Champ1 Cas9-KO Strategy

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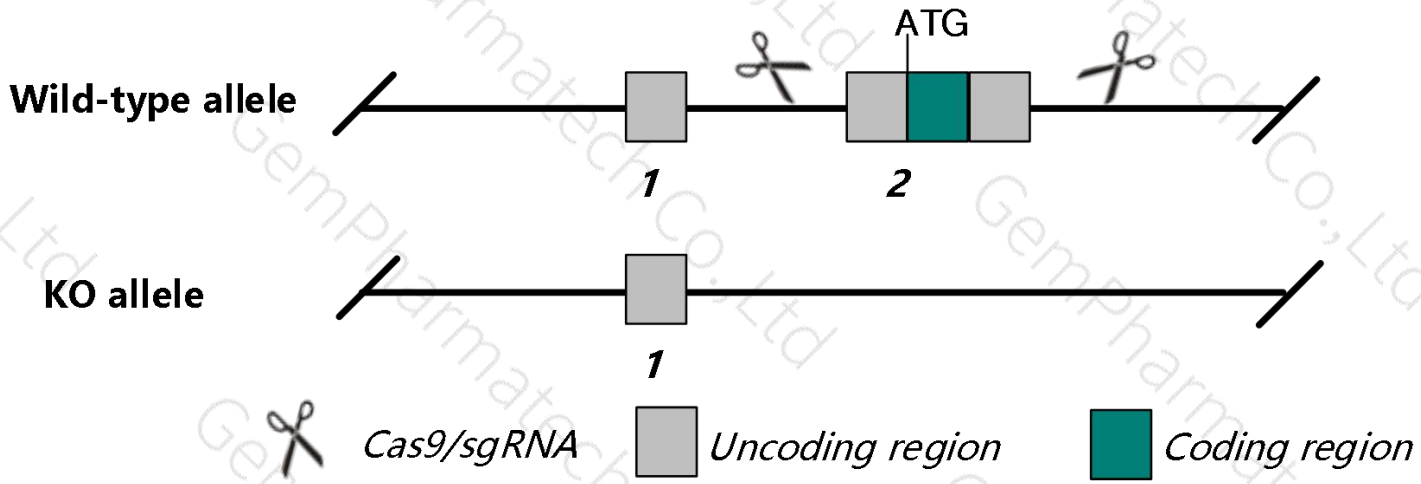
Project Overview

Project Name	***Champ1***
Project type	**Cas9-KO**
Strain background	**C57BL/6JGpt**

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Champ1* gene. The schematic diagram is as follows:



Technical routes

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➢ The *Champ1* gene has 2 transcripts. According to the structure of *Champ1* gene, exon2 of *Champ1-201* (ENSMUST00000051870.7) transcript is recommended as the knockout region.The region contains all of the coding sequenc Knock out the region will result in disruption of protein function.

➢ In this project we use CRISPR/Cas9 technology to modify *Champ1* gene. The brief process is as follows: gRNA was transcribed in vitro.Cas9 and gRNA 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

- The knockout region is near to the C-terminal of *Gm47277* gene and *Gm24698* and *Coprs* gene,this strategy may influence the regulatory function of the C-terminal of these genes.

- The *Champ1* 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information （NCBI）

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Champ1 chromosome alignment maintaining phosphoprotein 1 [*Mus musculus* (house mouse)]

Gene ID: 101994, updated on 14-Aug-2019

Summary

Official Symbol	Champ1 provided by MGI
Official Full Name	chromosome alignment maintaining phosphoprotein 1 provided by MGI
Primary source	MGI:MGI:1196398
See related	Ensembl:ENSMUSG00000047710
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	Zfp828; Znf828; AA675043; AI116001; mKIAA1802; D8Ertd457e; D8Ertd569e
Expression	Ubiquitous expression in whole brain E14.5 (RPKM 12.3), CNS E14 (RPKM 11.6) and 28 other tissues See more
Orthologs	human all

Genomic context

Location: 8 A1.1; 8 6.4 cM

See Champ1 in Genome Data Viewer

Exon count: 2

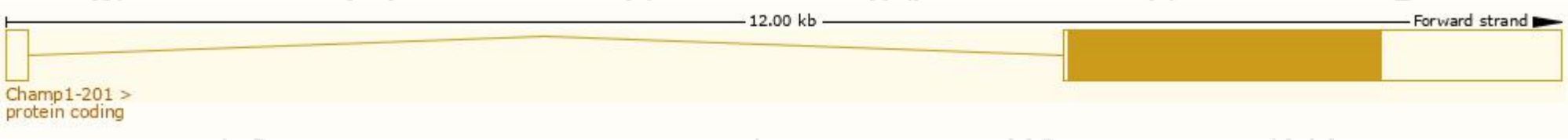
Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	8	NC_000074.6 (13869641..13881639)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	8	NC_000074.5 (13869641..13881639)

Transcript information（Ensembl）

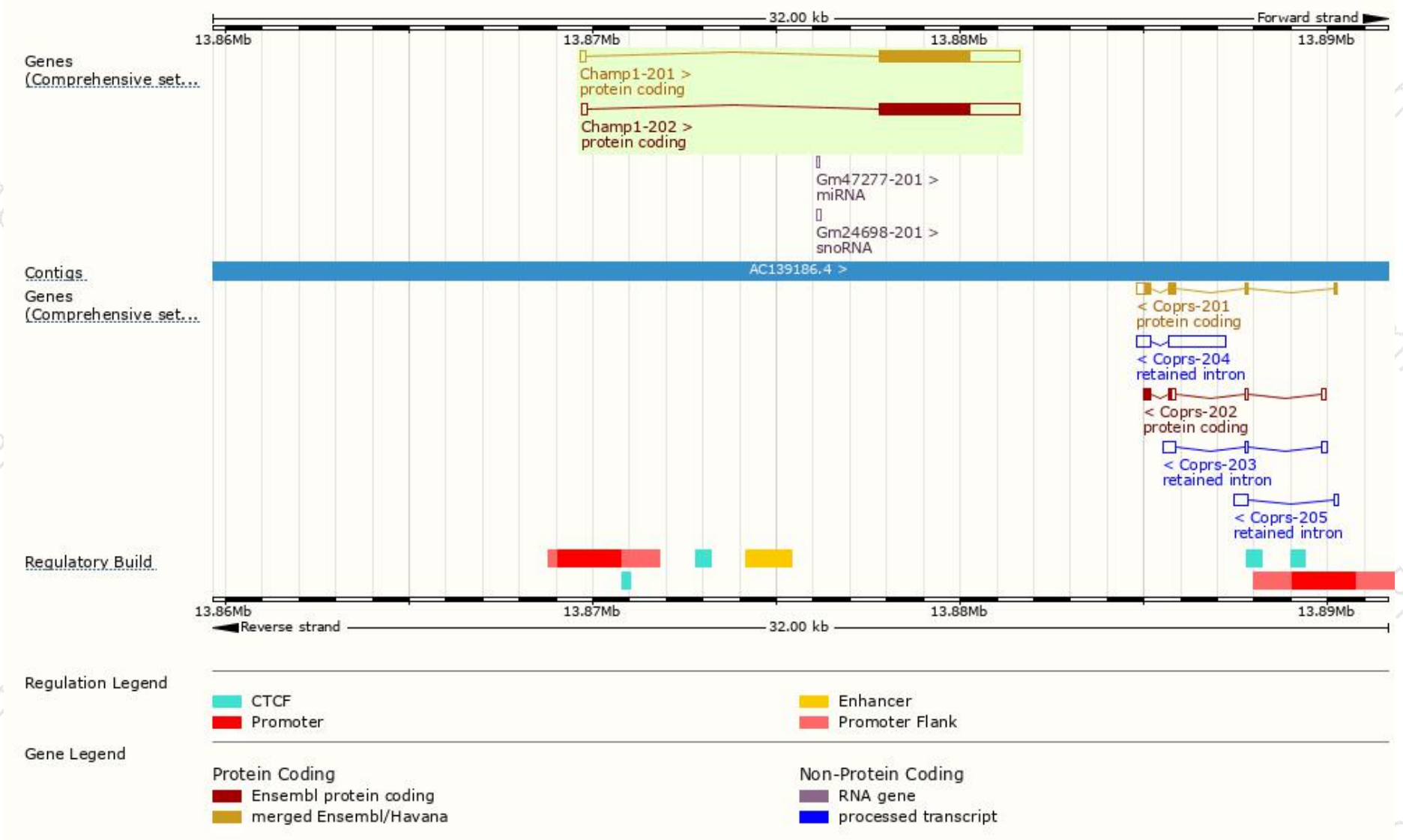
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Champ1-201	ENSMUST00000051870.7	4013	802aa	Protein coding	CCDS22116	Q8K327	TSL:1 GENCODE basic APPRIS P1
Champ1-202	ENSMUST00000128557.2	3977	802aa	Protein coding	CCDS22116	A0A140T8S5 Q8K327	TSL:1 GENCODE basic APPRIS P1

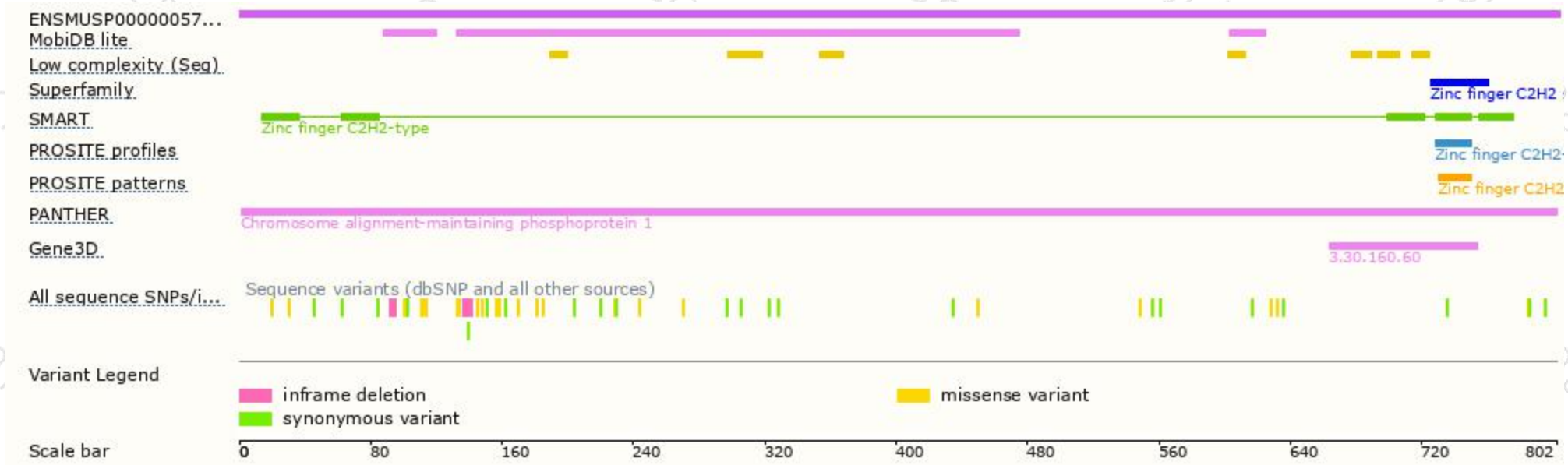
The strategy is based on the design of *Champ1-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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