

Zfp14 Cas9-KO Strategy

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Reviewer

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Design Date:

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Project Overview

Project Name

Zfp14

Project type

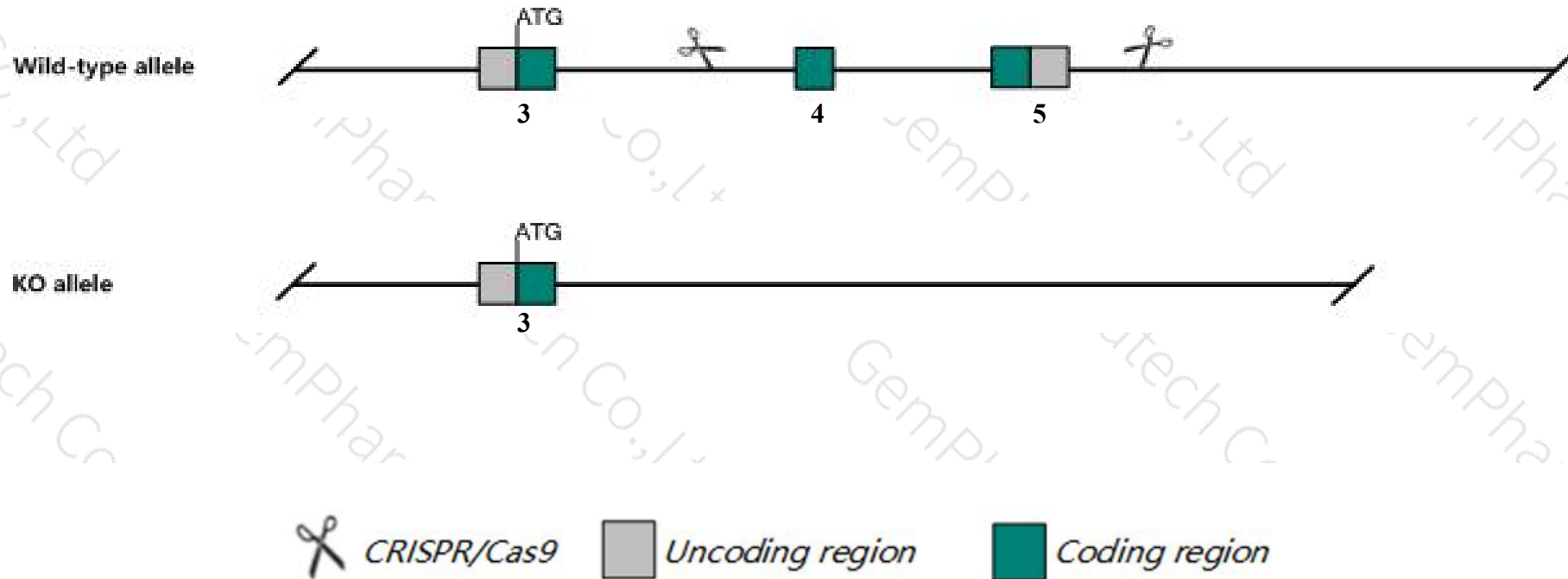
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Zfp14* gene. The schematic diagram is as follows:



- The *Zfp14* gene has 3 transcripts. According to the structure of *Zfp14* gene, exon4-exon5 of *Zfp14-203* (ENSMUST00000207873.1) transcript is recommended as the knockout region. The region contains most 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 *Zfp14* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Zfp14* gene is located on the Chr7. 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.

Zfp14 zinc finger protein 14 [*Mus musculus* (house mouse)]

Gene ID: 243906, updated on 24-Oct-2019

Summary

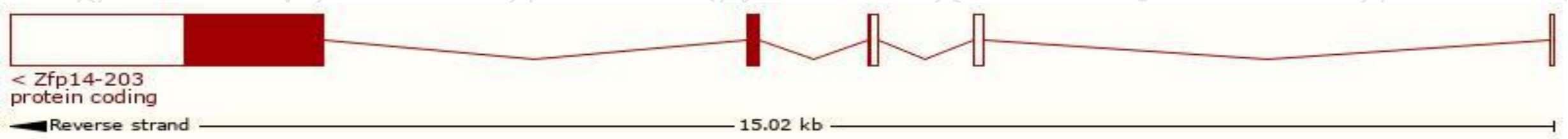
Official Symbol	Zfp14 provided by MGI
Official Full Name	zinc finger protein 14 provided by MGI
Primary source	MGI:MGI:99160
See related	Ensembl:ENSMUSG00000053985
Gene type	protein coding
RefSeq status	VALIDATED
Organism	<i>Mus musculus</i>
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Krox-9; Zfp-14; mKIAA1559; 4732429I09Rik
Expression	Broad expression in bladder adult (RPKM 3.1), CNS E18 (RPKM 2.8) and 26 other tissues See more
Orthologs	human all

Transcript information Ensembl

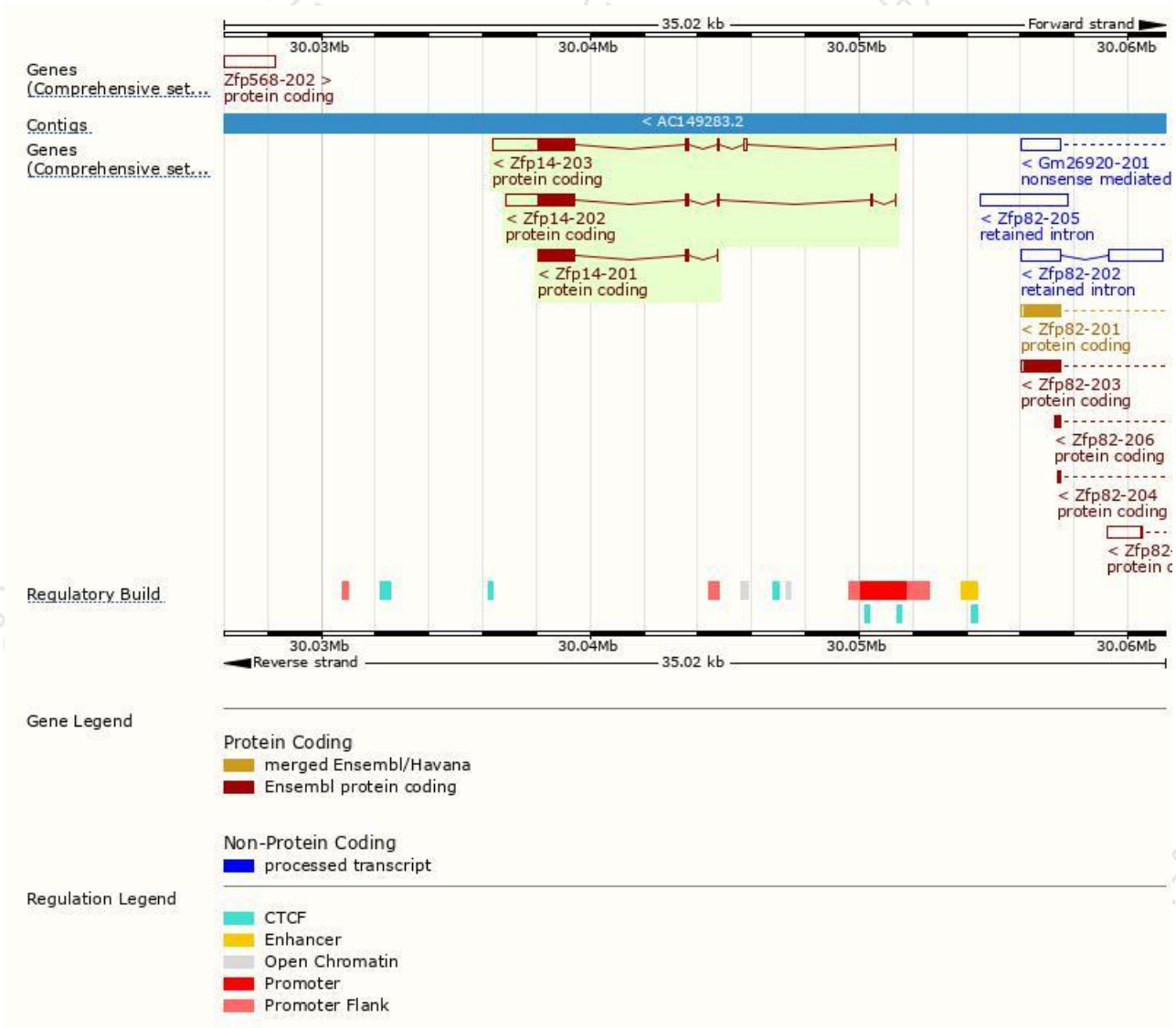
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Zfp14-203	ENSMUST00000207873.1	3411	501aa	Protein coding	CCDS39875	P10755	TSL:1 GENCODE basic APPRIS P1
Zfp14-202	ENSMUST00000207072.1	2835	501aa	Protein coding	CCDS39875	P10755	TSL:1 GENCODE basic APPRIS P1
Zfp14-201	ENSMUST00000077787.7	1506	501aa	Protein coding	CCDS39875	P10755	TSL:1 GENCODE basic APPRIS P1

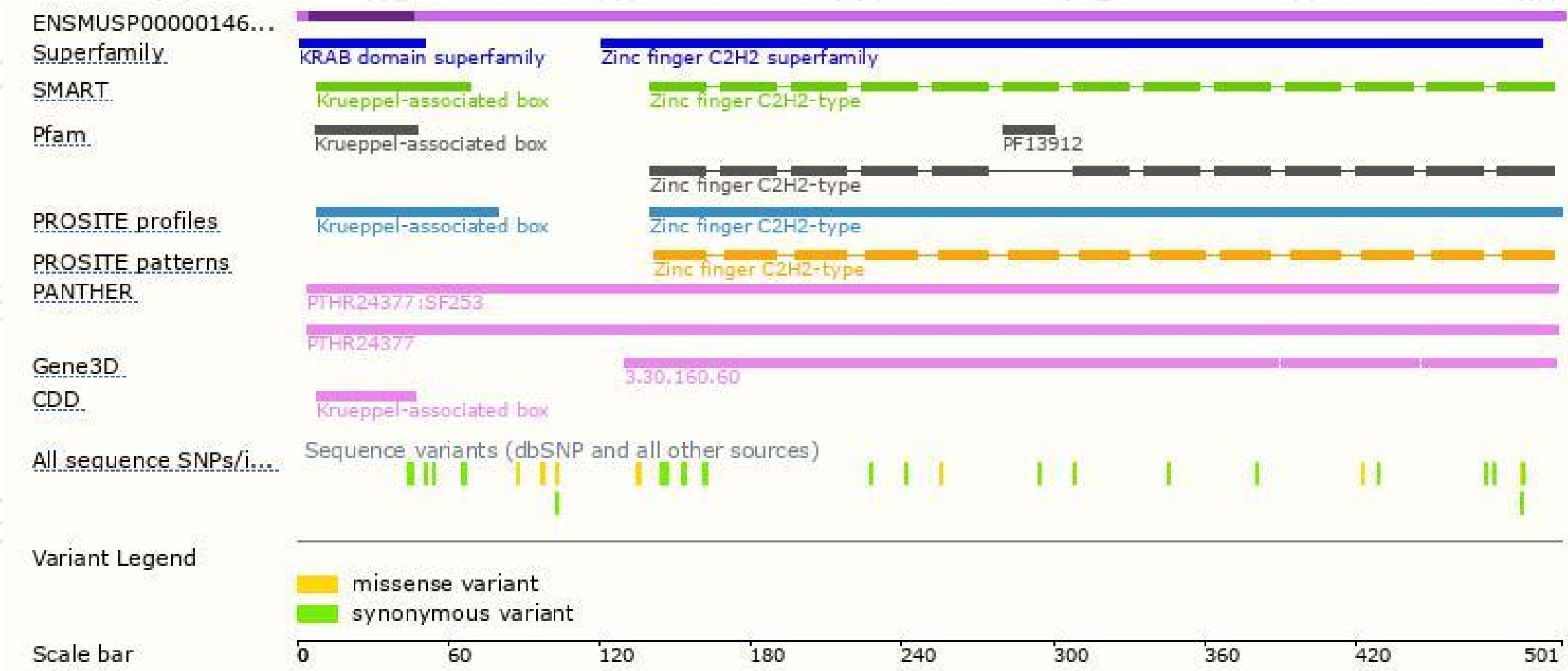
The strategy is based on the design of *Zfp14-203* transcript,The transcription is shown below



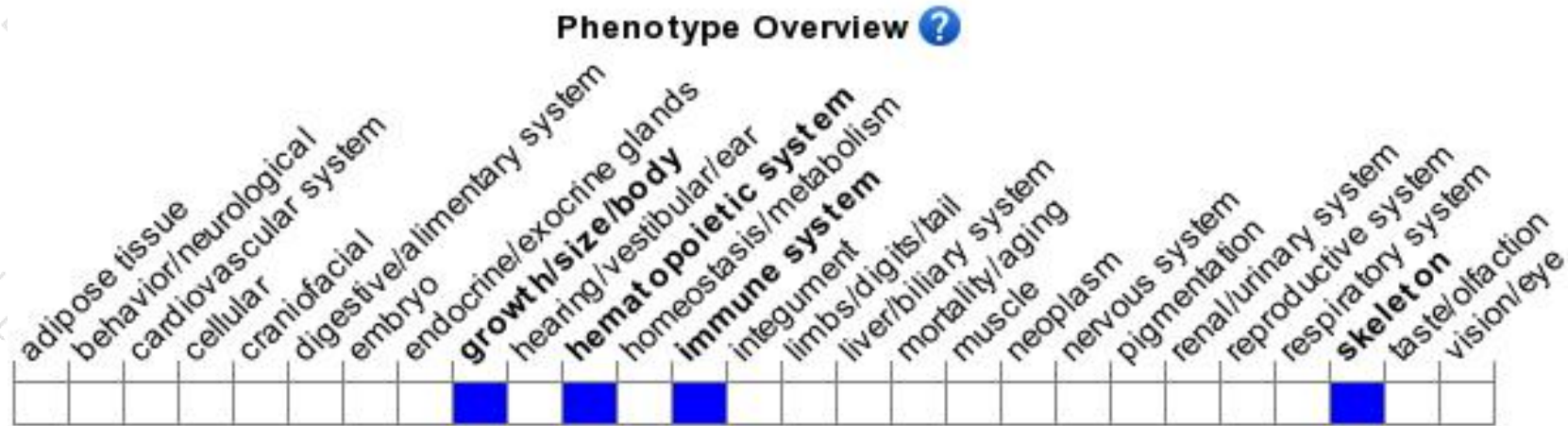
Genomic location distribution



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/>).

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

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