

Rnf182 Cas9-CKO Strategy

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Project Overview

Project Name

Rnf182

Project type

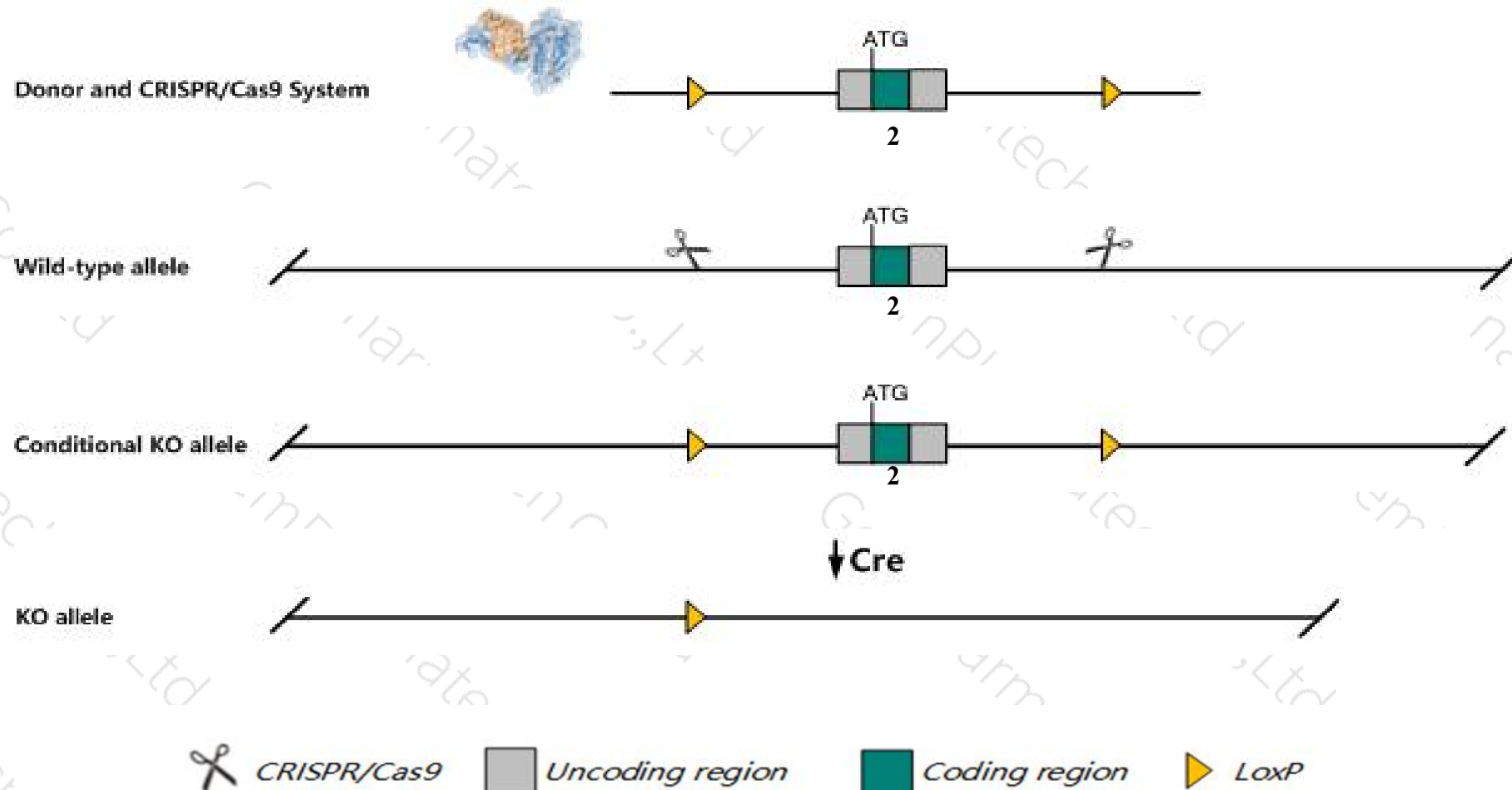
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Rnfl82* gene. The schematic diagram is as follows:



- The *Rnf182* gene has 2 transcripts. According to the structure of *Rnf182* gene, exon2 of *Rnf182-201* (ENSMUST00000059986.2) transcript is recommended as the knockout region. The region contains all of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Rnf182* 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 *Rnfl82* gene is located on the Chr13. 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)

Rnf182 ring finger protein 182 [Mus musculus (house mouse)]

Gene ID: 328234, updated on 31-Jan-2019

Summary



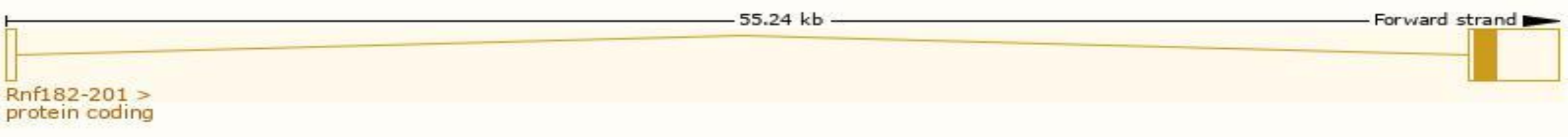
Official Symbol	Rnf182 provided by MGI
Official Full Name	ring finger protein 182 provided by MGI
Primary source	MGI:MGI:3045355
See related	Ensembl:ENSMUSG00000044164
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	C630023L15Rik
Expression	Biased expression in CNS E18 (RPKM 7.8), whole brain E14.5 (RPKM 6.5) and 8 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

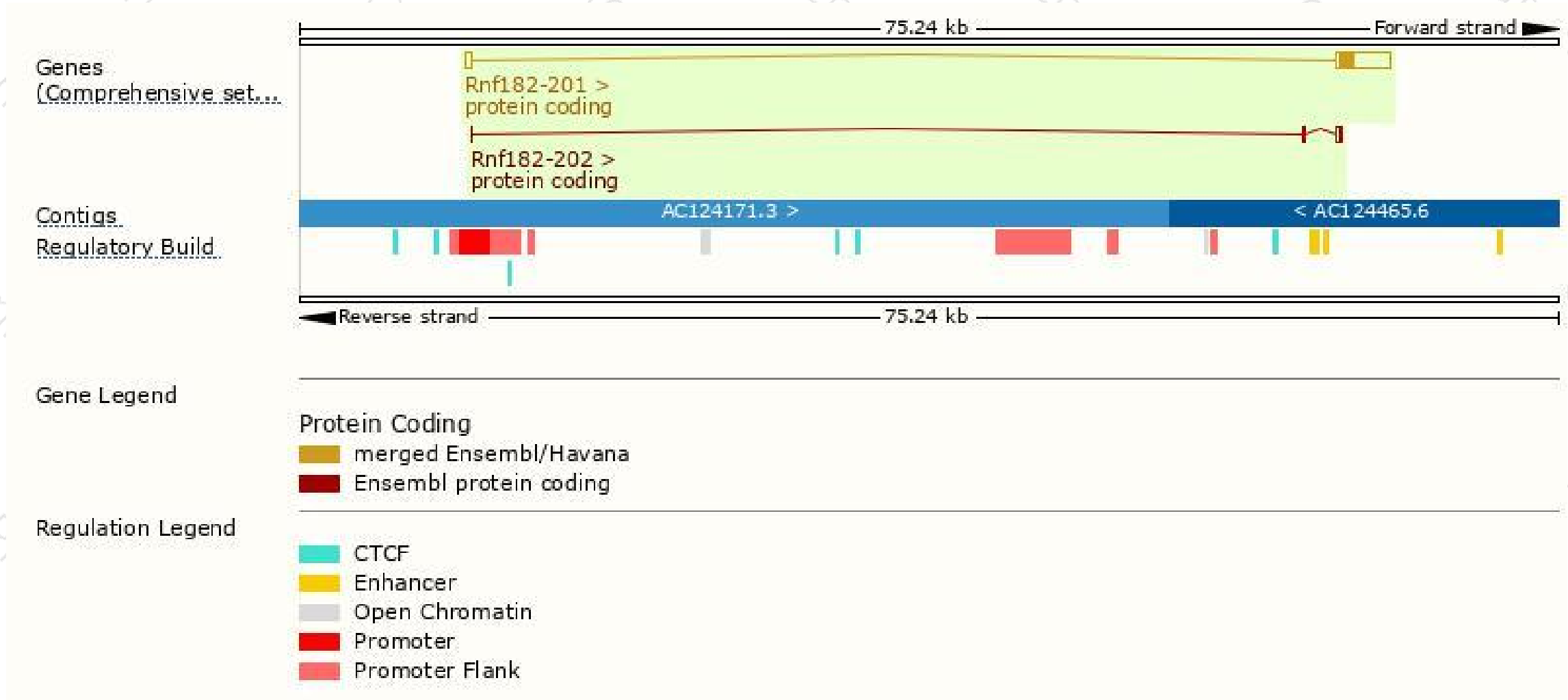
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Rnf182-201	ENSMUST00000059986.2	3544	247aa	Protein coding	CCDS26480	Q8C432	TSL:1 GENCODE basic APPRIS P1
Rnf182-202	ENSMUST00000161817.1	457	30aa	Protein coding	-	E0CZ87	CDS 3' incomplete TSL:3

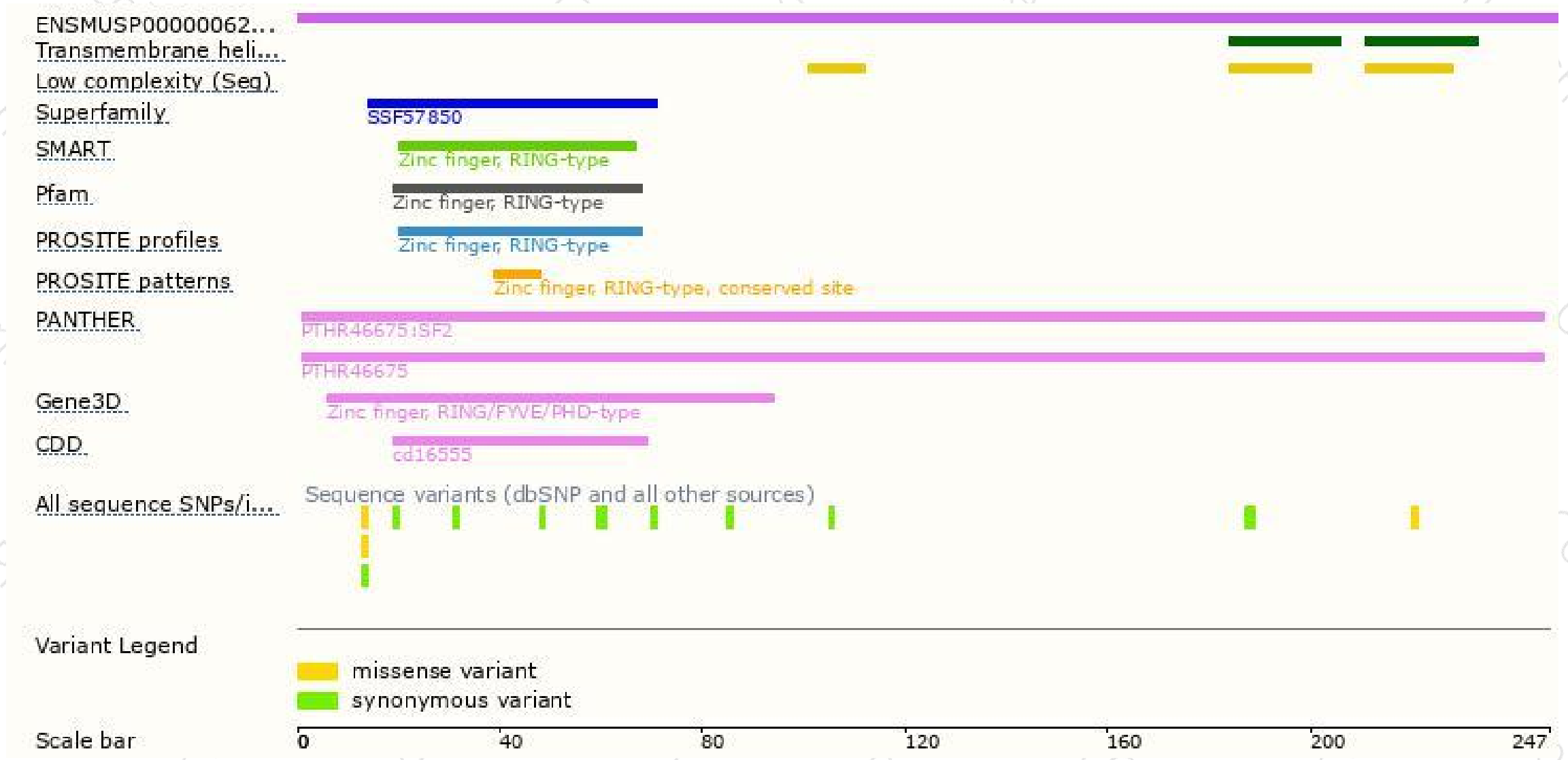
The strategy is based on the design of *Rnf182-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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