Ncan Cas9-CKO Strategy

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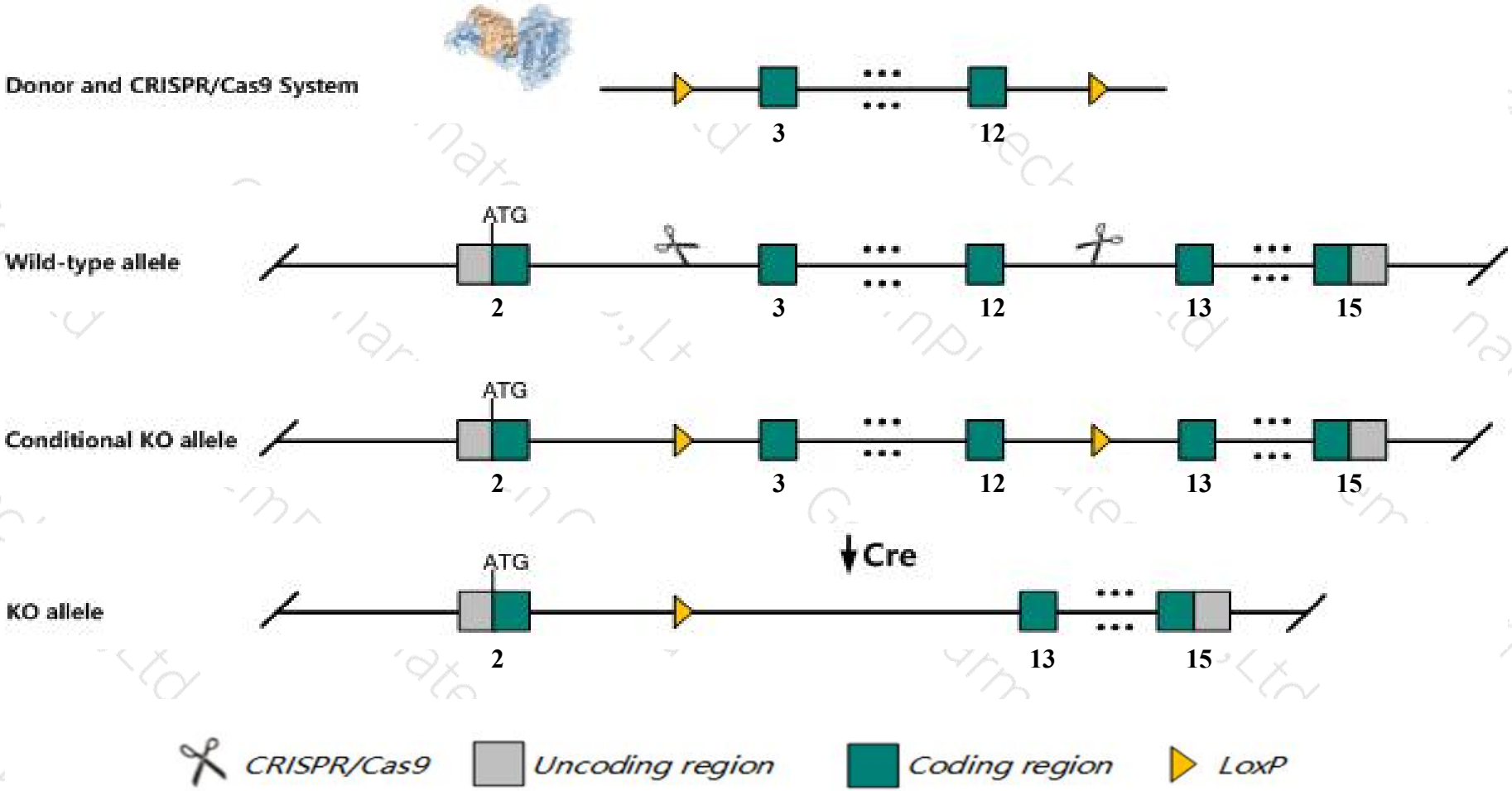
Design Date: 2019-7-29

Project Overview

Project Name	***Ncan***
Project type	**Cas9-CKO**
Strain background	**C57BL/6JGpt**

Conditional Knockout strategy

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



Technical routes

- The *Ncan* gene has 1 transcript. According to the structure of *Ncan* gene, exon3-exon12 of *Ncan-201* (ENSMUST00000002412.8) transcript is recommended as the knockout region.The region contains 3275bp coding sequence. Knock out the region will result in disruption of protein function.

- In this project we use CRISPR/Cas9 technology to modify *Ncan* gene. The brief process is as follows:CRISPR/Cas9 system and Donor 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.

- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

Notice

- According to the existing MGI data,Mice homozygous for targeted null mutations are viable and fertile and exhibit normal behavior and brain anatomy; however, mild defects in long term potentiation were noted.

- Because of *Mir7066* gene is located in *Ncan* gene, *Mir7066* gene will be deleted together in this strategy.

- The *Ncan* gene is located on the Chr8. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information （NCBI）

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Ncan neurocan [Mus musculus (house mouse)]

Gene ID: 13004, updated on 31-Jan-2019

Summary

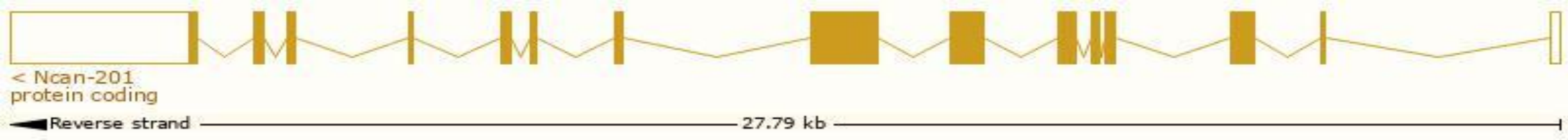
Official Symbol	Ncan provided by MGI
Official Full Name	neurocan provided by MGI
Primary source	MGI:MGI:104694
See related	Ensembl:ENSMUSG00000002341
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	C230035B04, Cspg3, Cspg3-rs, Tgfbit
Expression	Biased expression in whole brain E14.5 (RPKM 31.9), CNS E18 (RPKM 28.5) and 6 other tissues See more
Orthologs	human all

Transcript information（Ensembl）

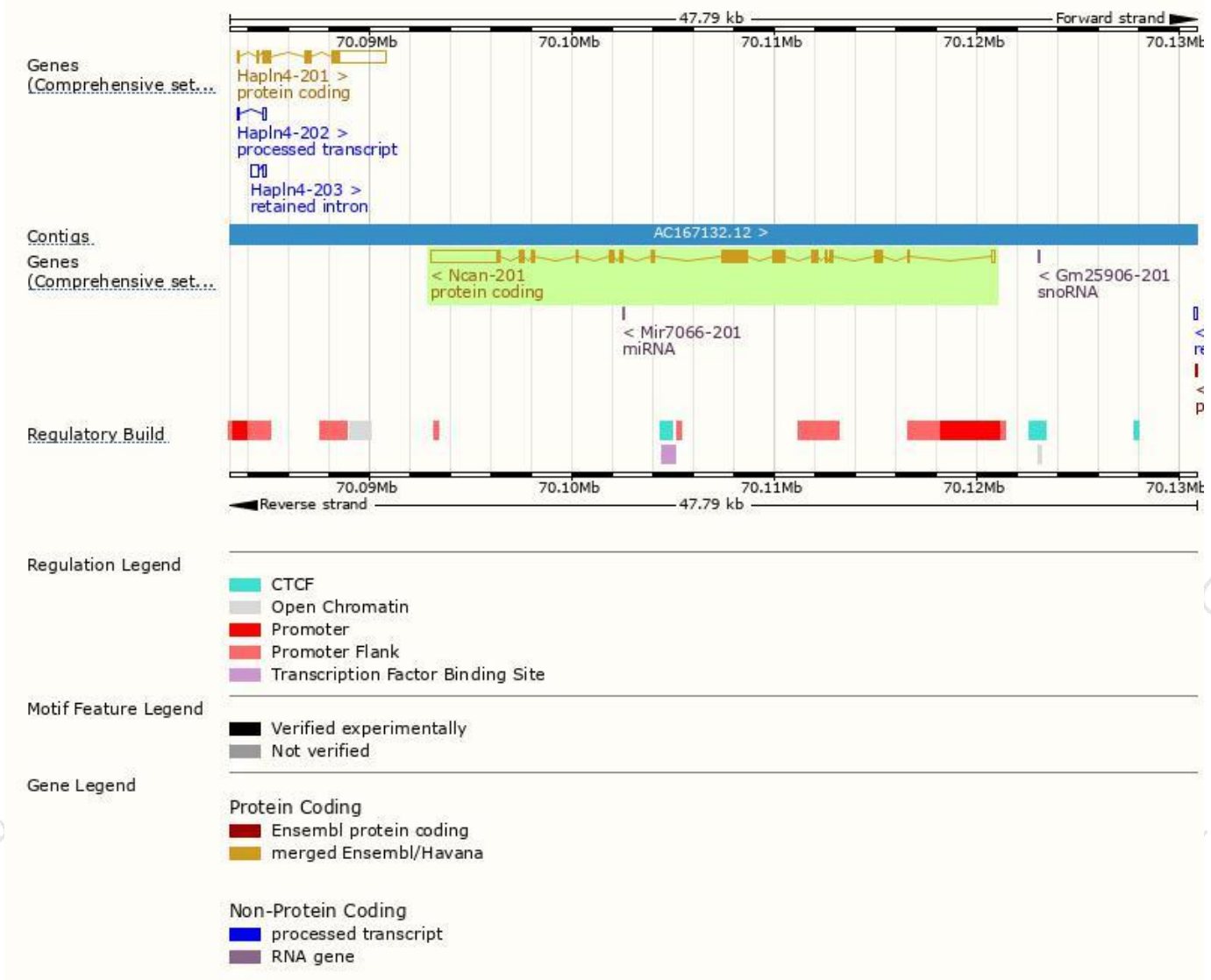
The gene has 1 transcript,and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ncan-201	ENSMUST00000002412.8	7195	1268aa	Protein coding	CCDS22358	A0A0R4IZX5	TSL:1 GENCODE basic APPRIS P1

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

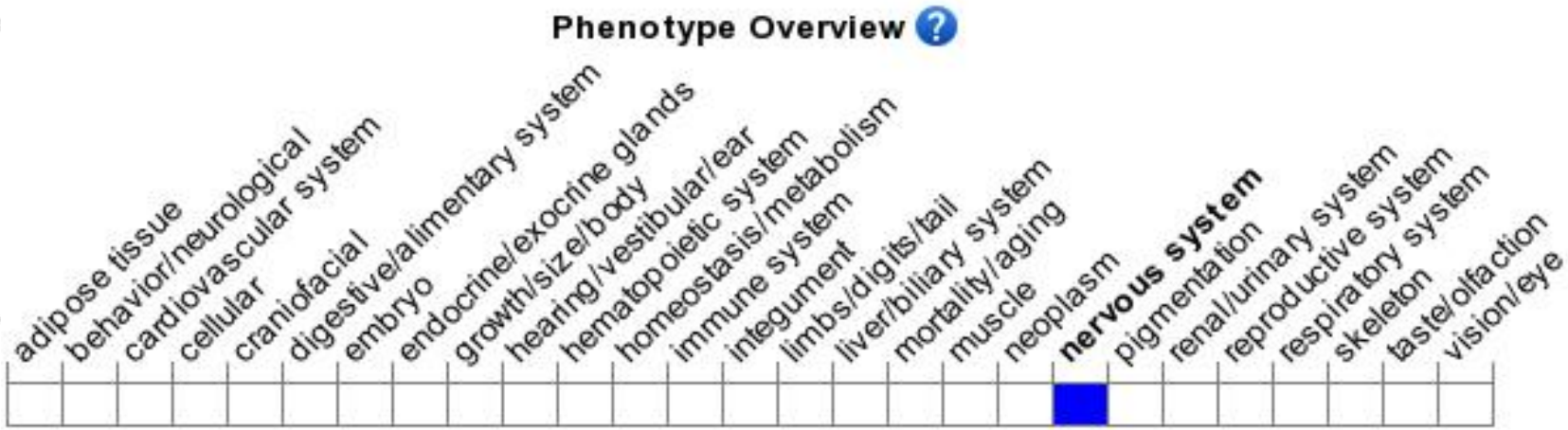


Genomic location distribution



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/).

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

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