

Neu3 Cas9-KO Strategy

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Reviewer:

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Project Overview

Project Name

Neu3

Project type

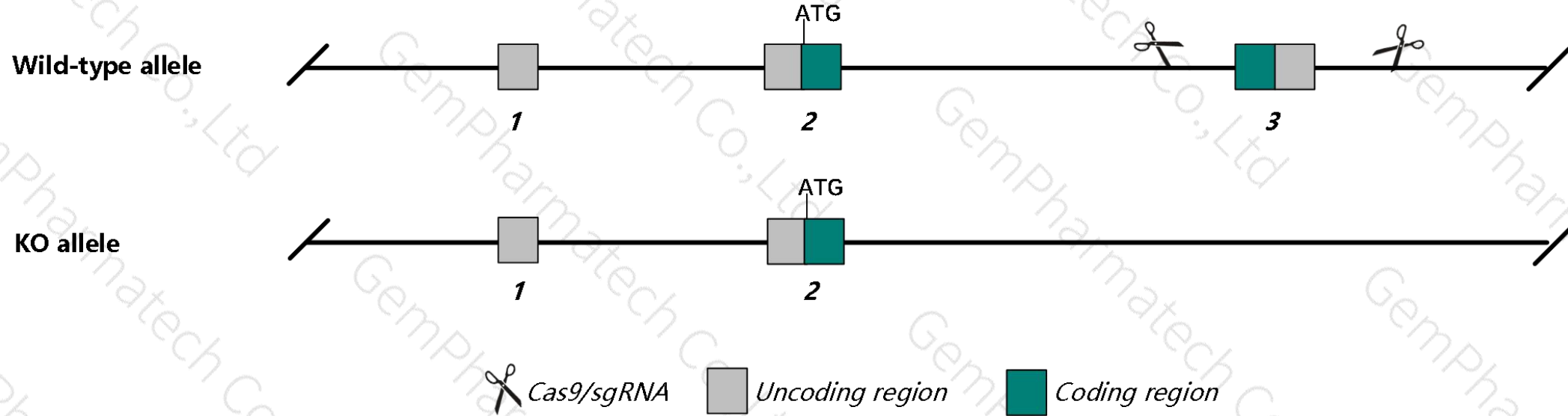
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Neu3* gene. The schematic diagram is as follows:



- The *Neu3* gene has 1 transcript. According to the structure of *Neu3* gene, exon3 of *Neu3-201* (ENSMUST00000036331.6) transcript is recommended as the knockout region. The region contains most of coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Neu3* gene. The brief process is as follows: CRISPR/Cas9 system we

- According to the existing MGI data, Mice homozygous for a knock-out allele exhibit decreased colon carcinogenesis induced by azoxymethane and dextran sodium sulfate.
- Some amino acids will remain at the N-terminus and some functions may be retained.
- The *Neu3* gene is located on the Chr7. 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)

Neu3 neuraminidase 3 [*Mus musculus* (house mouse)]

Gene ID: 50877, updated on 10-Oct-2019

Summary

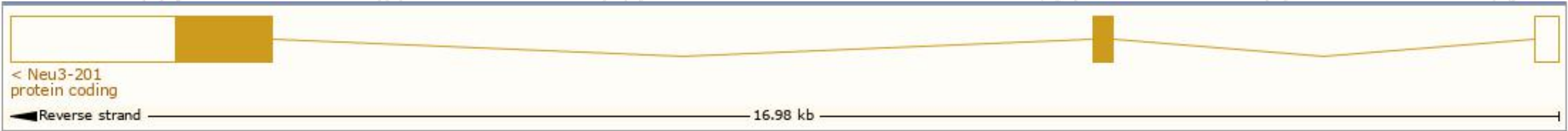
Official Symbol	Neu3 provided by MGI
Official Full Name	neuraminidase 3 provided by MGI
Primary source	MGI:MGI:1355305
See related	Ensembl:ENSMUSG00000035239
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Expression	Broad expression in thymus adult (RPKM 13.1), ovary adult (RPKM 8.7) and 19 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

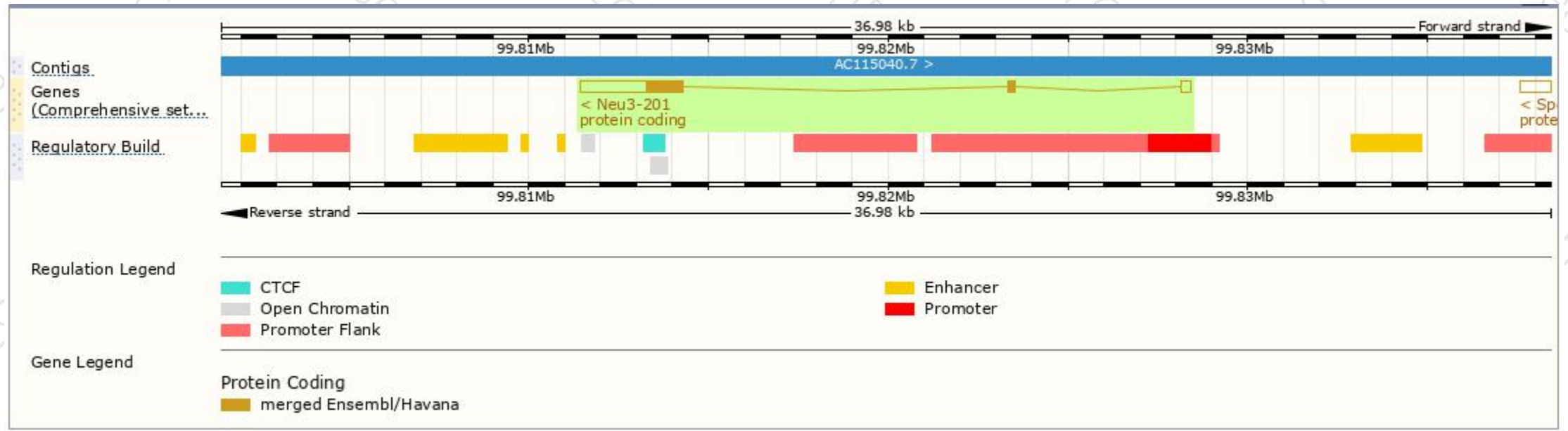
The gene has 1 transcript,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Neu3-201	ENSMUST00000036331.6	3341	418aa	Protein coding	CCDS21489	Q543I9 Q9JMH7	TSL:1 Gencode basic APPRIS P1

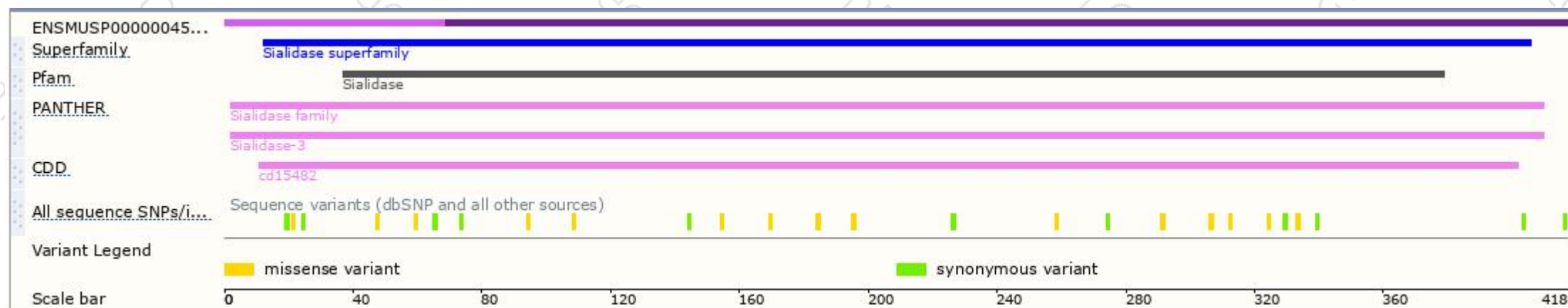
The strategy is based on the design of *Neu3-201* transcript,The transcription is shown below



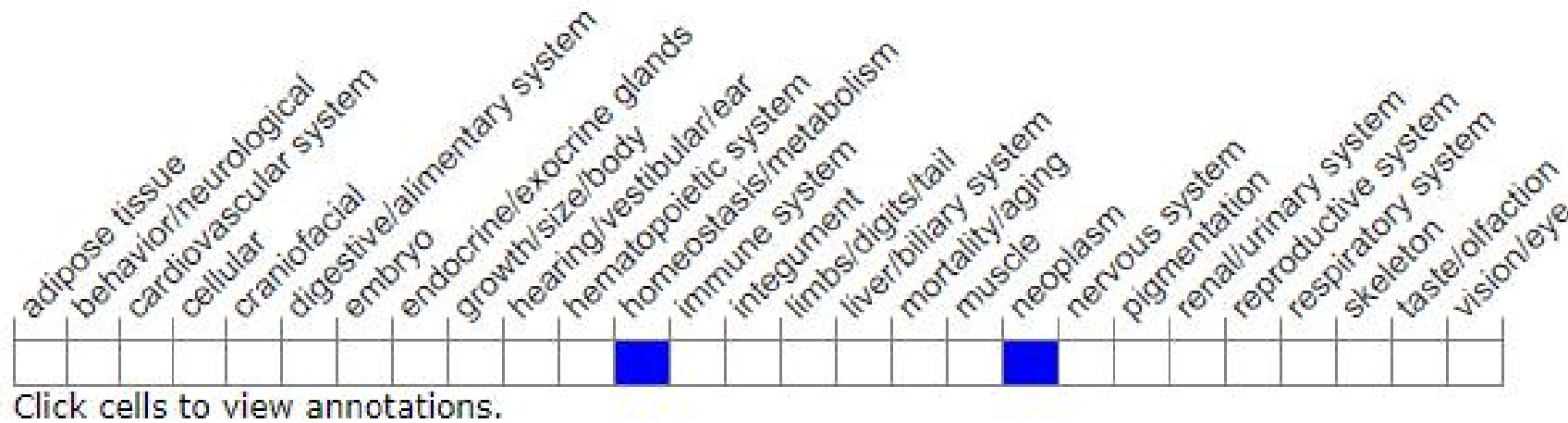
Genomic location distribution



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 a knock-out allele exhibit decreased colon carcinogenesis induced by azoxymethane and dextran sodium sulfate.

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

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