

# *Jade2* Cas9-KO Strategy

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# Project Overview

**Project Name**

*Jade2*

**Project type**

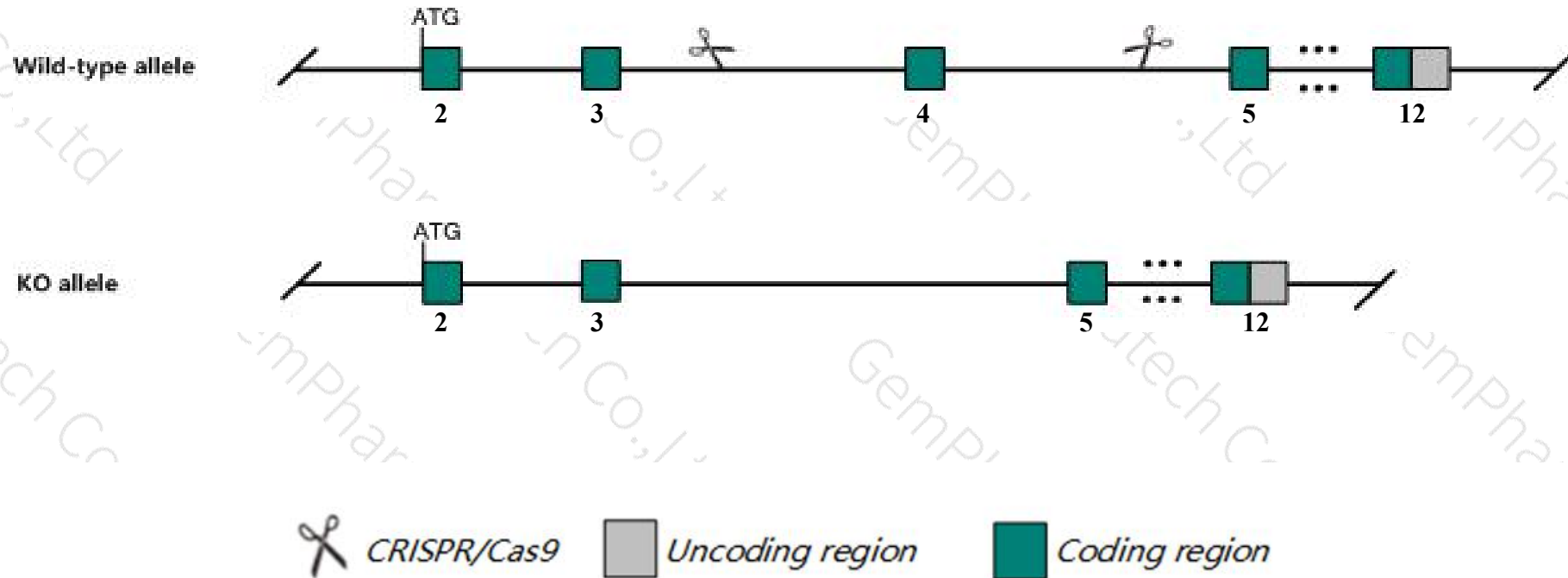
**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Jade2* gene. The schematic diagram is as follows:



- The *Jade2* gene has 4 transcripts. According to the structure of *Jade2* gene, exon4 of *Jade2-201* (ENSMUST00000020655.13) transcript is recommended as the knockout region. The region contains 158bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Jade2* gene. The brief process is as follows: CRISPR/Cas9 system

- Transcript 204 may not be affected.
- The *Jade2* gene is located on the Chr11. 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)

## Jade2 jade family PHD finger 2 [ *Mus musculus* (house mouse) ]

Gene ID: 76901, updated on 12-Aug-2019

### Summary

Official Symbol	Jade2 provided by <a href="#">MGI</a>
Official Full Name	jade family PHD finger 2 provided by <a href="#">MGI</a>
Primary source	<a href="#">MGI:MGI:1924151</a>
See related	<a href="#">Ensembl:ENSMUSG00000020387</a>
Gene type	protein coding
RefSeq status	VALIDATED
Organism	<a href="#">Mus musculus</a>
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Phf15; AI480685; mKIAA0239; 1200017K05Rik
Expression	Broad expression in thymus adult (RPKM 14.2), lung adult (RPKM 11.2) and 20 other tissues <a href="#">See more</a>
Orthologs	<a href="#">human</a> <a href="#">all</a>

### Genomic context

Location: 11; 11 B1.3

Exon count: 17

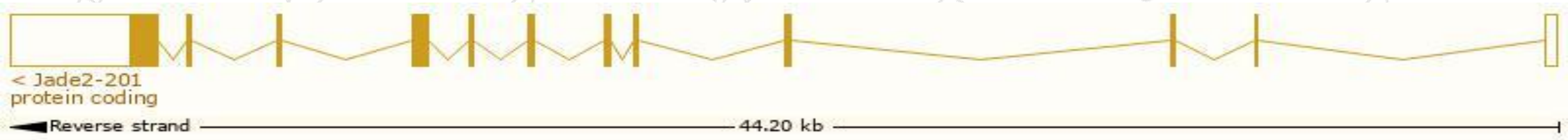
See Jade2 in [Genome Data Viewer](#)

# Transcript information (Ensembl)

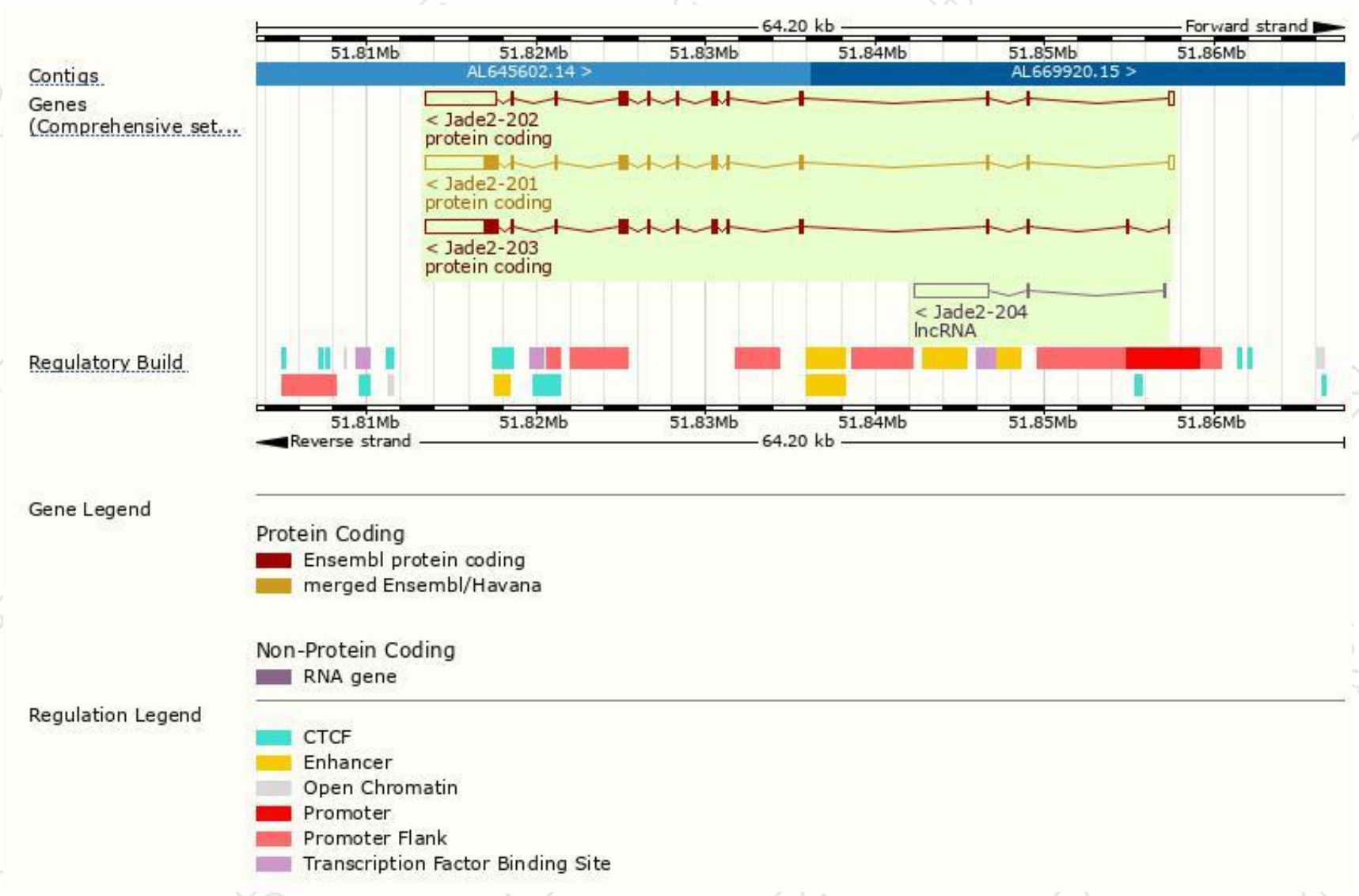
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Jade2-201	<a href="#">ENSMUST00000020655.13</a>	6273	<a href="#">829aa</a>	Protein coding	<a href="#">CCDS24662</a>	<a href="#">Q6ZQF7</a>	TSL:1 GENCODE basic APPRIS P2
Jade2-203	<a href="#">ENSMUST00000109091.1</a>	6099	<a href="#">829aa</a>	Protein coding	<a href="#">CCDS24662</a>	<a href="#">Q6ZQF7</a>	TSL:1 GENCODE basic APPRIS P2
Jade2-202	<a href="#">ENSMUST00000109090.7</a>	6230	<a href="#">576aa</a>	Protein coding	-	<a href="#">A2AA59</a>	TSL:1 GENCODE basic APPRIS ALT2
Jade2-204	<a href="#">ENSMUST00000151277.1</a>	4552	No protein	lncRNA	-	-	TSL:1

The strategy is based on the design of *Jade2-201* transcript,The transcription is shown below

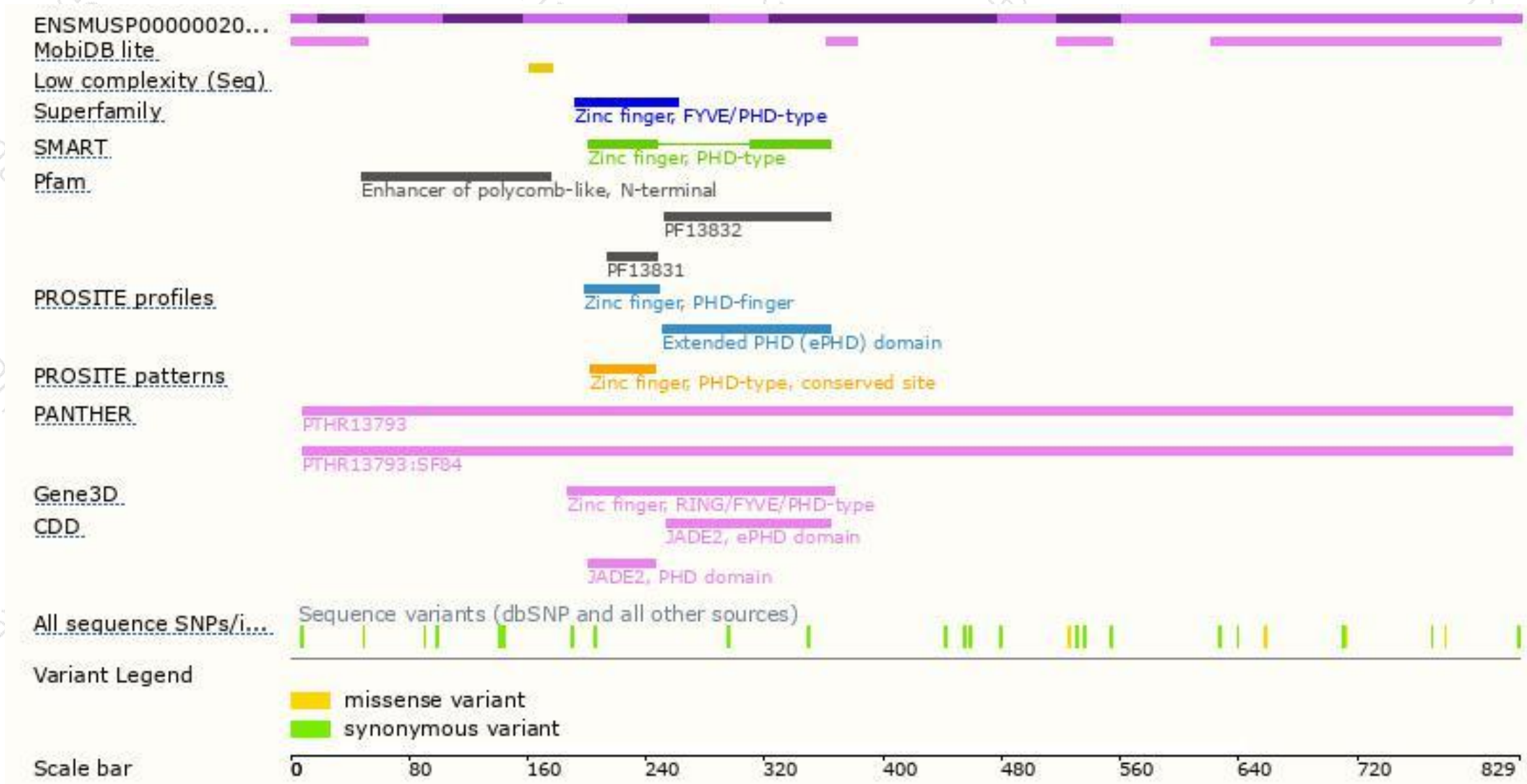


# Genomic location distribution





# Protein domain



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

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