

Zp1 Cas9-KO Strategy

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Project Overview

Project Name

Zp1

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

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



- The *Zp1* gene has 2 transcripts. According to the structure of *Zp1* gene, exon2-exon9 of *Zp1-201* (ENSMUST00000025641.1) transcript is recommended as the knockout region. The region contains 1379bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Zp1* gene. The brief process is as follows: CRISPR/Cas9 system w

- According to the existing MGI data, Female homozygous mutants produce oocytes with abnormal zona pellucida. Fecundity is significantly reduced, probably due to precocious hatching.
- The *Zp1* gene is located on the Chr19. 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.

Gene information (NCBI)

Zp1 zona pellucida glycoprotein 1 [Mus musculus (house mouse)]

Gene ID: 22786, updated on 31-Jan-2019

Summary



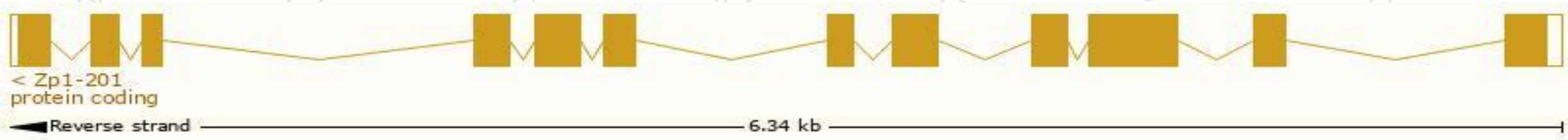
Official Symbol	Zp1 provided by MGI
Official Full Name	zona pellucida glycoprotein 1 provided by MGI
Primary source	MGI:MGI:103073
See related	Ensembl:ENSMUSG000000024734
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Expression	Restricted expression toward ovary adult (RPKM 22.1) See more
Orthologs	human all

Transcript information (Ensembl)

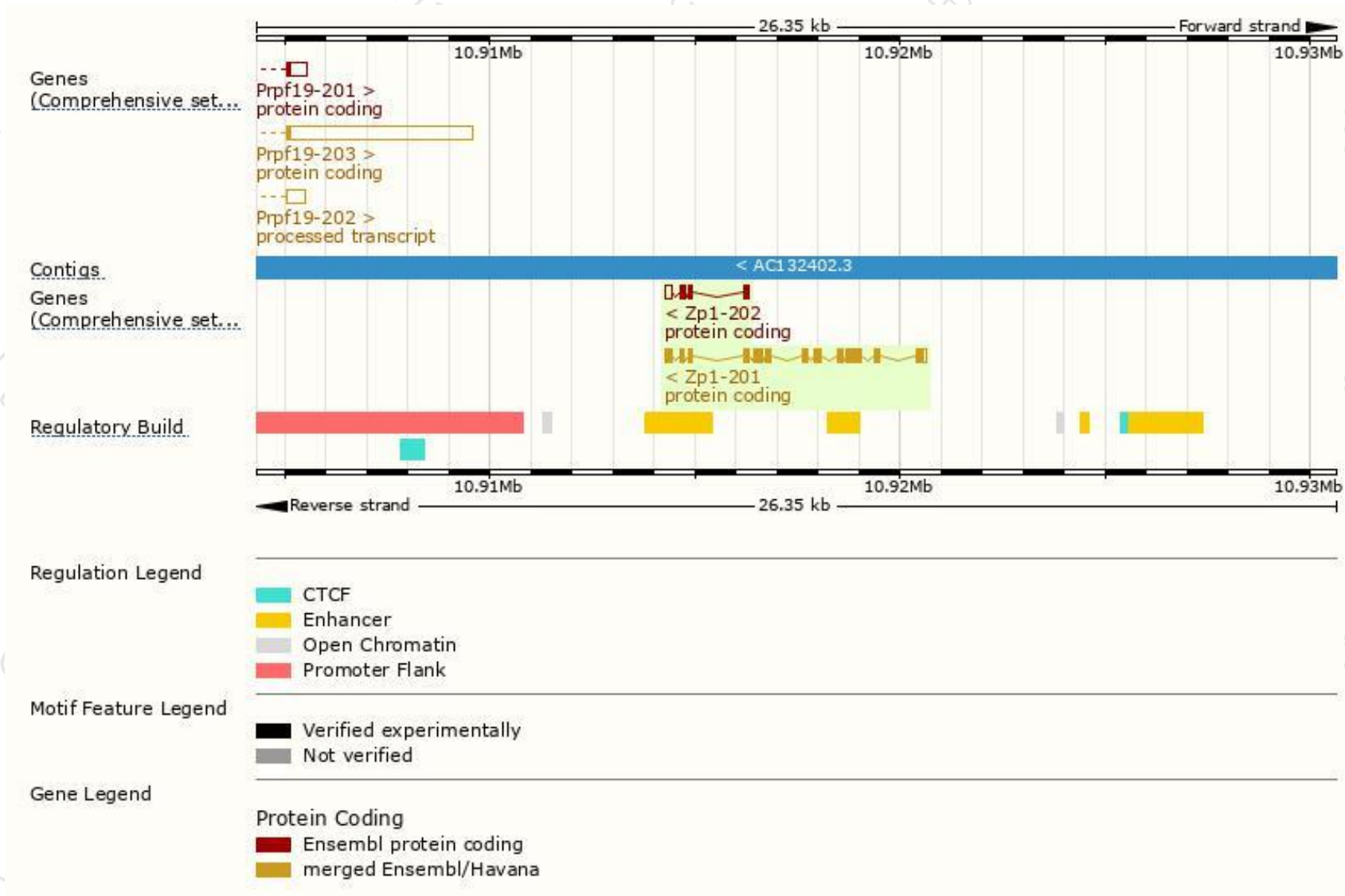
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Zp1-201	ENSMUST00000025641.1	1963	623aa	Protein coding	CCDS37918	Q059L1 Q62005	TSL:1 GENCODE basic APPRIS P1
Zp1-202	ENSMUST00000168708.7	481	95aa	Protein coding	-	F7CW32	CDS 5' incomplete TSL:3

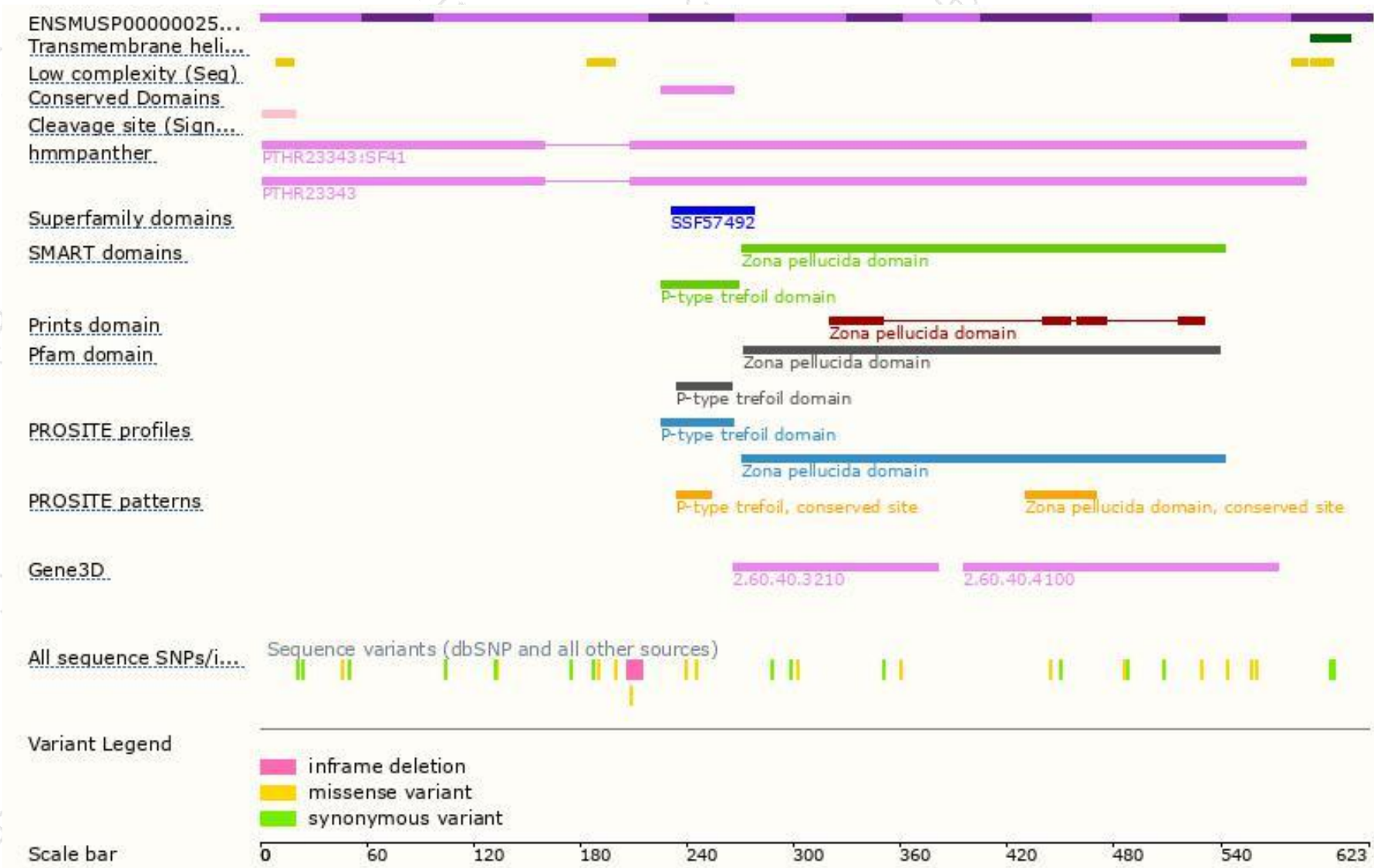
The strategy is based on the design of *Zp1-201* transcript,The transcription is shown below



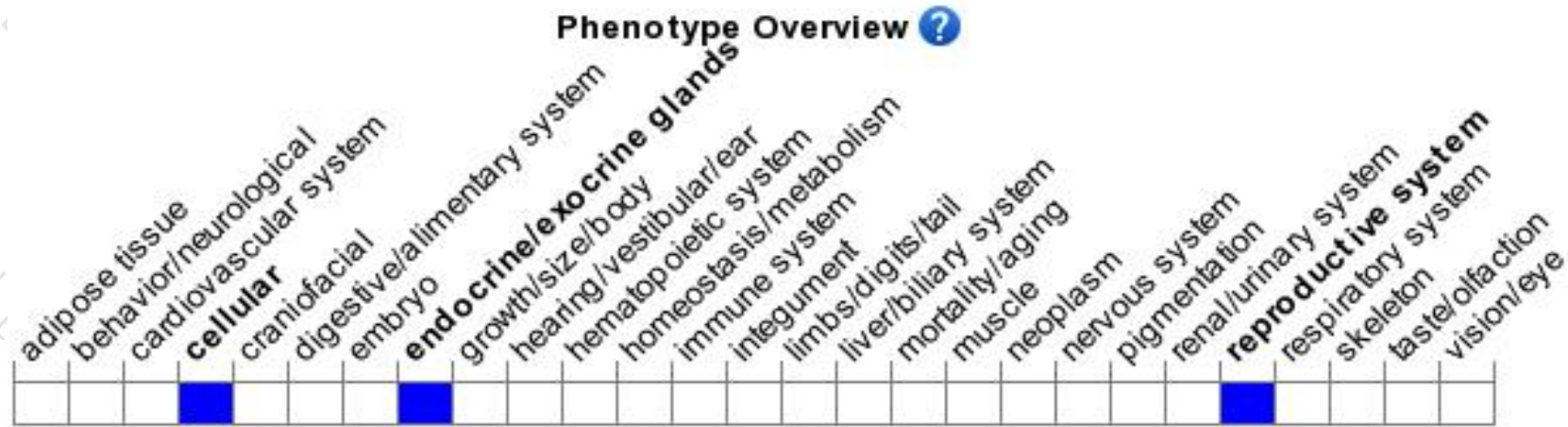
Genomic location distribution



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, Female homozygous mutants produce oocytes with abnormal zona pellucida. Fecundity is significantly reduced, probably due to precocious hatching.

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

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