Pqbp1 Cas9-KO Strategy Rohalmakech Co.

Designer: Condo de Co

Mohamar Ch

Qiong Zhou

Project Overview



Project Name

Pqbp1

Project type

Cas9-KO

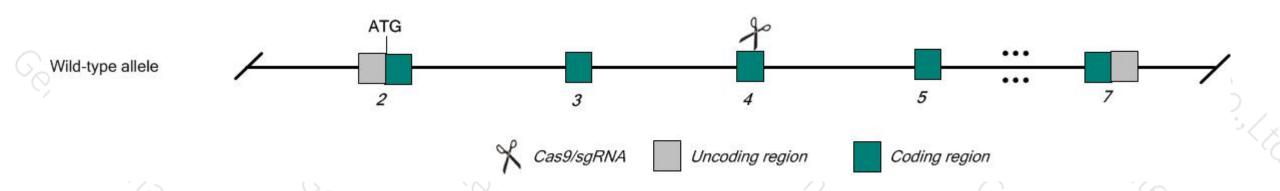
Strain background

C57BL/6J

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Pqbp1* gene. The schematic diagram is as follows:



Technical routes



- The *Pqbp1* gene has 5 transcripts. According to the structure of *Pqbp1* gene, partial exon4 of *Pqbp1*-203 (ENSMUST00000115655.7)transcript is recommended as the knockout region. The region contains key coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Pqbp1* gene. The brief process is as follows: sgRNA was transcribed in vitro.Cas9 and sgRNA were microinjected into the fertilized eggs of C57BL/6J 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/6J mice.

Notice



- The *Pqbp1* gene is located on the ChrX. 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)



Pqbp1 polyglutamine binding protein 1 [Mus musculus (house mouse)]

Gene ID: 54633, updated on 31-Jan-2019

Summary

Official Symbol Pgbp1 provided by MGI

Official Full Name polyglutamine binding protein 1 provided by MGI

Primary source MGI:MGI:1859638

See related Ensembl: ENSMUSG00000031157

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 Sfc2; npw38; PQBP-1

Expression Ubiquitous expression in CNS E14 (RPKM 52.5), CNS E11.5 (RPKM 48.3) and 27 other tissues See more

Orthologs <u>human</u> all

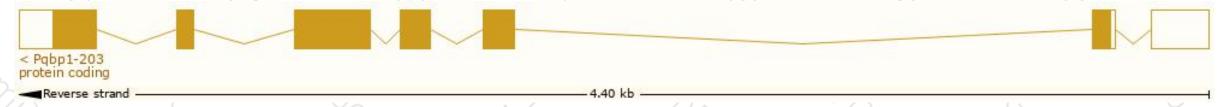
Transcript information (Ensembl)



The gene has 5 transcripts, and all transcripts are shown below:

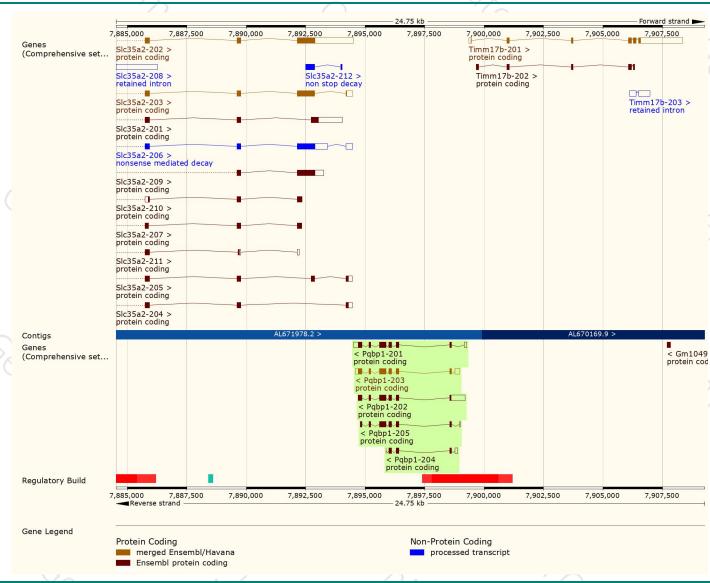
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Name 👙	Transcript ID 👙	bp 🍦	Protein 🌲	Biotype 🍦	CCDS 🍦	UniProt 🍦	Flags
Pqbp1-202	ENSMUST00000115654.7	1374	<u>263aa</u>	Protein coding	<u>CCDS29977</u> ₽	<u>A2AER7</u> @ <u>Q91VJ5</u> @	TSL:1 GENCODE basic APPRIS P1
Pqbp1-203	ENSMUST00000115655.7	1148	<u>263aa</u>	Protein coding	CCDS29977 ₽	<u>A2AER7</u> Ø <u>Q91VJ5</u> Ø	TSL:1 GENCODE basic APPRIS P1
Pqbp1-201	ENSMUST00000033497.8	1104	<u>263aa</u>	Protein coding	CCDS29977 ₽	<u>A2AER7</u> @ <u>Q91√J5</u> @	TSL:1 GENCODE basic APPRIS P1
Pqbp1-205	ENSMUST00000156741.7	763	236aa	Protein coding	-	<u>A2AER8</u> ₽	CDS 3' incomplete TSL:5
Pqbp1-204	ENSMUST00000154552.1	436	<u>100aa</u>	Protein coding	92	<u>A2AER9</u> ₽	CDS 3' incomplete TSL:3

The strategy is based on the design of Pqbp1-203 transcript, The transcription is shown below



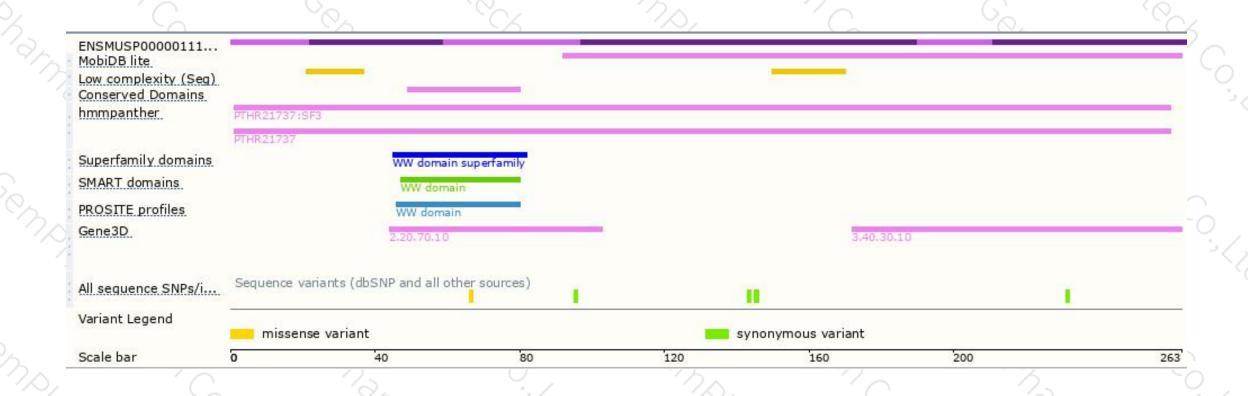
Genomic location (Ensembl)





Protein domain (Ensembl)





If you have any questions, you are welcome to inquire. Tel: 025-5864 1534





