

Ndufa9 Cas9-CKO Strategy

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Reviewer:

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Design Date:

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Project Overview

Project Name

Ndufa9

Project type

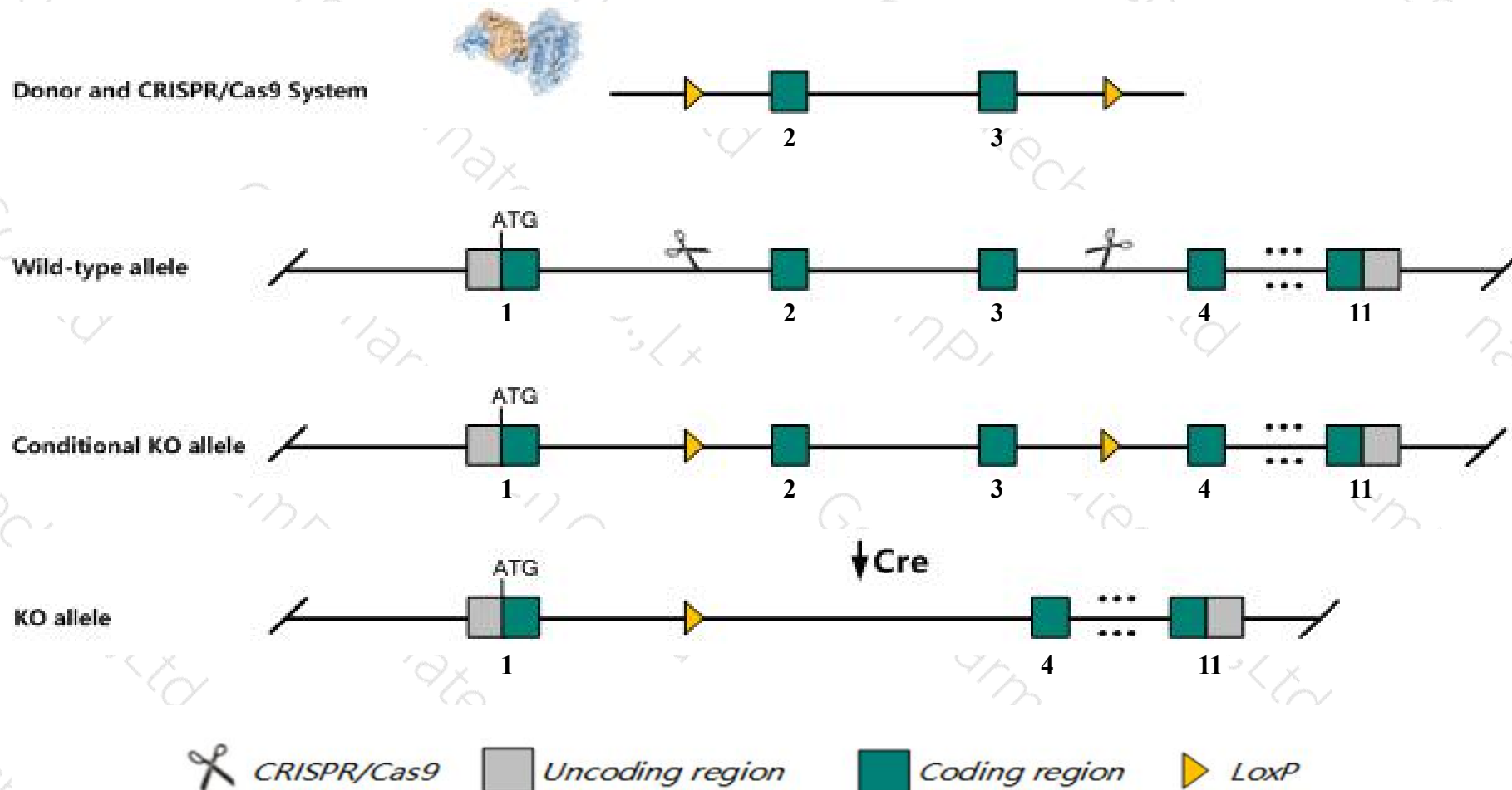
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Ndufa9* gene. The schematic diagram is as follows:



- The *Ndufa9* gene has 3 transcripts. According to the structure of *Ndufa9* gene, exon2-exon3 of *Ndufa9*-203 (ENSMUST00000205002.2) transcript is recommended as the knockout region. The region contains 269bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Ndufa9* 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

- The *Ndufa9* gene is located on the Chr6. 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)

Ndufa9 NADH:ubiquinone oxidoreductase subunit A9 [Mus musculus (house mouse)]

Gene ID: 66108, updated on 7-Apr-2019

Summary



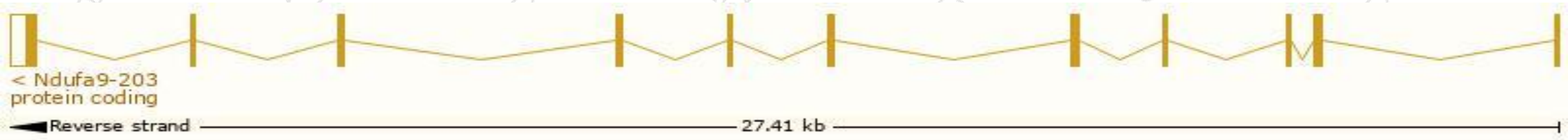
Official Symbol	Ndufa9 provided by MGI
Official Full Name	NADH:ubiquinone oxidoreductase subunit A9 provided by MGI
Primary source	MGI:MGI:1913358
See related	Ensembl:ENSMUSG000000000399
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	1010001N11Rik
Expression	Ubiquitous expression in heart adult (RPKM 197.9), kidney adult (RPKM 121.9) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

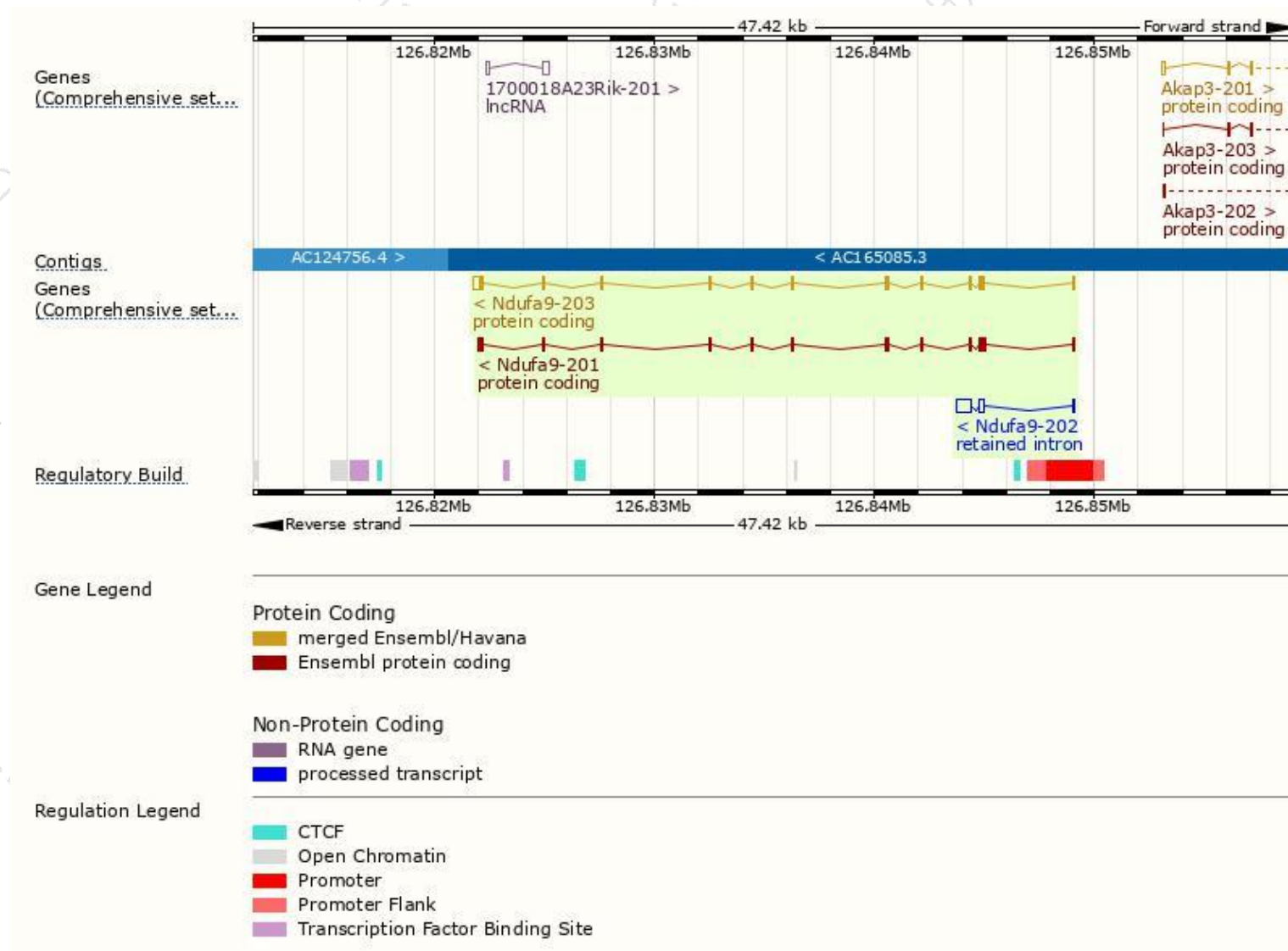
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ndufa9-203	ENSMUST00000205002.2	1445	377aa	Protein coding	CCDS20557	Q9DC69	TSL:1 GENCODE basic APPRIS P2
Ndufa9-201	ENSMUST00000088194.6	1238	373aa	Protein coding	-	A0A0R3P9C8	TSL:5 GENCODE basic APPRIS ALT2
Ndufa9-202	ENSMUST00000200890.1	919	No protein	Retained intron	-	-	TSL:2

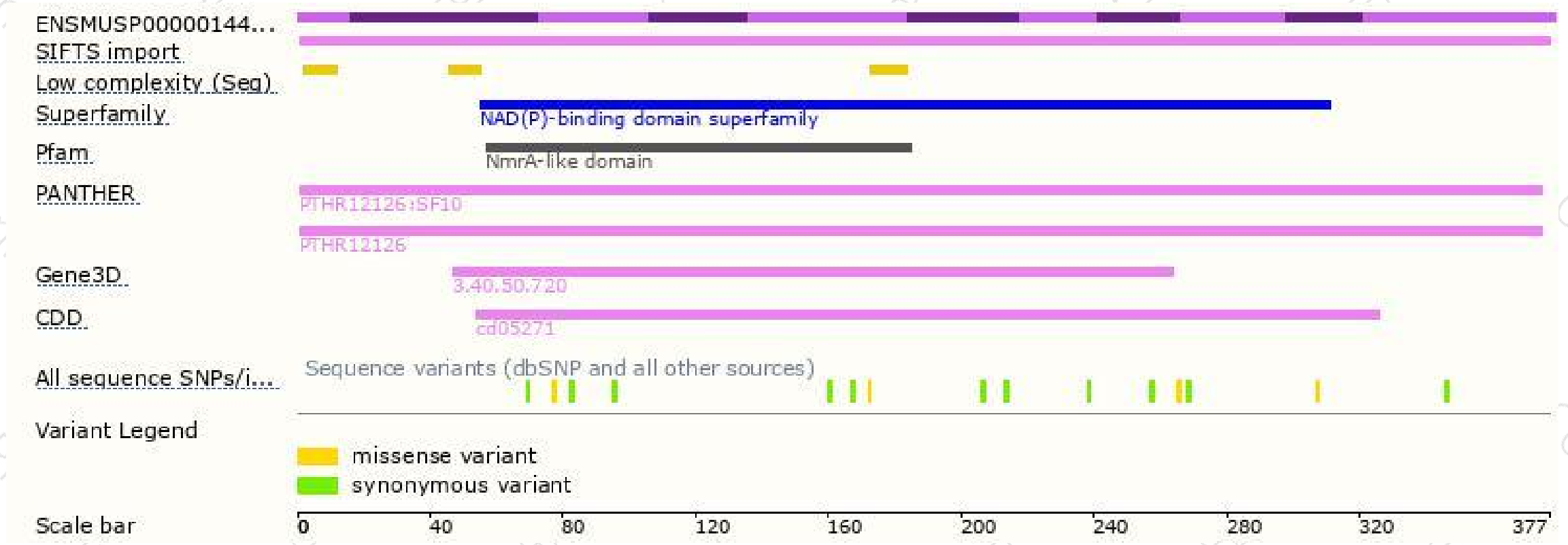
The strategy is based on the design of *Ndufa9-203* transcript,The transcription is shown below



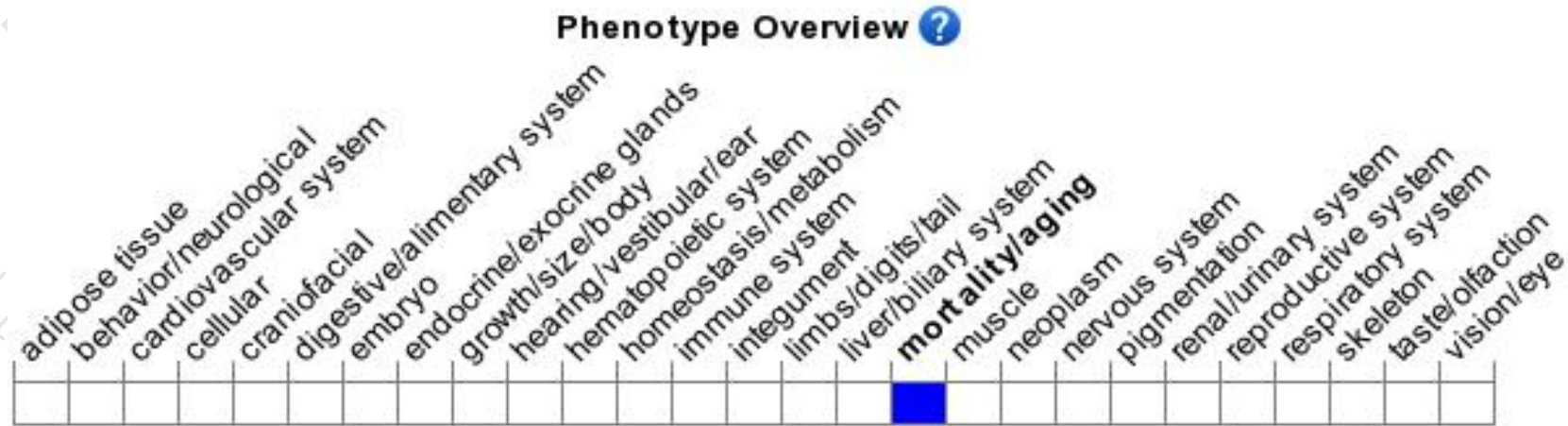
Genomic location distribution



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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