

Dnajb12 Cas9-CKO Strategy

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Project Overview

Project Name

Dnajb12

Project type

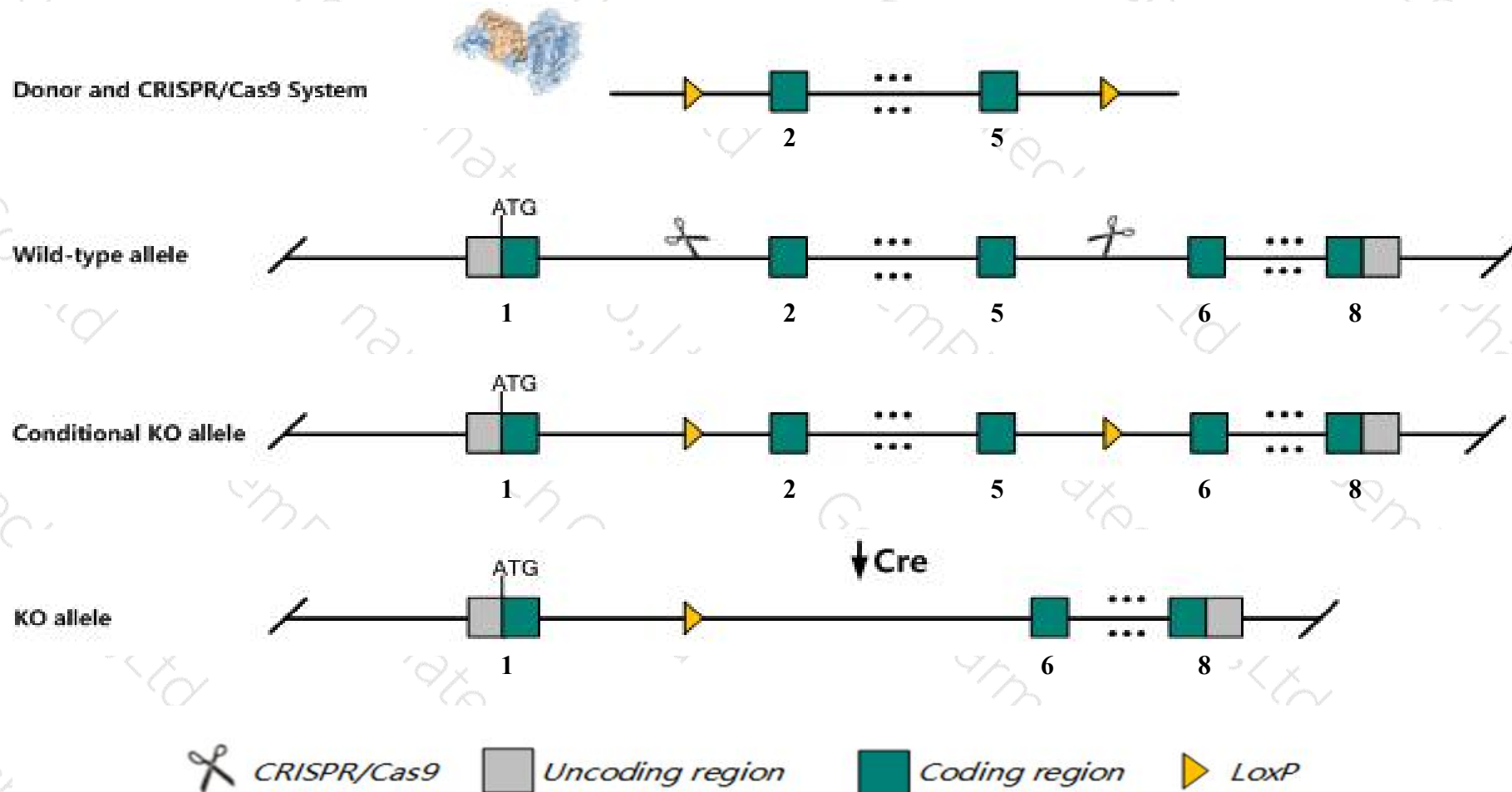
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Dnajb12* gene. The schematic diagram is as follows:



- The *Dnajb12* gene has 5 transcripts. According to the structure of *Dnajb12* gene, exon2-exon5 of *Dnajb12*-204 (ENSMUST00000146590.7) transcript is recommended as the knockout region. The region contains 593bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Dnajb12* 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 *Dnajb12* gene is located on the Chr10. 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)

Dnajb12 DnaJ heat shock protein family (Hsp40) member B12 [Mus musculus (house mouse)]

Gene ID: 56709, updated on 13-Mar-2020

Summary



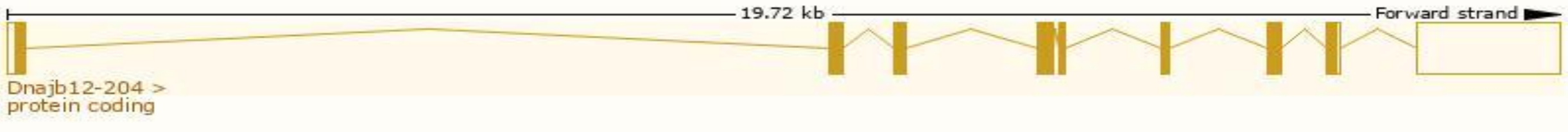
Official Symbol	Dnajb12 provided by MGI
Official Full Name	DnaJ heat shock protein family (Hsp40) member B12 provided by MGI
Primary source	MGI:MGI:1931881
See related	Ensembl:ENSMUSG00000020109
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Dj10, mDj10
Expression	Ubiquitous expression in kidney adult (RPKM 29.5), adrenal adult (RPKM 24.3) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

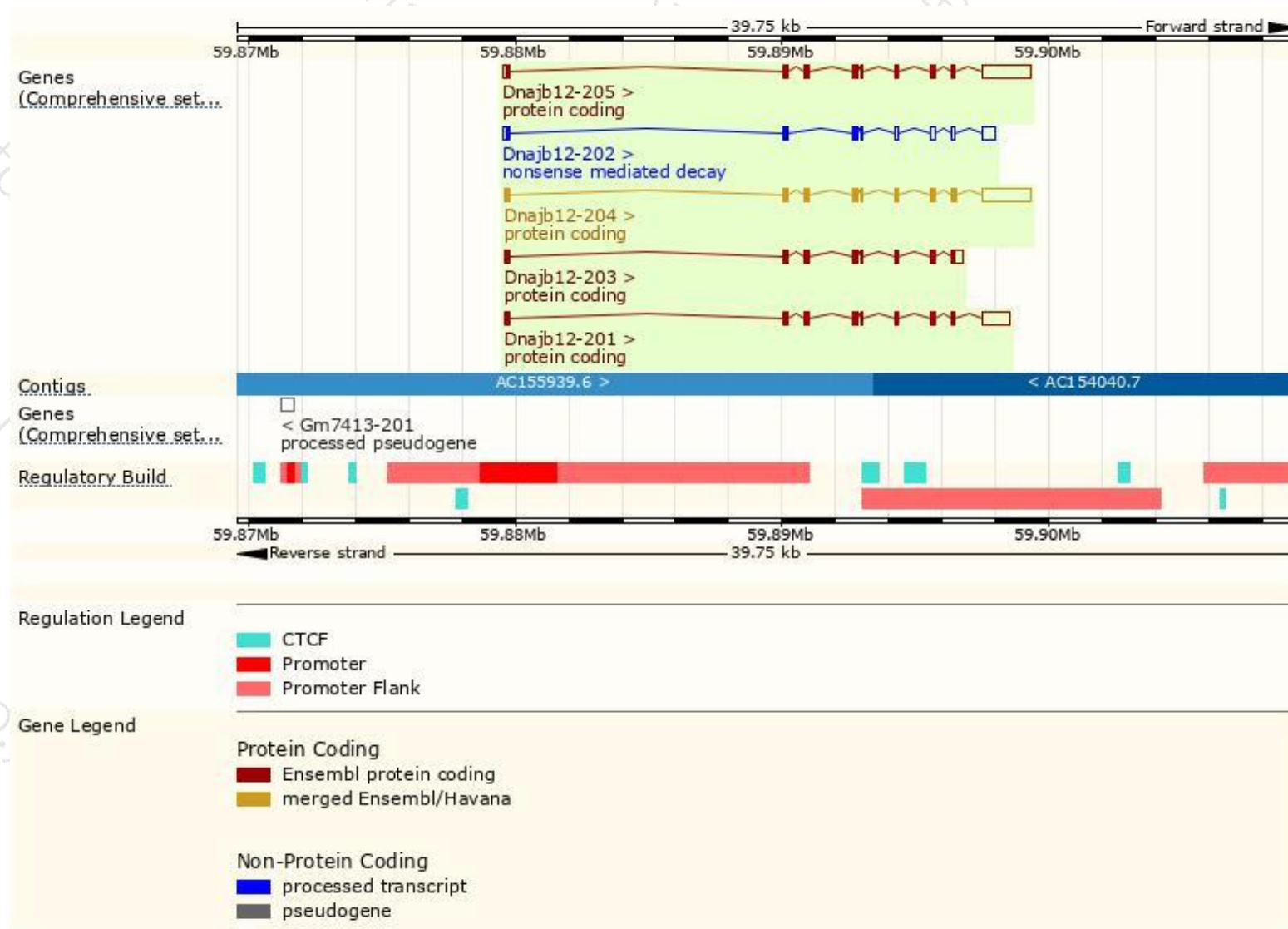
The gene has 5 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Dnajb12-201	ENSMUST00000020309.6	2238	376aa	Protein coding	CCDS23867	Q9QYI4	TSL:1 GENCODE basic APPRIS P2
Dnajb12-202	ENSMUST00000131810.7	1647	126aa	Nonsense mediated decay	-	D6RHR9	TSL:1
Dnajb12-203	ENSMUST00000142819.7	1495	376aa	Protein coding	CCDS23867	Q9QYI4	TSL:1 GENCODE basic APPRIS P2
Dnajb12-204	ENSMUST00000146590.7	3070	376aa	Protein coding	CCDS23867	Q9QYI4	TSL:1 GENCODE basic APPRIS P2
Dnajb12-205	ENSMUST00000147914.7	3041	378aa	Protein coding	-	Q8C4C9	TSL:1 GENCODE basic APPRIS ALT2

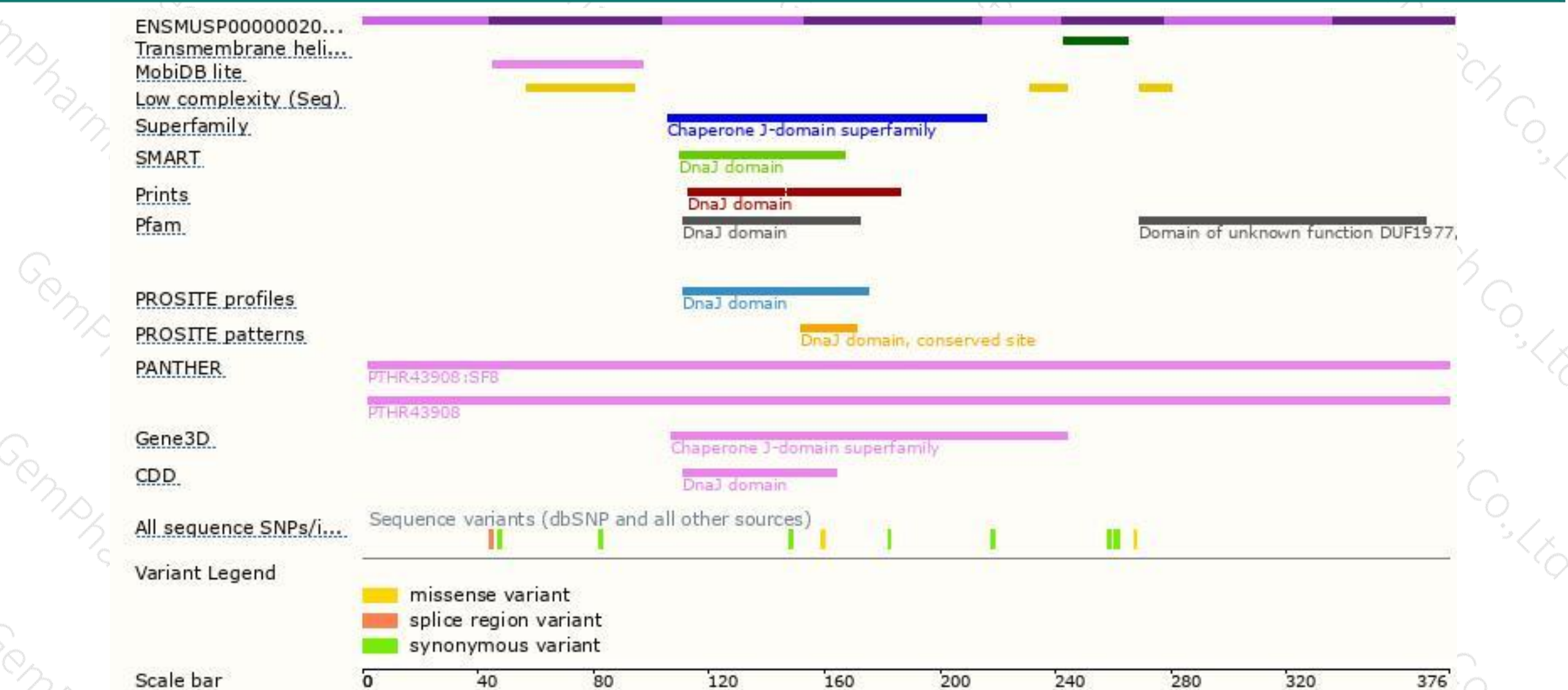
The strategy is based on the design of *Dnajb12-204* transcript,The transcription is shown below



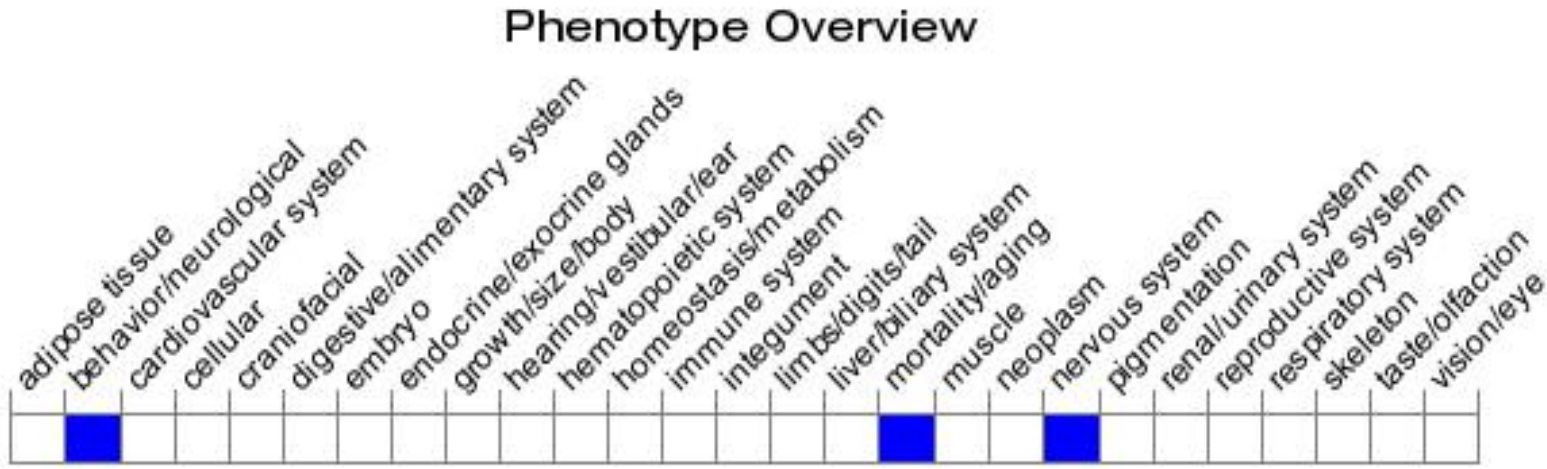
Genomic location distribution



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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