

Zfp316 Cas9-KO Strategy

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Reviewer

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Project Overview

Project Name

Zfp316

Project type

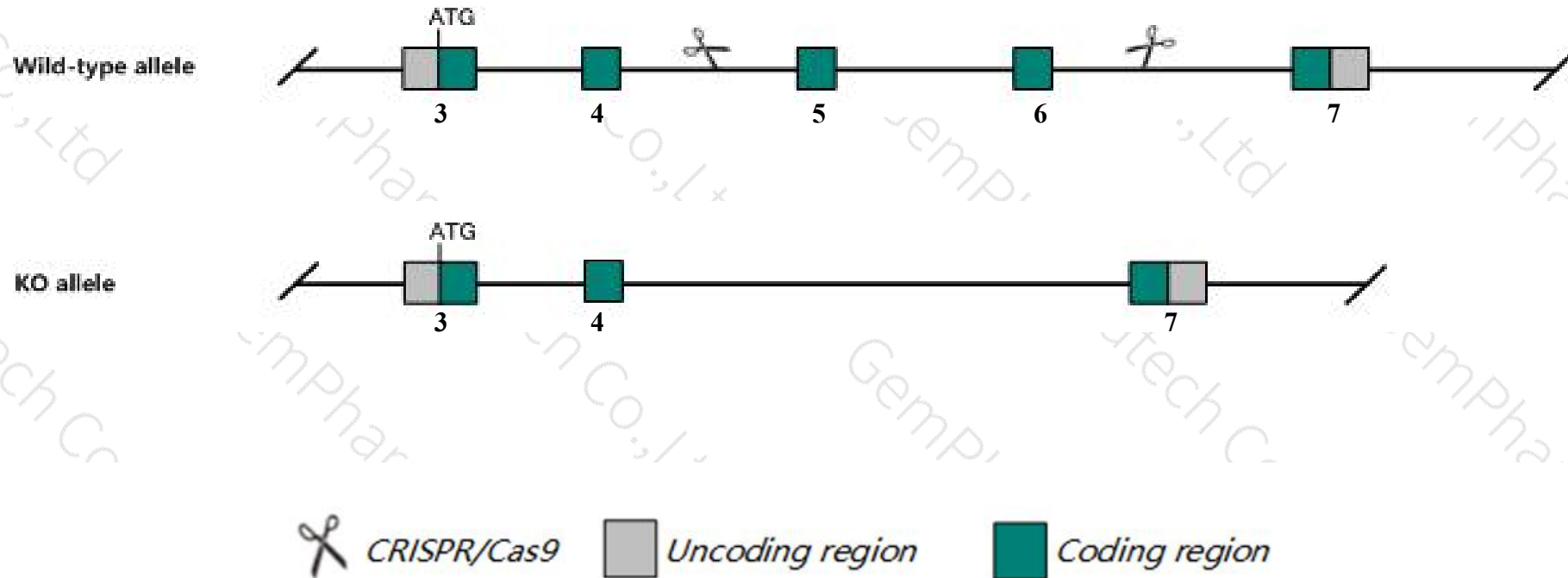
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Zfp316* gene. The schematic diagram is as follows:



- The *Zfp316* gene has 3 transcripts. According to the structure of *Zfp316* gene, exon5-exon6 of *Zfp316-202* (ENSMUST00000161448.7) transcript is recommended as the knockout region. The region contains 241bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Zfp316* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Zfp316* gene is located on the Chr5. 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.

Zfp316 zinc finger protein 316 [*Mus musculus* (house mouse)]

Gene ID: 54201, updated on 12-Aug-2019

Summary

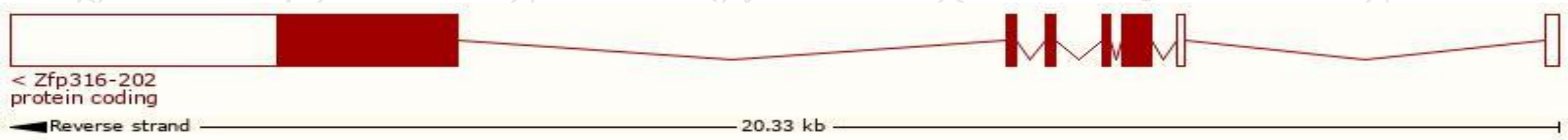
Official Symbol	Zfp316 provided by MGI
Official Full Name	zinc finger protein 316 provided by MGI
Primary source	MGI:MGI:1860402
See related	Ensembl:ENSMUSG00000046658
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Mzf3; Emzf1; Znf316
Expression	Ubiquitous expression in ovary adult (RPKM 8.9), adrenal adult (RPKM 7.2) and 24 other tissues See more
Orthologs	human all

Transcript information Ensembl

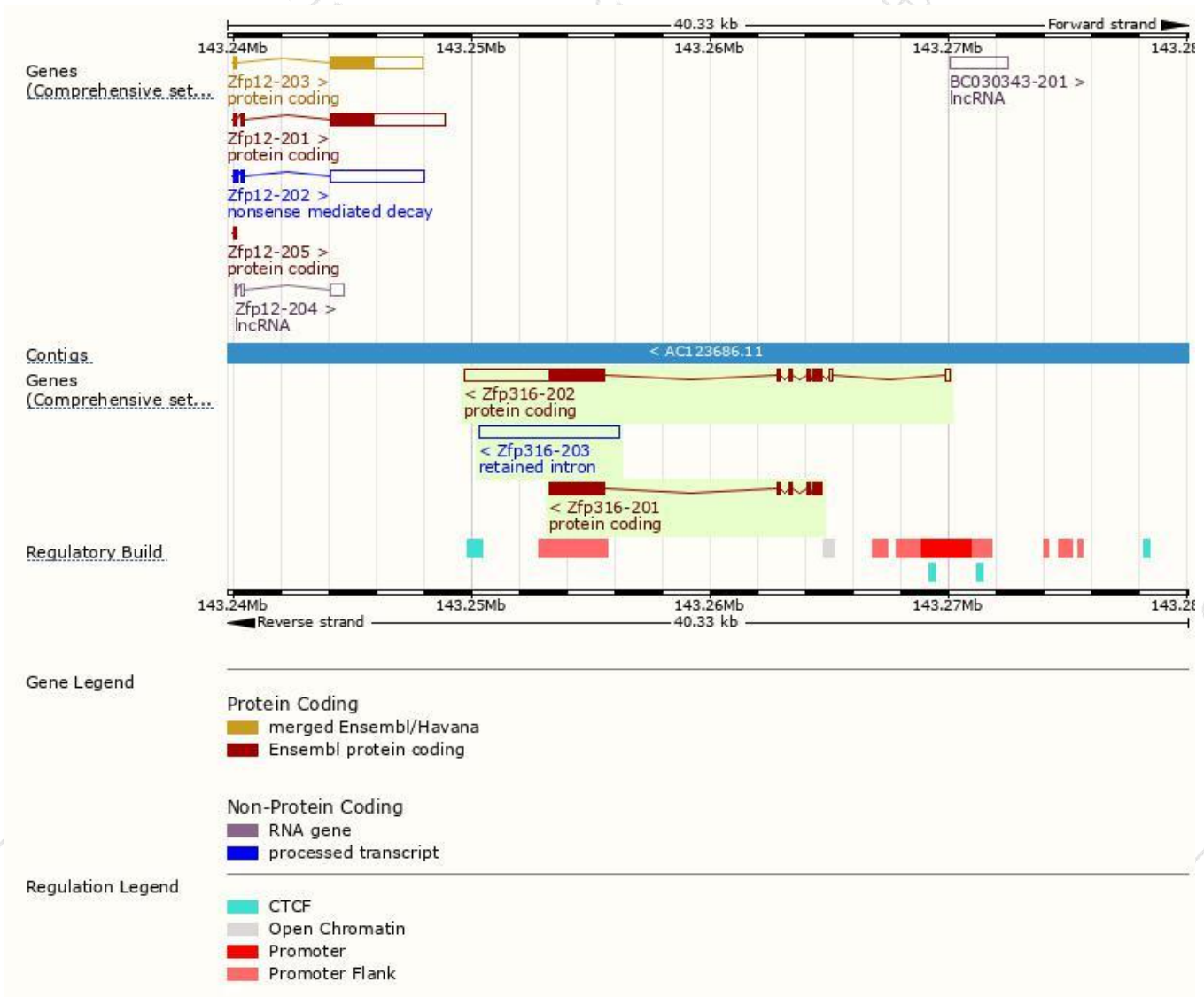
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Zfp316-202	ENSMUST00000161448.7	6891	1017aa	Protein coding	CCDS19838	G3XA34	TSL:1 GENCODE basic APPRIS P1
Zfp316-201	ENSMUST00000051665.8	3054	1017aa	Protein coding	CCDS19838	G3XA34	TSL:1 GENCODE basic APPRIS P1
Zfp316-203	ENSMUST00000197349.1	5912	No protein	Retained intron	-	-	TSL:NA

The strategy is based on the design of *Zfp316-202* transcript,The transcription is shown below



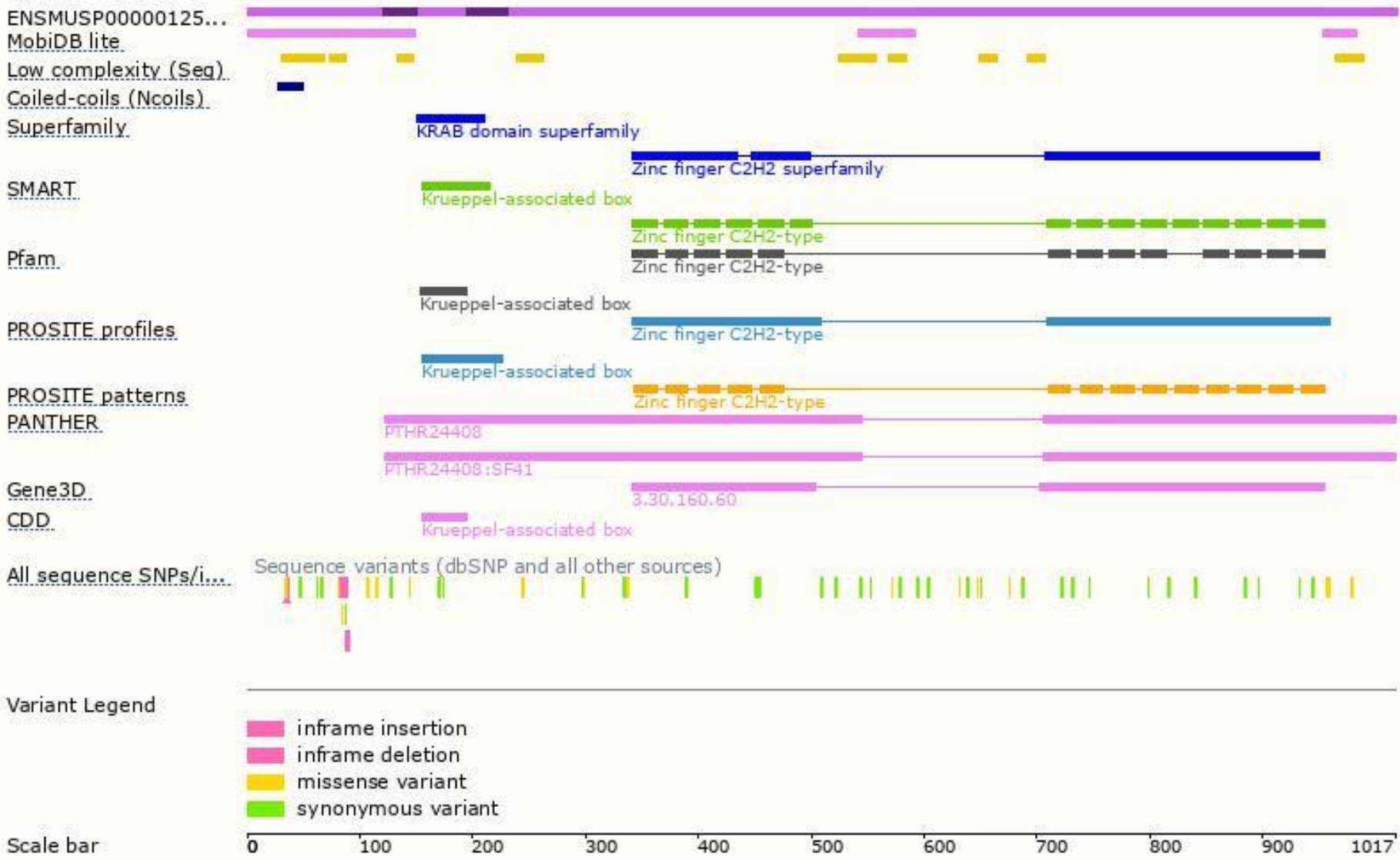
Genomic location distribution



Protein domain



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

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