

# ***Fnip1*** Cas9-KO Strategy

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# Project Overview

**Project Name**

***Fnip1***

**Project type**

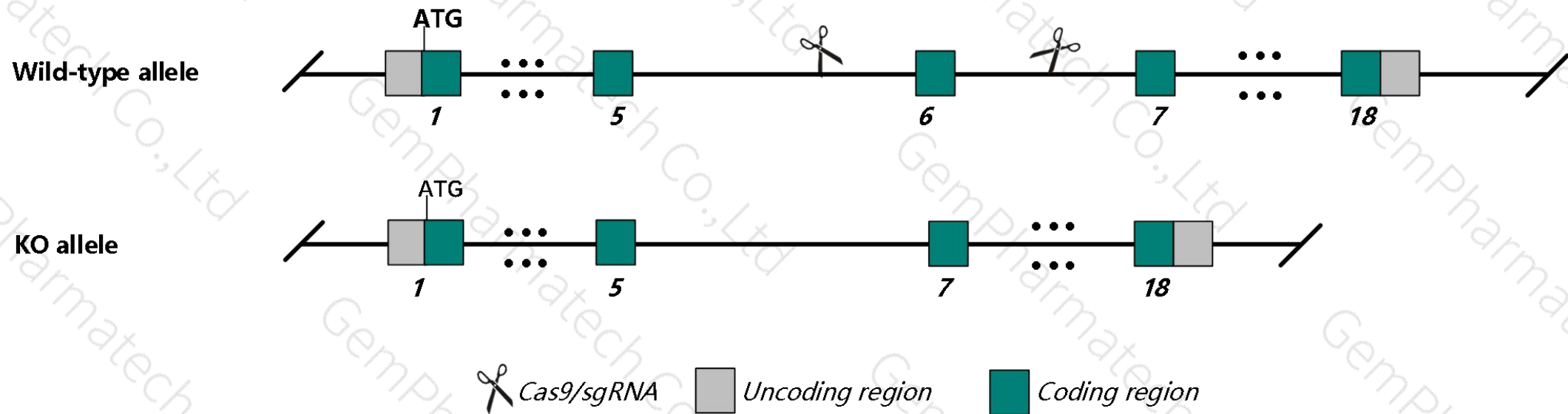
**Cas9-KO**

**Strain background**

**C57BL/6J**

# Knockout strategy

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



- The *Fnip1* gene has 2 transcripts. According to the structure of *Fnip1* gene, exon6 of *Fnip1-201* (ENSMUST00000046835.13) transcript is recommended as the knockout region. The region contains 92bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Fnip1* gene. The brief process is as follows: sgRNA was transcribed in vitro. Cas9 and sgRNA were microinjected into the fertilized eggs of C57BL/6J 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/6J mice.

- According to the existing MGI data, mice homozygous for an ENU-induced or targeted allele exhibit arrested B cell development at the pre-B cell stage with increased B cell apoptosis.
- The *Fnip1* gene is located on the Chr11. 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 gene transcription and translation processes, all risks cannot be predicted under existing information.



# Gene information (NCBI)

## Fnip1 folliculin interacting protein 1 [Mus musculus (house mouse)]

Gene ID: 216742, updated on 31-Jan-2019

### Summary



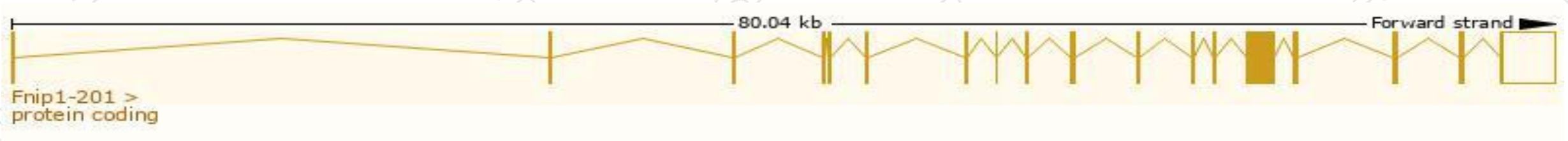
<b>Official Symbol</b>	Fnip1 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	folliculin interacting protein 1 provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:2444668</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000035992</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	VALIDATED
<b>Organism</b>	<a href="#">Mus musculus</a>
<b>Lineage</b>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
<b>Also known as</b>	A730024A03Rik, AI838773, AW557298
<b>Expression</b>	Ubiquitous expression in bladder adult (RPKM 4.7), placenta adult (RPKM 4.6) and 28 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

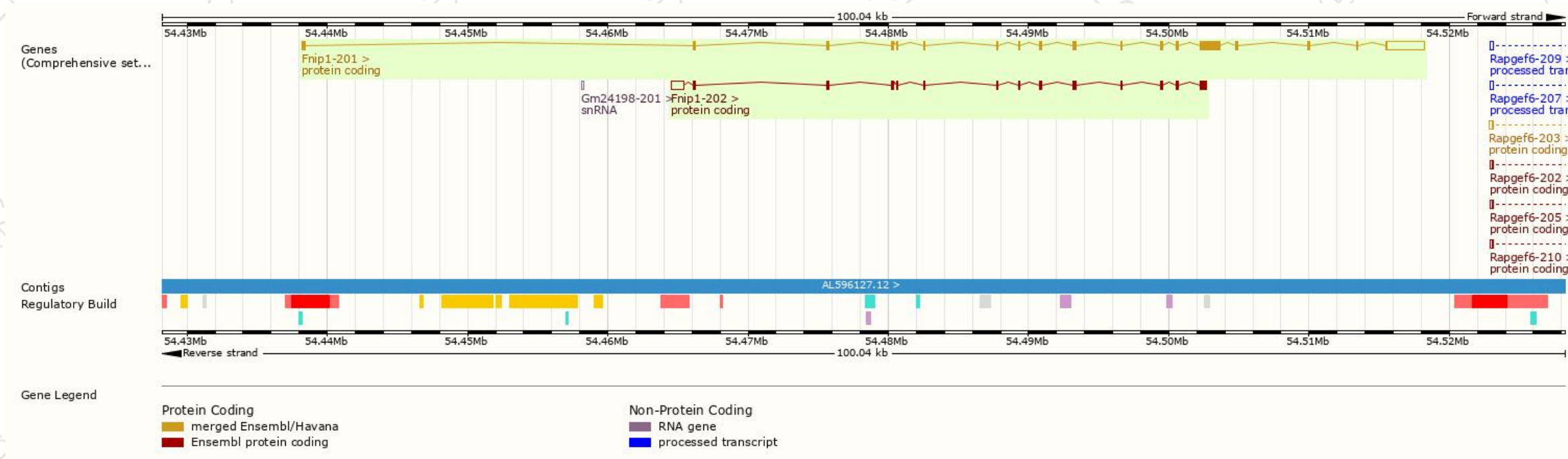
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Fnip1-201	<a href="#">ENSMUST00000046835.13</a>	6255	<a href="#">1165aa</a>	Protein coding	<a href="#">CCDS24698</a>	<a href="#">Q68FD7</a>	TSL:1 GENCODE basic APPRIS P1
Fnip1-202	<a href="#">ENSMUST00000143650.1</a>	2738	<a href="#">631aa</a>	Protein coding	-	<a href="#">F6RJ64</a>	CDS 3' incomplete TSL:1

The strategy is based on the design of *Fnip1-201* transcript,The transcription is shown below

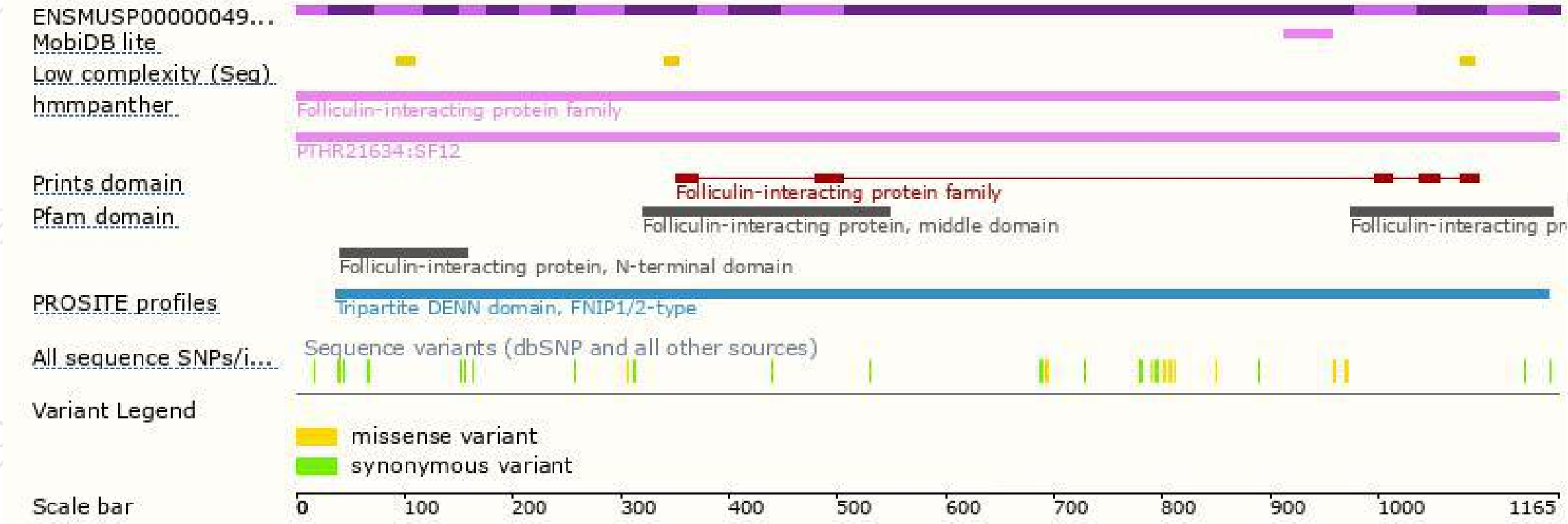


# Genomic location distribution

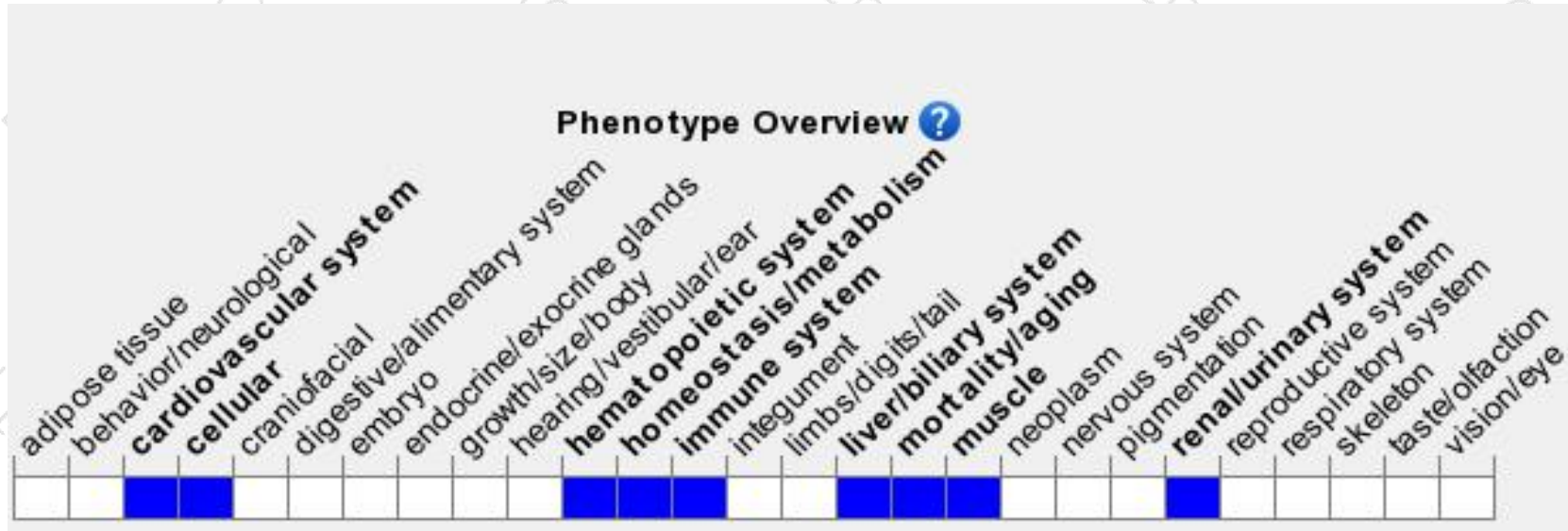




# Protein domain



# Mouse phenotype description(MGI )



*Phenotypes affected by the gene are marked in blue. Data quoted from MGI database(<http://www.informatics.jax.org/>).*

According to the existing MGI data, Mice homozygous for an ENU-induced or targeted allele exhibit arrested B cell development at the pre-B cell stage with increased B cell apoptosis.

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

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