

Creb314 Cas9-KO Strategy

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Project Overview

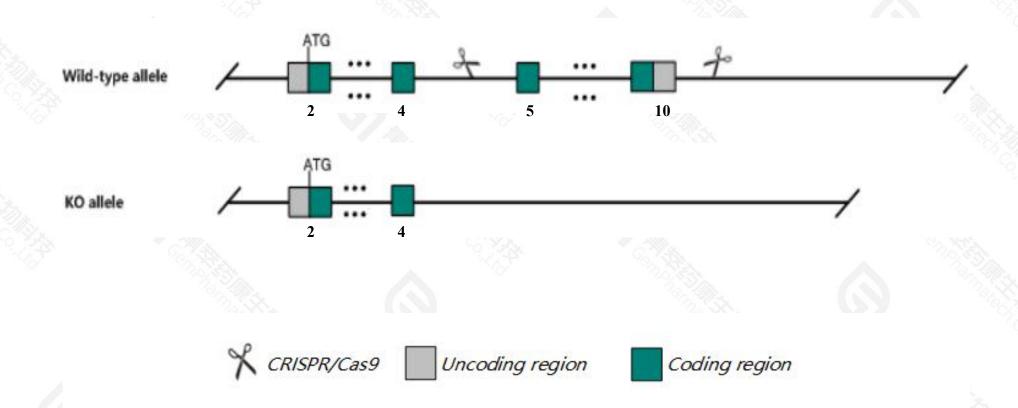


Project Name	Creb3l4			
Project type	Cas9-KO			
Strain background	C57BL/6JGpt			

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Creb314 gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Creb3l4* gene has 2 transcripts. According to the structure of *Creb3l4* gene, exon5-exon10 of *Creb3l4*-201(ENSMUST00000029547.10) transcript is recommended as the knockout region. The region contains translation stop codon sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Creb314* 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.

Notice



- > According to the existing MGI data, homozygous null mice display oligozoospermia but have normal fertility and sperm morphology and motility.
- ➤ The *Creb314* gene is located on the Chr3. 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)

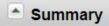


Creb3l4 cAMP responsive element binding protein 3-like 4 [Mus musculus (house mouse)]

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Gene ID: 78284, updated on 16-Oct-2021



Official Symbol Creb3l4 provided by MGI

Official Full Name cAMP responsive element binding protein 3-like 4 provided by MGI

Primary source MGI:MGI:1916603

See related Ensembl: ENSMUSG00000027938

Gene type protein coding
RefSeq status REVIEWED
Organism Mus musculus

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae;

Murinae; Mus; Mus

Also known as JAL; CREB4; acre1; AIBZIP; TISP40; AV040530; AV258827; 1700012K17Rik; 5330432F22Rik

Summary This gene encodes a CREB (cyclic AMP-responsive element-binding) protein with a transmembrane domain which localizes it to the ER membrane. The

encoded protein may play a role in adiposity and male germ cell development. Homozygous knockout mice for this gene show increased adipogenesis, elevated testicular germ cell apoptosis and defects in spermatogenesis. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Apr

2015]

Expression Biased expression in testis adult (RPKM 127.4), colon adult (RPKM 11.4) and 1 other tissue See more

Orthologs human all

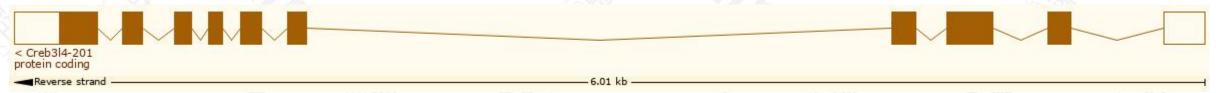
Transcript information (Ensembl)



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

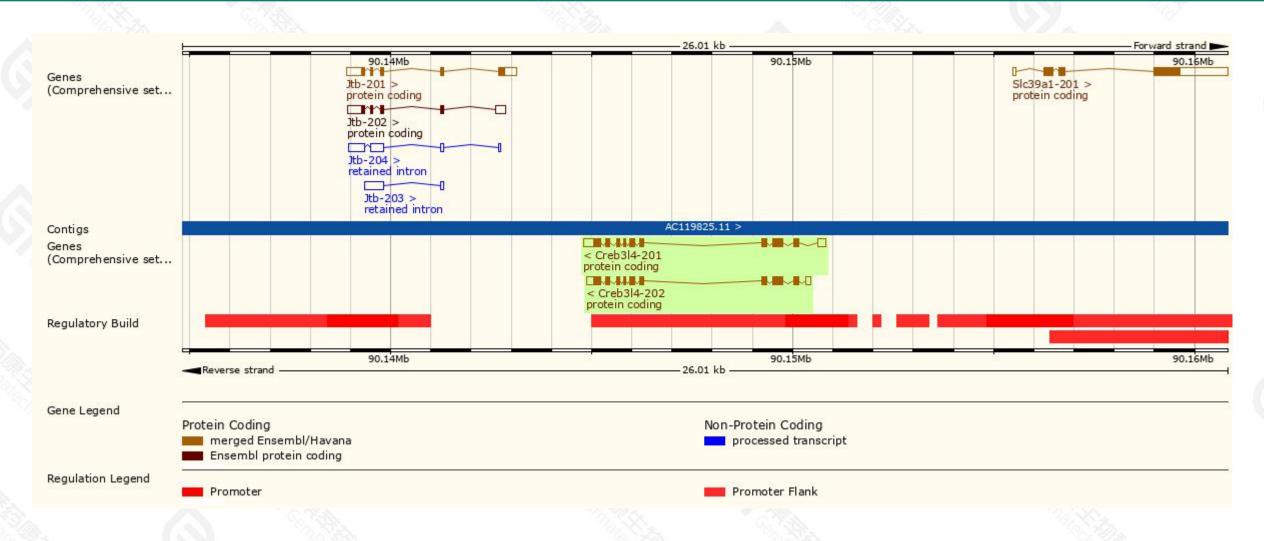
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Creb3l4-201	ENSMUST00000029547.10	1553	370aa	Protein coding	CCDS17524		TSL:1, GENCODE basic, APPRIS P1,
Creb3l4-202	ENSMUST00000107369.2	1404	370aa	Protein coding	CCDS17524		TSL:1, GENCODE basic, APPRIS P1,

The strategy is based on the design of *Creb314-201* transcript, the transcription is shown below:



Genomic location distribution





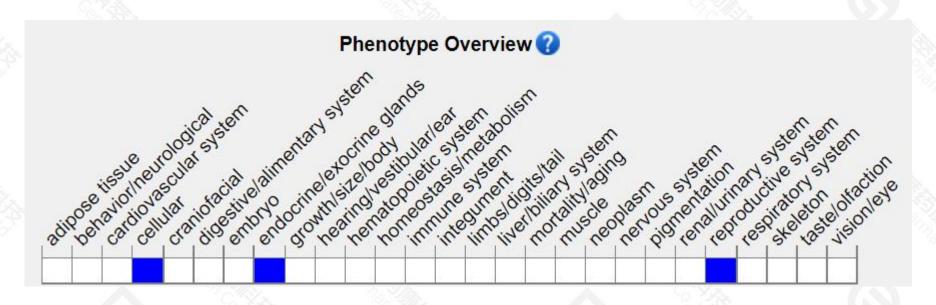
Protein domain





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, homozygous null mice display oligozoospermia but have normal fertility and sperm morphology and motility.



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

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