

# *Nfu1* Cas9-KO Strategy

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**Reviewer: Shuang Zhang**

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# Project Overview

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**Project Name**

*Nfu1*

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**Project type**

**Cas9-KO**

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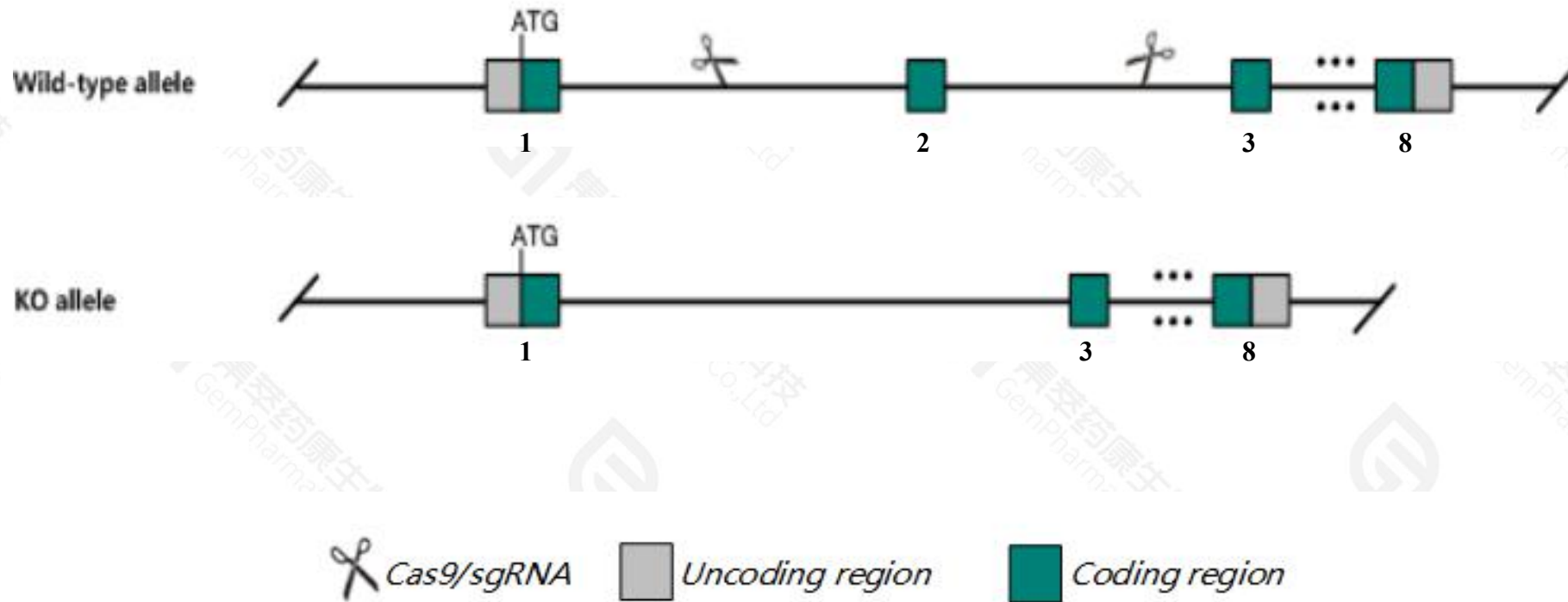
**Strain background**

**C57BL/6JGpt**

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# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Nfu1* gene. The schematic diagram is as follows:



- The *Nfu1* gene has 9 transcripts. According to the structure of *Nfu1* gene, exon2 of *Nfu1-201*(ENSMUST00000032060.15) transcript is recommended as the knockout region. The region contains 101bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Nfu1* gene. The brief process is as follows: sgRNA was transcribed in vitro. Cas9 and sgRNA 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 *Nfu1* 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

# Gene information (NCBI)

## Nfu1 NFU1 iron-sulfur cluster scaffold [Mus musculus (house mouse)]

Gene ID: 56748, updated on 10-Oct-2020

### Summary



**Official Symbol** Nfu1 provided by [MGI](#)

**Official Full Name** NFU1 iron-sulfur cluster scaffold provided by [MGI](#)

**Primary source** [MGI:MGI:1913290](#)

**See related** [Ensembl:ENSMUSG00000029993](#)

**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** 0610006G17Rik, CGI-3, CGI-33, Hi, Hirip5

**Expression** Ubiquitous expression in kidney adult (RPKM 20.2), CNS E18 (RPKM 18.7) and 28 other tissues [See more](#)

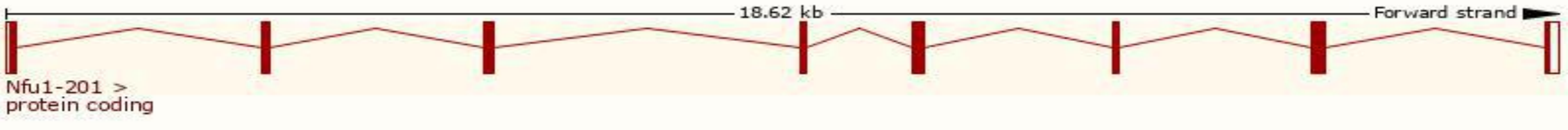
**Orthologs** [human](#) [all](#)

# Transcript information (Ensembl)

The gene has 9 transcripts,all transcripts are shown below:

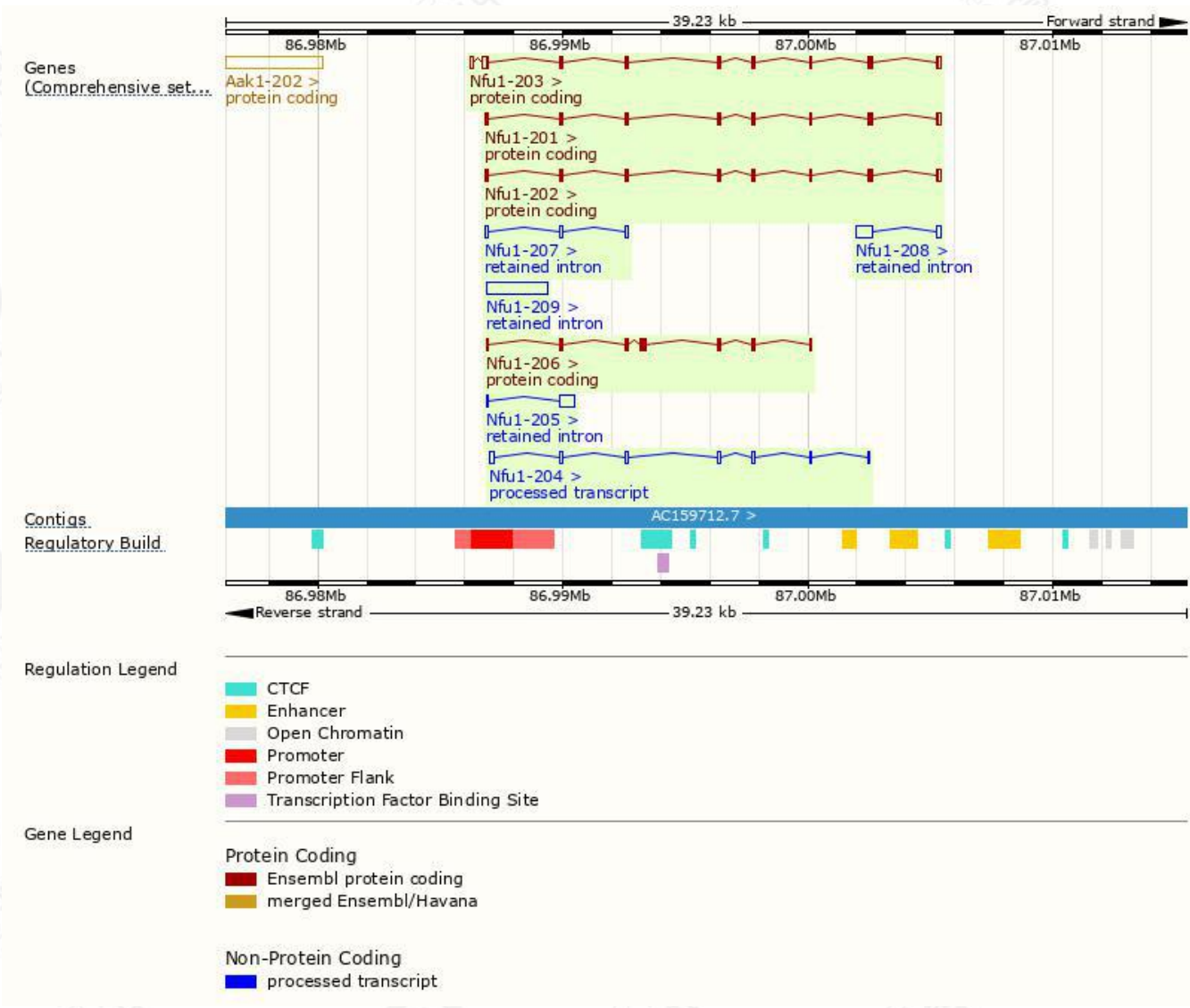
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Nfu1-203	<a href="#">ENSMUST00000120240.8</a>	1195	<a href="#">256aa</a>	Protein coding	<a href="#">CCDS51836</a>		TSL:5 , GENCODE basic , APPRIS P3 ,
Nfu1-202	<a href="#">ENSMUST00000117583.8</a>	950	<a href="#">255aa</a>	Protein coding	<a href="#">CCDS85088</a>		TSL:1 , GENCODE basic , APPRIS ALT2 ,
Nfu1-201	<a href="#">ENSMUST00000032060.15</a>	948	<a href="#">256aa</a>	Protein coding	<a href="#">CCDS51836</a>		TSL:1 , GENCODE basic , APPRIS P3 ,
Nfu1-206	<a href="#">ENSMUST00000144776.7</a>	777	<a href="#">255aa</a>	Protein coding	-		CDS 3' incomplete , TSL:3 ,
Nfu1-204	<a href="#">ENSMUST00000127819.2</a>	716	No protein	Processed transcript	-		TSL:3 ,
Nfu1-209	<a href="#">ENSMUST00000205018.2</a>	2508	No protein	Retained intron	-		TSL:NA ,
Nfu1-208	<a href="#">ENSMUST00000204103.2</a>	829	No protein	Retained intron	-		TSL:2 ,
Nfu1-205	<a href="#">ENSMUST00000140343.3</a>	655	No protein	Retained intron	-		TSL:2 ,
Nfu1-207	<a href="#">ENSMUST00000203512.3</a>	371	No protein	Retained intron	-		TSL:2 ,

The strategy is based on the design of *Nfu1-201* transcript,the transcription is shown below:



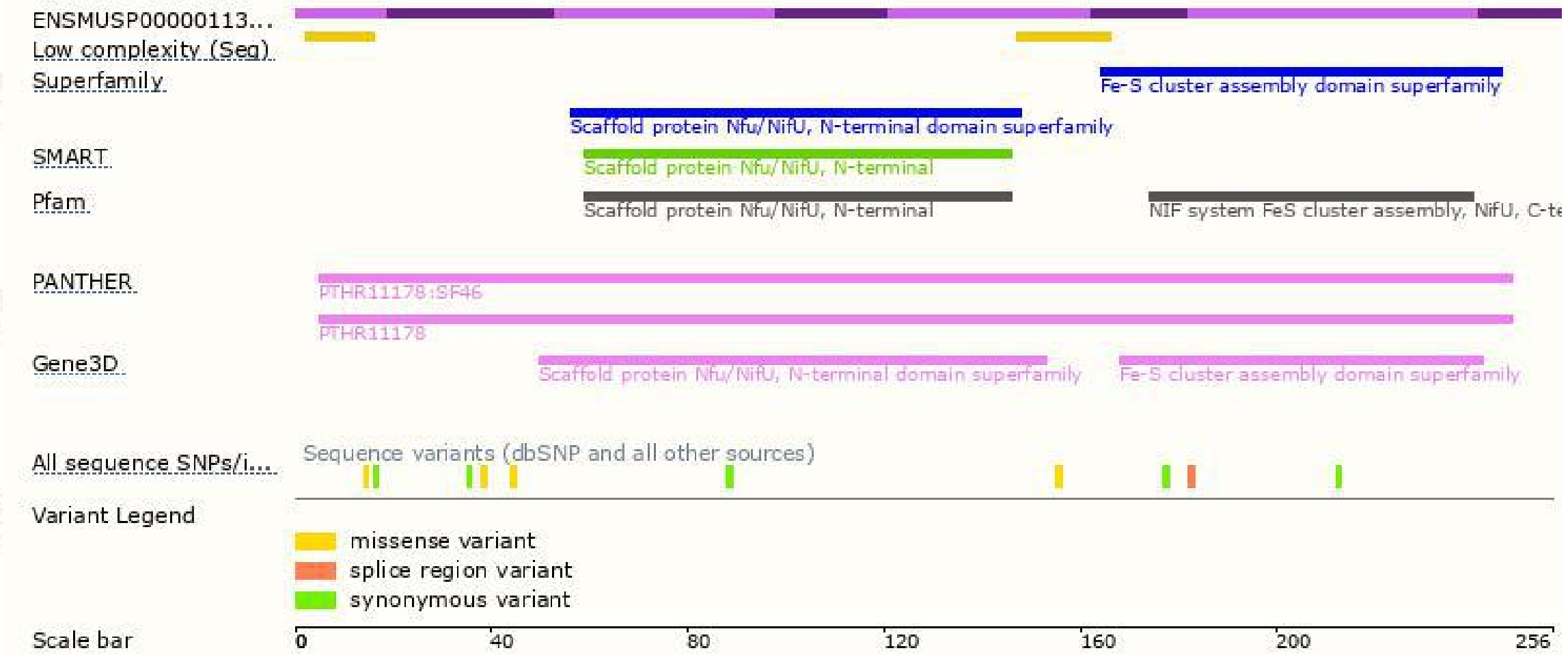


# Genomic location distribution





# Protein domain



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

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