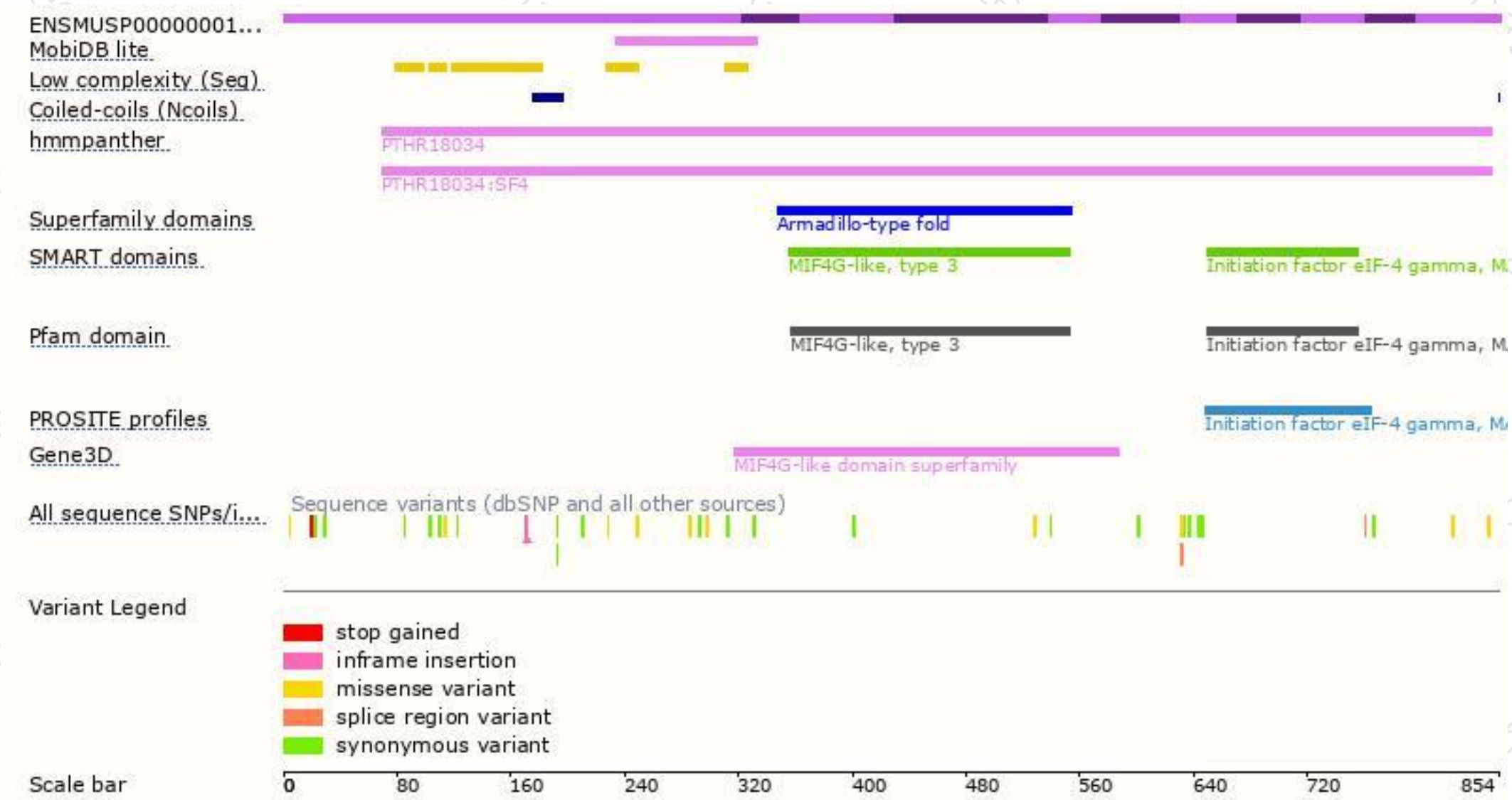
The strategy is based on the design of *Nom1-201* transcript,The transcription is shown below



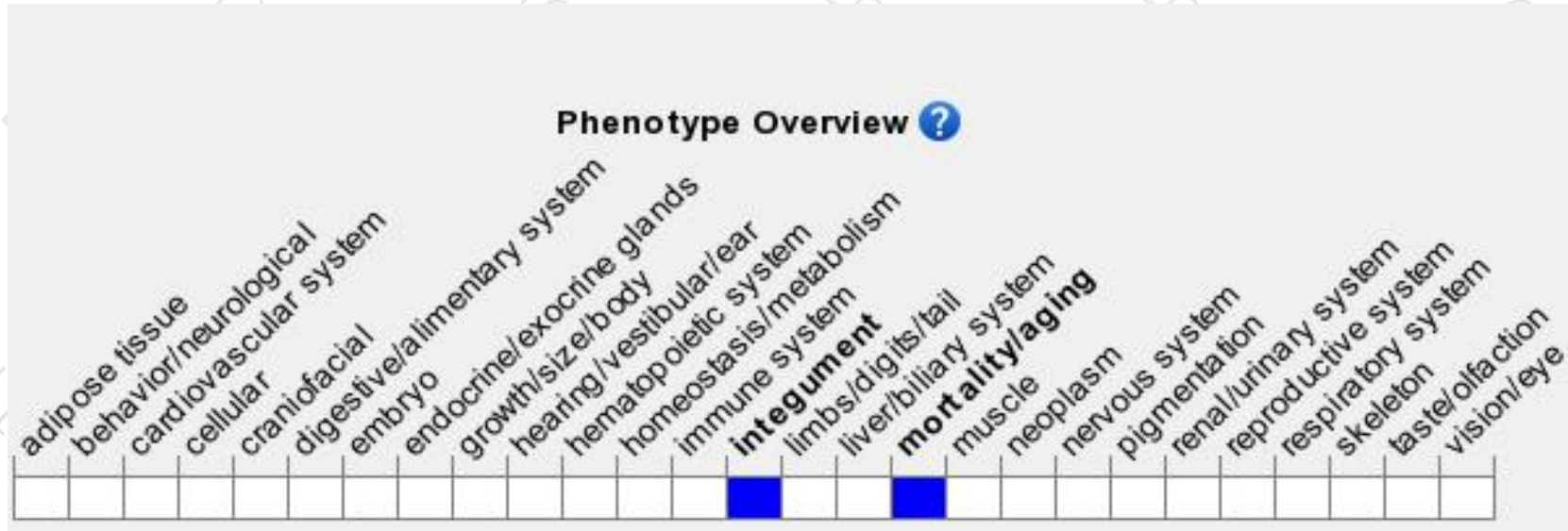
Genomic location distribution



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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