

# *Creb3l3* Cas9-KO Strategy

Designer: Qiong Zhou

# Project Overview

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**Project Name**

*Creb3l3*

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**Project type**

**Cas9-KO**

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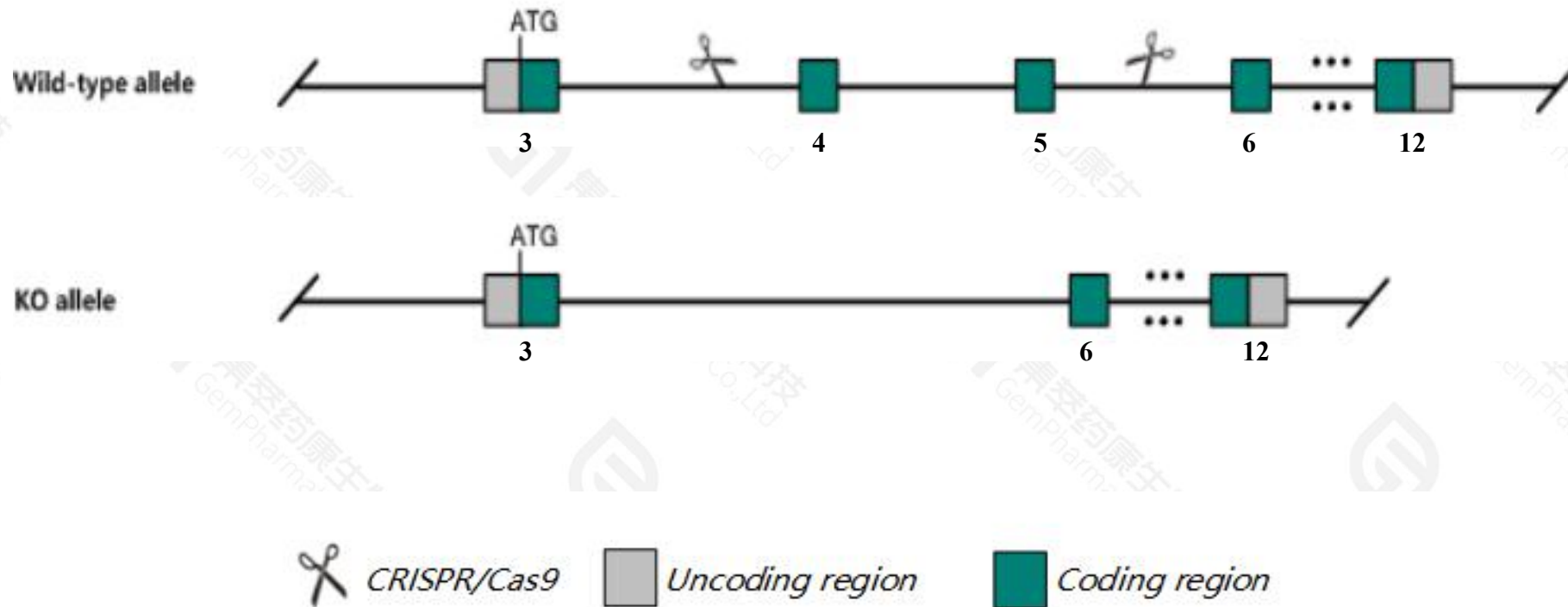
**Strain background**

**C57BL/6JGpt**

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# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Creb3l3* gene. The schematic diagram is as follows:



# Technical routes

- The *Creb3l3* gene has 4 transcripts. According to the structure of *Creb3l3* gene, exon4-exon5 of *Creb3l3-201*(ENSMUST00000117422.2) transcript is recommended as the knockout region. The region contains 430bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Creb3l3* gene. The brief process is as follows: CRISPR/Cas9 system 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.

- According to the existing MGI data, mice homozygous for a knock-out allele are viable and healthy but exhibit reduced expression of acute phase response proteins following treatment with tunicamycin that induces ER stress. Mice homozygous for a different knock-out allele exhibit resistance to sulpyrine-induced shock.
- Transcript *Creb3l3-202* may not be affected.
- The *Creb3l3* gene is located on the Chr10. 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.



## Creb3l3 cAMP responsive element binding protein 3-like 3 [Mus musculus (house mouse)]

Gene ID: 208677, updated on 1-Mar-2021

### Summary



**Official Symbol** Creb3l3 provided by [MGI](#)

**Official Full Name** cAMP responsive element binding protein 3-like 3 provided by [MGI](#)

**Primary source** [MGI:MGI:2384786](#)

**See related** [Ensembl:ENSMUSG00000035041](#)

**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** BC010786, CREB-, CREB-H, D10Bur1, D10Bur1e

**Expression** Biased expression in duodenum adult (RPKM 141.6), small intestine adult (RPKM 102.5) and 6 other tissues [See more](#)

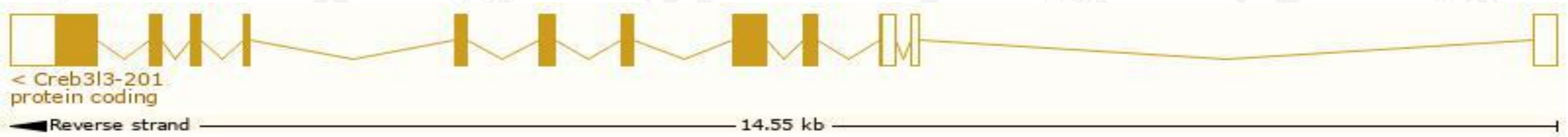
**Orthologs** [human](#) [all](#)

# Transcript information (Ensembl)

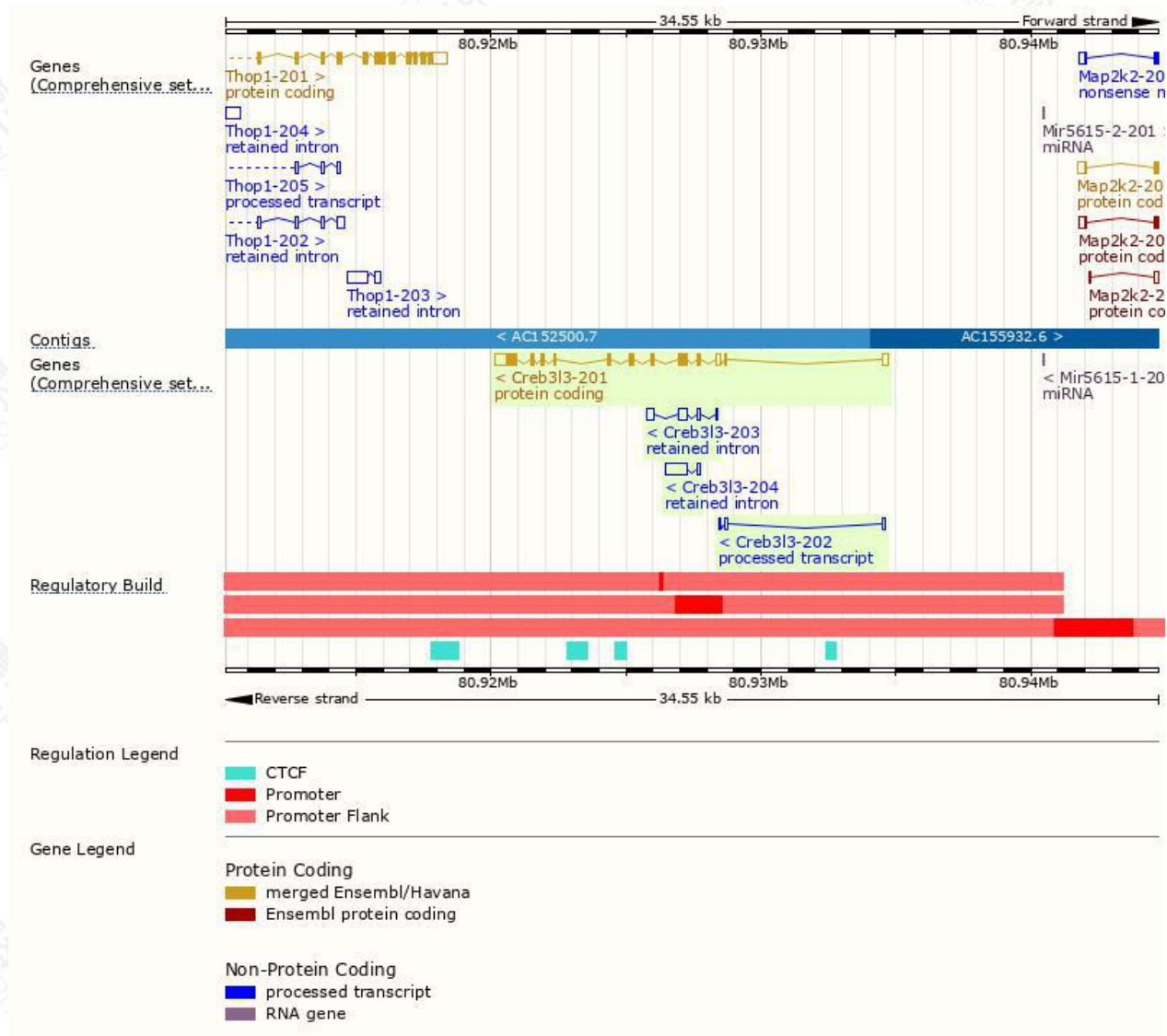
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Creb3l3-201	<a href="#">ENSMUST00000117422.2</a>	2294	<a href="#">479aa</a>	Protein coding	<a href="#">CCDS24043</a>		TSL:1 , GENCODE basic , APPRIS P1 ,
Creb3l3-202	<a href="#">ENSMUST00000125907.2</a>	275	No protein	Processed transcript	-		TSL:5 ,
Creb3l3-204	<a href="#">ENSMUST00000143969.2</a>	922	No protein	Retained intron	-		TSL:3 ,
Creb3l3-203	<a href="#">ENSMUST00000127374.2</a>	744	No protein	Retained intron	-		TSL:2 ,

The strategy is based on the design of *Creb3l3-201* transcript,the transcription is shown below:

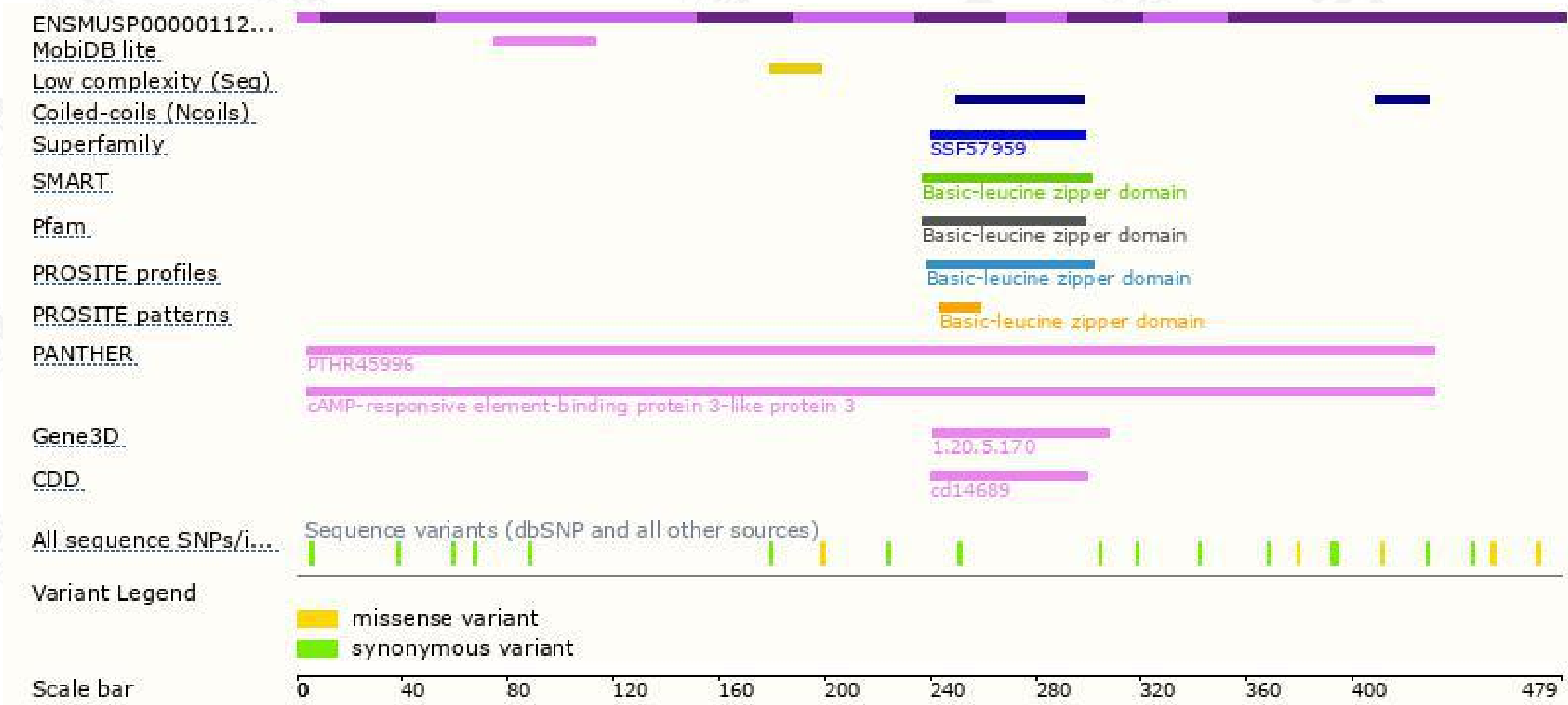


# Genomic location distribution

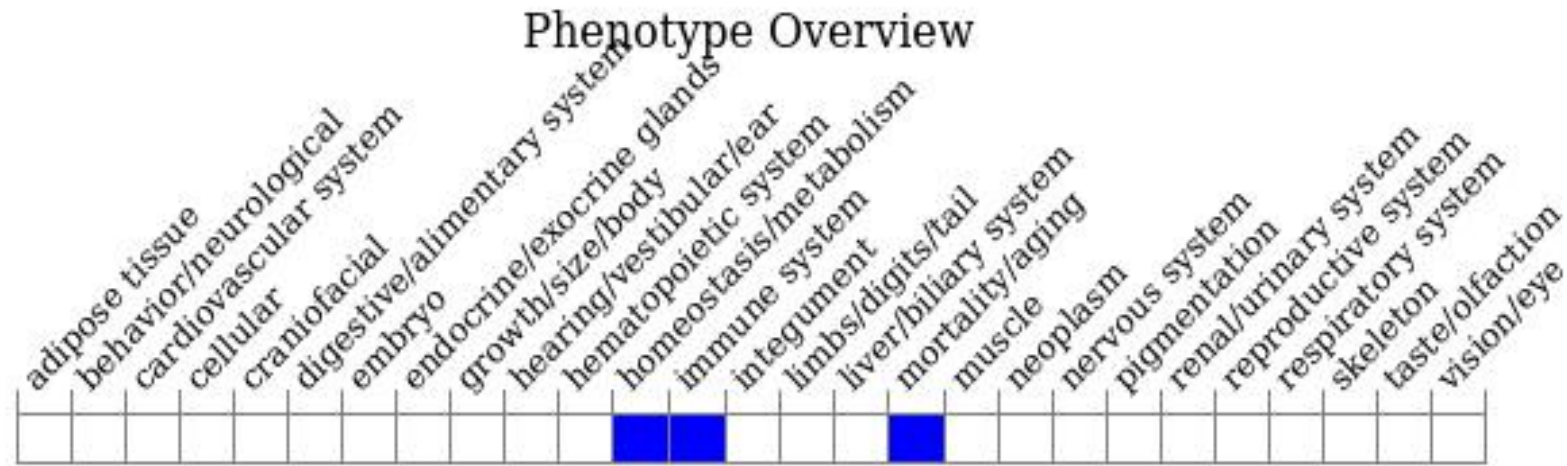




# Protein domain



# 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 a knock-out allele are viable and healthy but exhibit reduced expression of acute phase response proteins following treatment with tunicamycin that induces ER stress. Mice homozygous for a different knock-out allele exhibit resistance to sulpyrine-induced shock.

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