Zfp42 Cas9-CKO Strategy

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Design Date: 2019-8-4

Project Overview

Project Name	***Zfp42***
Project type	**Cas9-CKO**
Strain background	**C57BL/6JGpt**

Conditional Knockout strategy

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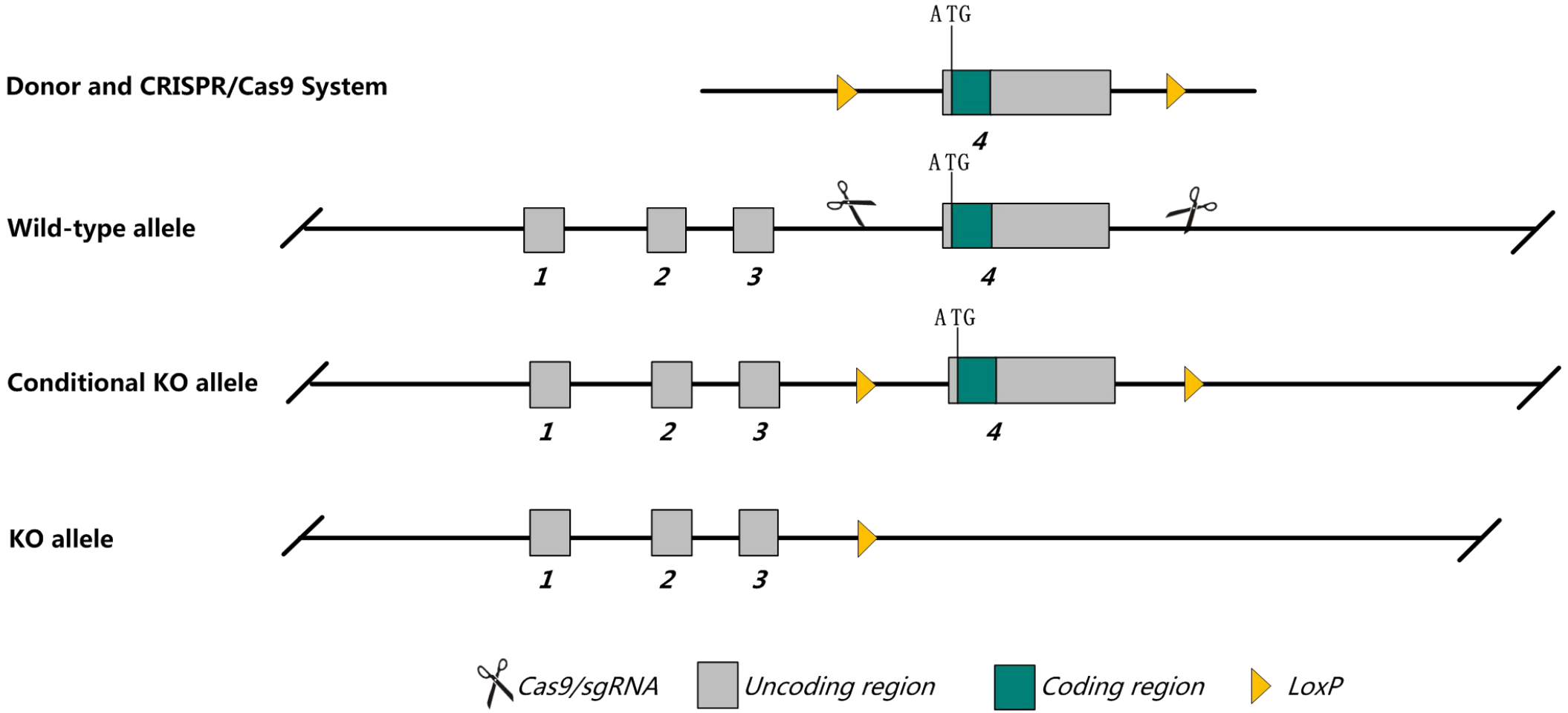
This model will use CRISPR/Cas9 technology to edit the *Zfp42* gene. The schematic diagram is as follows:



Technical routes

- The *Zfp42* gene has 3 transcripts. According to the structure of *Zfp42* gene, exon4 of *Zfp42-201* (ENSMUST00000082120.4) transcript is recommended as the knockout region.The region contains all 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 *Zfp42* gene. The brief process is as follows:CRISPR/Cas9 system and Donor 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 flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

Notice

- According to the existing MGI data,Mice homozygous for one knock-out allele exhibit premature age-related male germ cell loss, abnormal sperm morphology, and mild testicular atrophy.

- The *Zfp42* gene is located on the Chr8. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information （NCBI）

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Zfp42 zinc finger protein 42 [Mus musculus (house mouse)]

Gene ID: 22702, updated on 31-Jan-2019

Summary

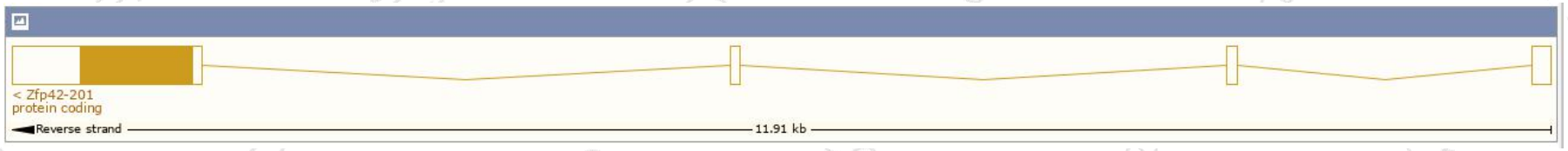
Official Symbol	Zfp42 provided by MGI
Official Full Name	zinc finger protein 42 provided byMGI
Primary source	MGI:MGI:99187
See related	Ensembl:ENSMUSG00000051176
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Rex-1, Rex1, Zfp-42
Expression	Restricted expression toward placenta adult (RPKM 4.2)See more
Orthologs	human all

Transcript information（Ensembl）

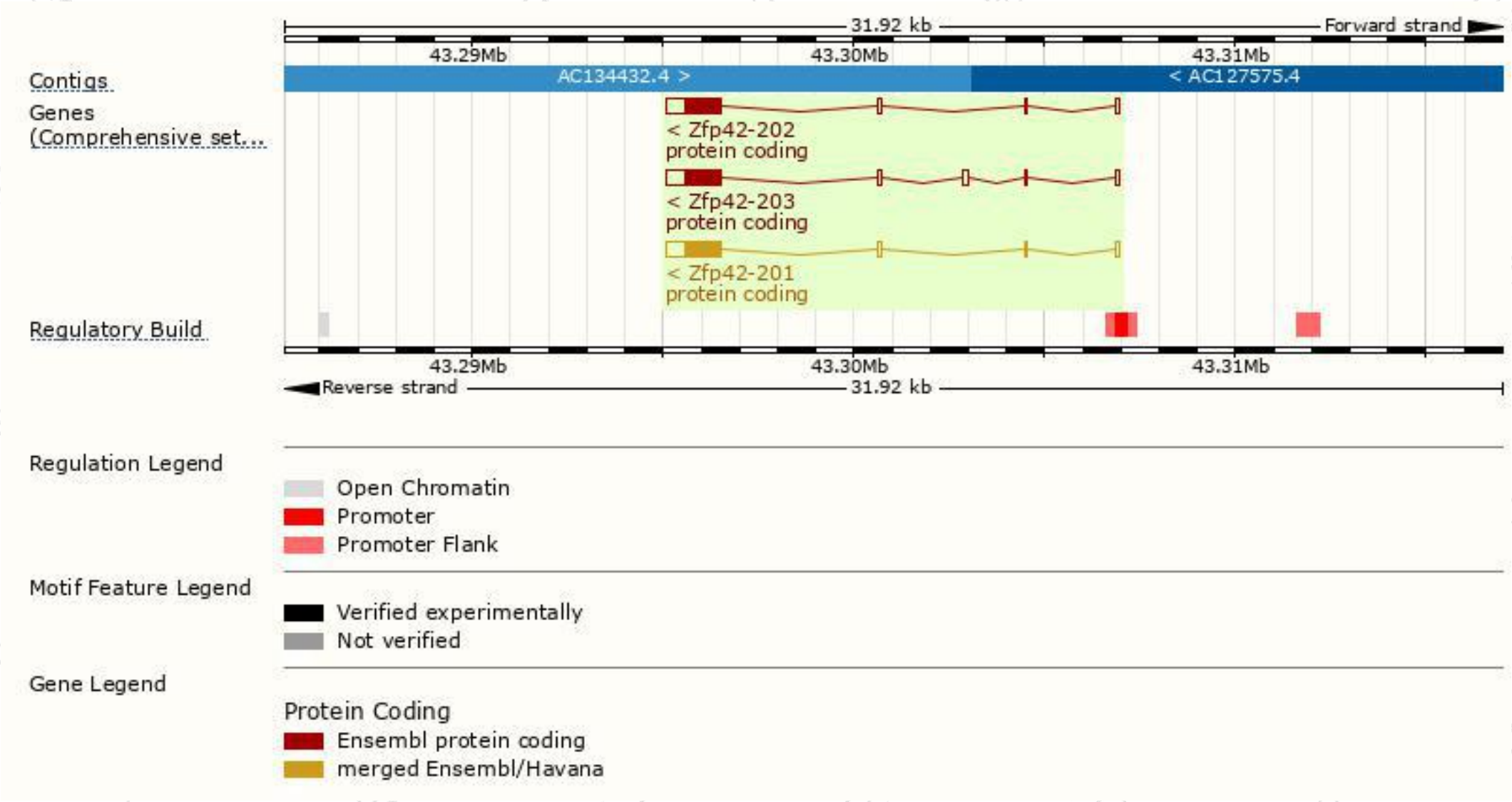
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Zfp42-203	ENSMUST00000211248.1	1837	288aa	Protein coding	CCDS22267	E9QK22	TSL:1 GENCODE basic APPRIS P1
Zfp42-201	ENSMUST00000082120.4	1772	288aa	Protein coding	CCDS22267	E9QK22	TSL:1 GENCODE basic APPRIS P1
Zfp42-202	ENSMUST00000209356.1	1744	288aa	Protein coding	CCDS22267	E9QK22	TSL:1 GENCODE basic APPRIS P1

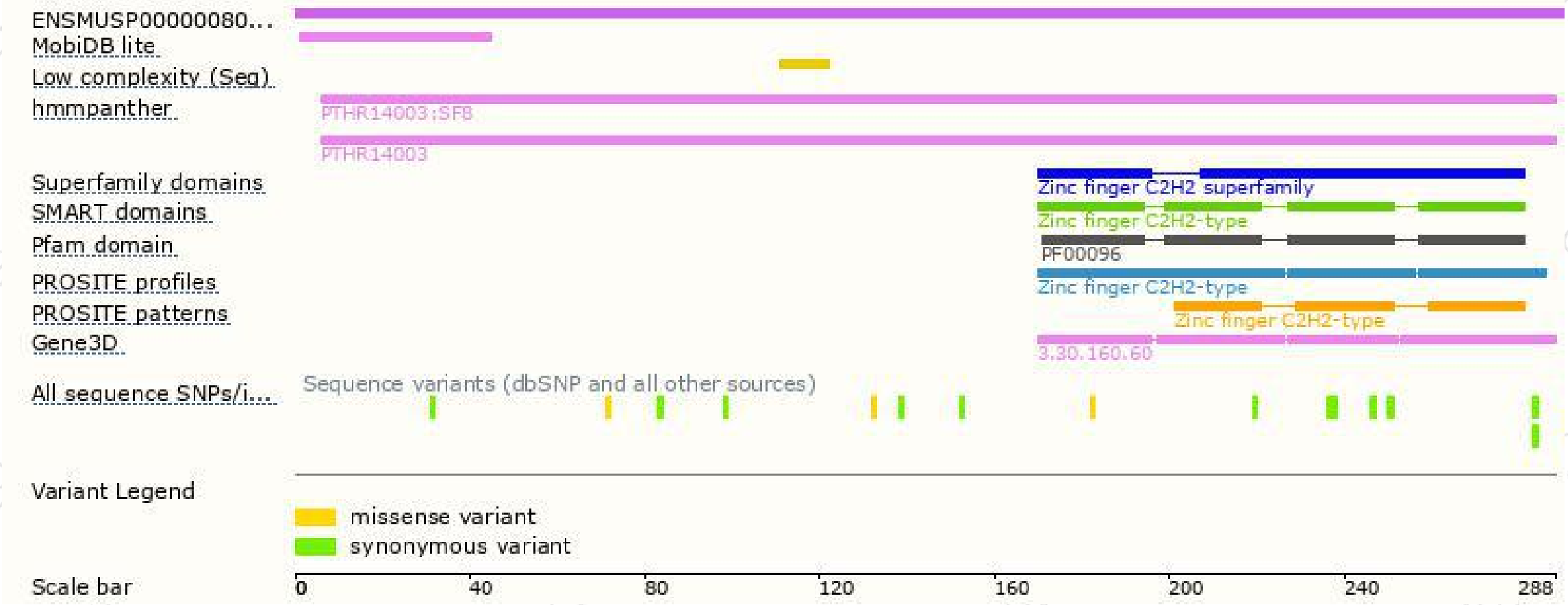
The strategy is based on the design of *Zfp42-201* transcript,The transcription is shown below



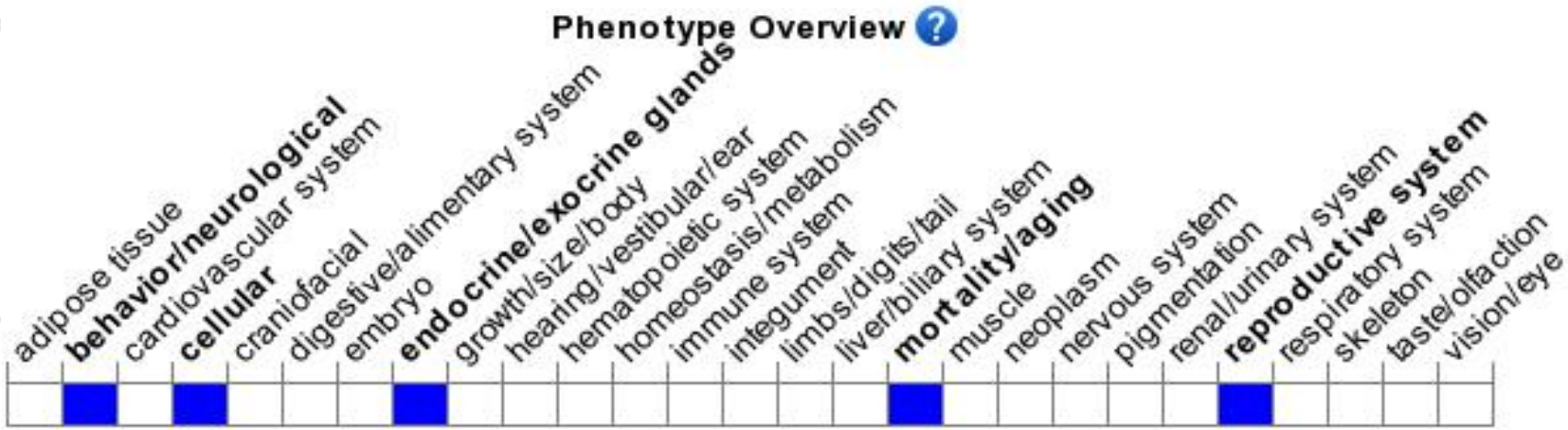
Genomic location distribution



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 one knock-out allele exhibit premature age-related male germ cell loss, abnormal sperm morphology, and mild testicular atrophy.

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

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