

# *T015738 Zfp219* Cas9-KO Strategy

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# Project Overview

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**Project Name**

*Zfp219*

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**Project type**

**Cas9-KO**

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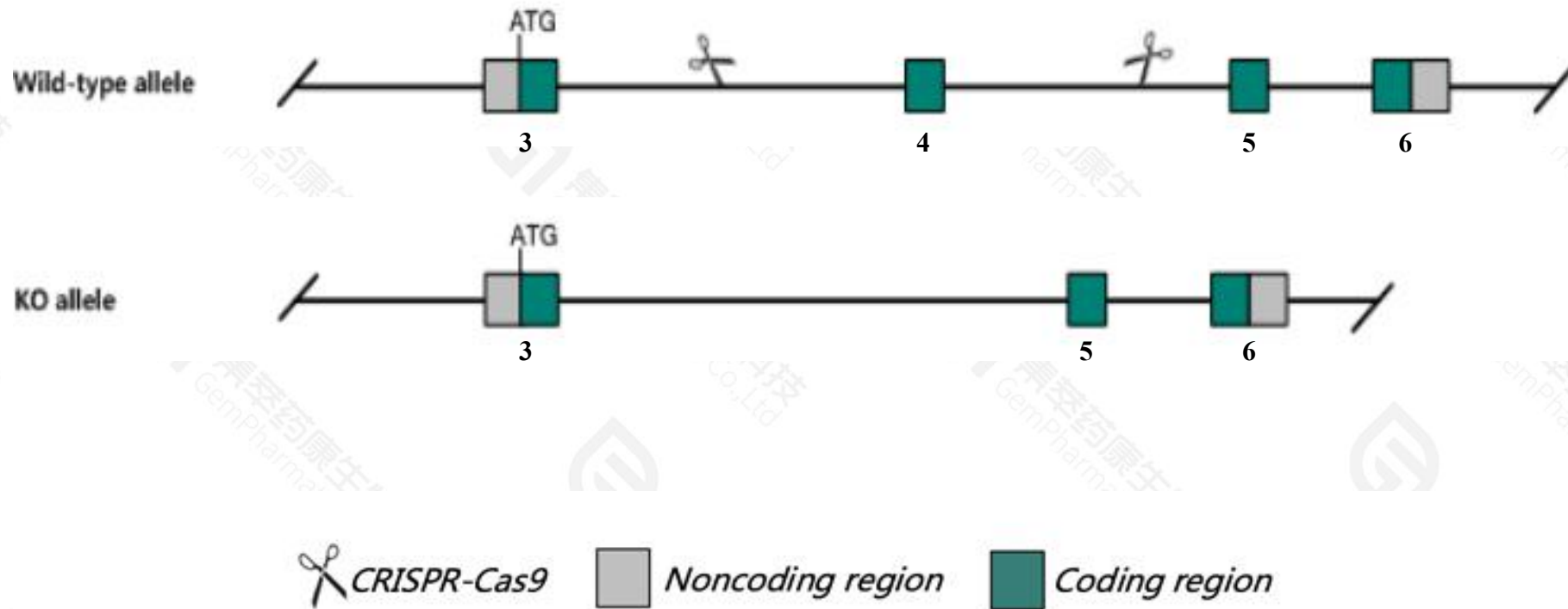
**Strain background**

**C57BL/6JGpt**

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# Knockout strategy

This model will use CRISPR-Cas9 technology to edit the *Zfp219* gene. The schematic diagram is as follows:



- The *Zfp219* gene has 13 transcripts. According to the structure of *Zfp219* gene, exon4 of *Zfp219*-202(ENSMUST00000166169.4) transcript is recommended as the knockout region. The region contains 1435bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR-Cas9 technology to modify *Zfp219* gene. The brief process is as follows: CRISPR-Cas9 system 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 *Zfp219* gene is located on the Chr14. 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.

## Zfp219 zinc finger protein 219 [Mus musculus (house mouse)]

Gene ID: 69890, updated on 23-Jun-2022

### Summary



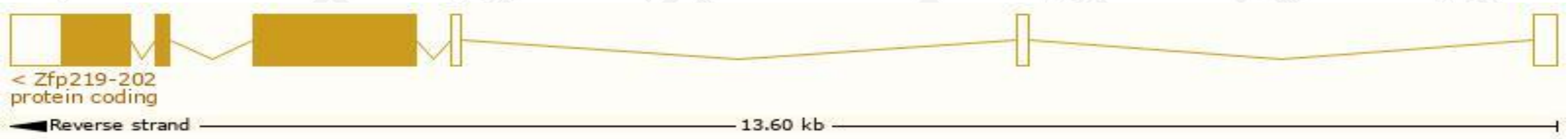
<b>Official Symbol</b>	Zfp219 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	zinc finger protein 219 provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:1917140</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000049295</a>
<b>Gene type</b>	protein coding
<b>RefSeq 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>	2010302A17Rik, Znf219
<b>Expression</b>	Ubiquitous expression in thymus adult (RPKM 27.5), CNS E11.5 (RPKM 23.7) 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 13 transcripts,all transcripts are shown below:

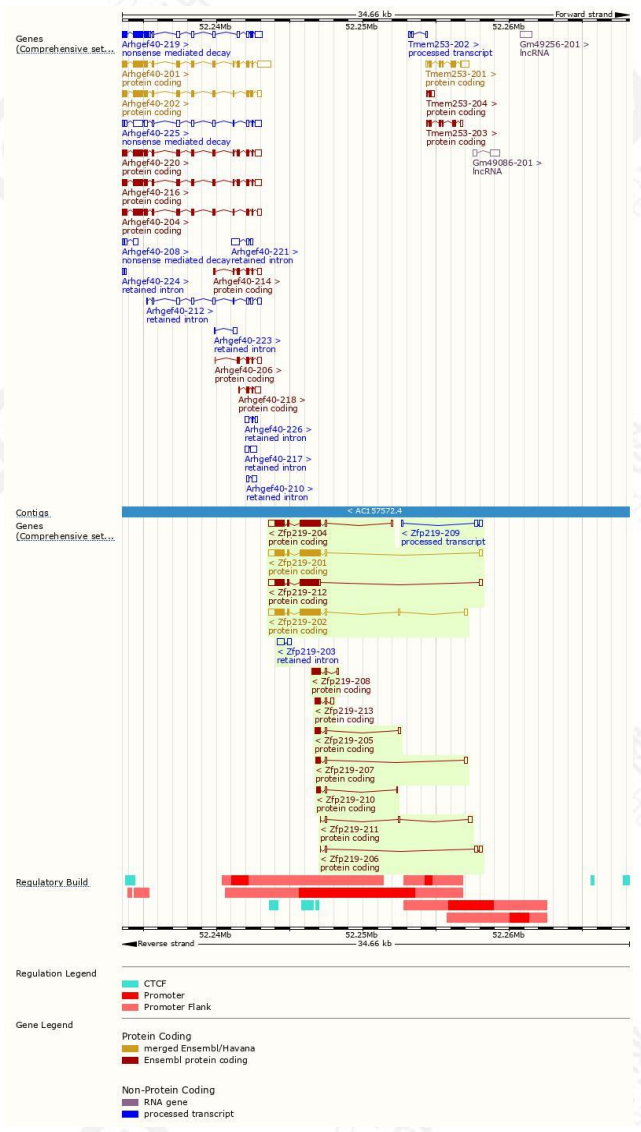
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Zfp219-202	<a href="#">ENSMUST00000166169.4</a>	3017	<a href="#">726aa</a>	Protein coding	<a href="#">CCDS36916</a>		TSL:1 , GENCODE basic , APPRIS P2 ,
Zfp219-201	<a href="#">ENSMUST00000067549.15</a>	2933	<a href="#">726aa</a>	Protein coding	<a href="#">CCDS36916</a>		TSL:1 , GENCODE basic , APPRIS P2 ,
Zfp219-204	<a href="#">ENSMUST00000226522.2</a>	2804	<a href="#">726aa</a>	Protein coding	<a href="#">CCDS36916</a>		GENCODE basic , APPRIS P2 ,
Zfp219-212	<a href="#">ENSMUST00000228580.2</a>	2812	<a href="#">681aa</a>	Protein coding	-		GENCODE basic , APPRIS ALT2 ,
Zfp219-208	<a href="#">ENSMUST00000226964.2</a>	787	<a href="#">201aa</a>	Protein coding	-		CDS 3' incomplete ,
Zfp219-213	<a href="#">ENSMUST00000228747.2</a>	719	<a href="#">141aa</a>	Protein coding	-		CDS 3' incomplete ,
Zfp219-205	<a href="#">ENSMUST00000226527.2</a>	657	<a href="#">131aa</a>	Protein coding	-		CDS 3' incomplete ,
Zfp219-207	<a href="#">ENSMUST00000226605.2</a>	630	<a href="#">109aa</a>	Protein coding	-		CDS 3' incomplete ,
Zfp219-206	<a href="#">ENSMUST00000226554.2</a>	495	<a href="#">10aa</a>	Protein coding	-		CDS 3' incomplete ,
Zfp219-211	<a href="#">ENSMUST00000228162.2</a>	480	<a href="#">14aa</a>	Protein coding	-		CDS 3' incomplete ,
Zfp219-210	<a href="#">ENSMUST00000228051.2</a>	431	<a href="#">95aa</a>	Protein coding	-		CDS 3' incomplete ,
Zfp219-209	<a href="#">ENSMUST00000227420.2</a>	465	No protein	Processed transcript	-		
Zfp219-203	<a href="#">ENSMUST00000226474.2</a>	747	No protein	Retained intron	-		

The strategy is based on the design of *Zfp219-202* transcript,the transcription is shown below:



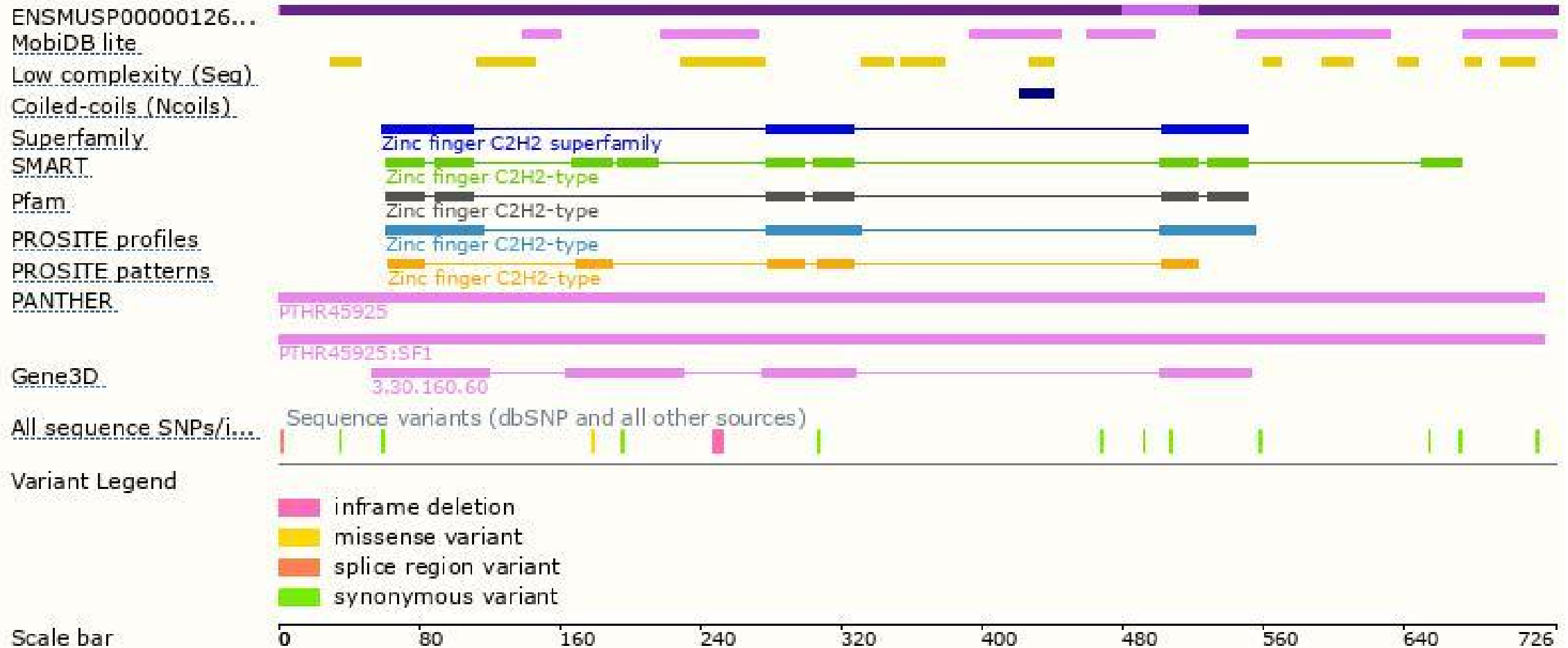


# Genomic location distribution





# Protein domain



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

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