

Donald Color Bmx Cas9-KO Strategy Rohalmakech Co.

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Condand Stock Designer: Lixin Lv

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Project Overview



Project Name

Bmx

Project type

Cas9-KO

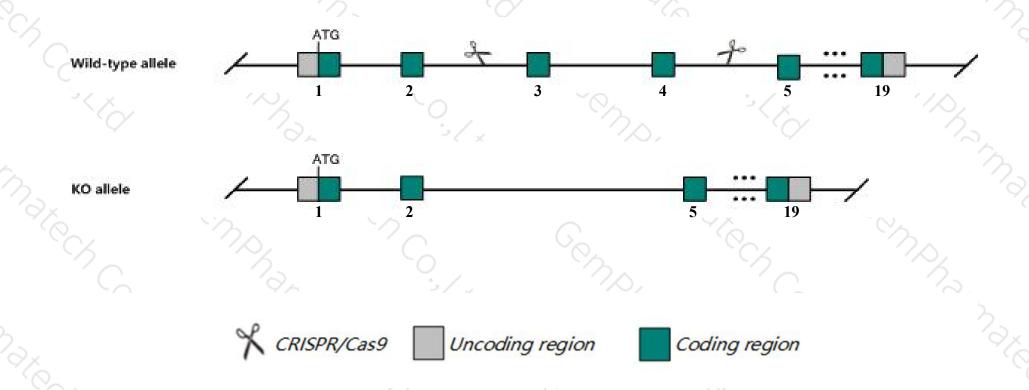
Strain background

C57BL/6JGpt

Knockout strategy



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



Technical routes



- ➤ The *Bmx* gene has 2 transcripts. According to the structure of *Bmx* gene, exon3-exon4 of *Bmx-202*(ENSMUST00000112265.8) transcript is recommended as the knockout region. The region contains 187bp coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Bmx* gene. The brief process is as follows: CRISPR/Cas9 system v

Notice



- ➤ According to the existing MGI data, Mice homozygous for a null mutation are fertile and have a normal life span but are protected from arthritis in a passive transfer model.
- The *Bmx* 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)



Bmx BMX non-receptor tyrosine kinase [Mus musculus (house mouse)]

Gene ID: 12169, updated on 31-Jan-2019

Summary

☆ ?

Official Symbol Bmx provided by MGI

Official Full Name BMX non-receptor tyrosine kinase provided by MGI

Primary source MGI:MGI:1101778

See related Ensembl:ENSMUSG00000031377

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 Etk, Etk/Bmx, Tyro8

Expression Broad expression in bladder adult (RPKM 1.1), lung adult (RPKM 0.9) and 22 other tissuesSee more

Orthologs human all

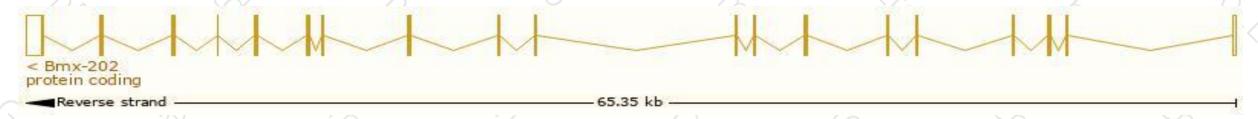
Transcript information (Ensembl)



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

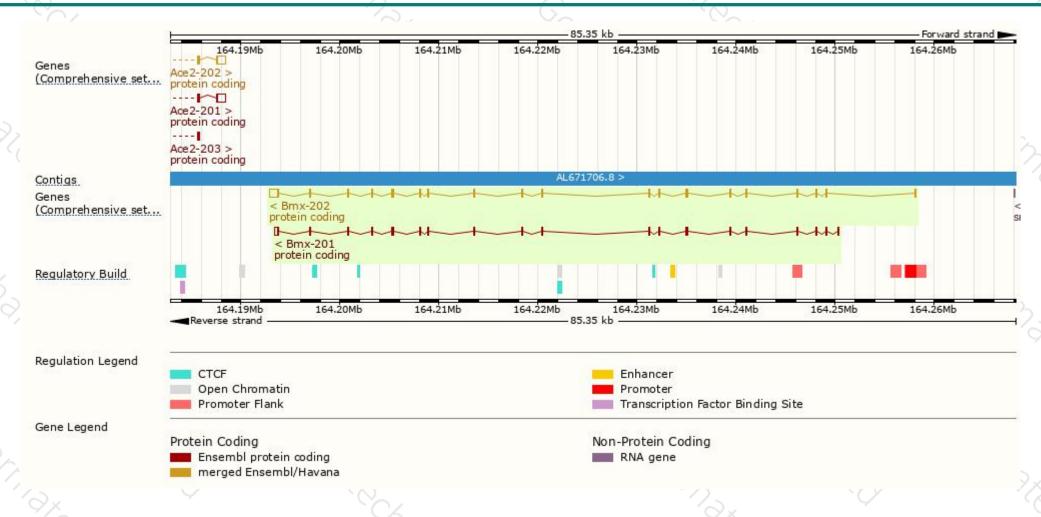
Name 🍦	Transcript ID .	bp 🍦	Protein 🍦	Biotype	CCDS 🍦	UniProt 🍦	Flags		
Bmx-202	ENSMUST00000112265.8	3005	<u>655aa</u>	Protein coding	CCDS30519₽	B1AUL6译	TSL:1	GENCODE basic	APPRIS P2
Bmx-201	ENSMUST00000112263.1	2433	<u>651aa</u>	Protein coding	150	B1AUL5@P97504@	TSL:5	GENCODE basic	APPRIS ALT2

The strategy is based on the design of Bmx-202 transcript, The transcription is shown below



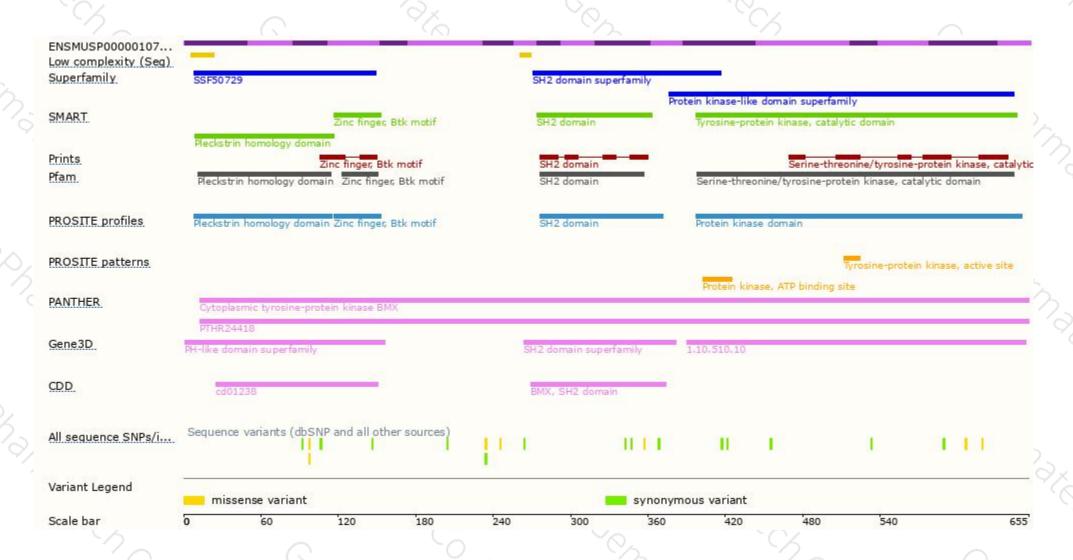
Genomic location distribution





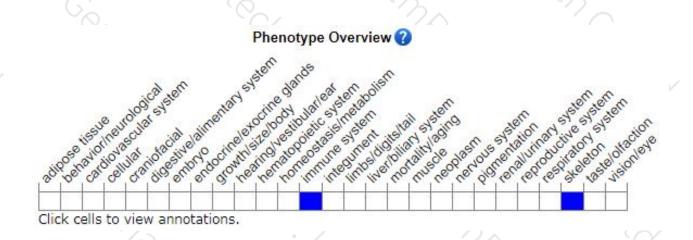
Protein domain





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 null mutation are fertile and have a normal life span but are protected from arthritis in a passive transfer model.



If you have any questions, you are welcome to inquire. Tel: 400-9660890





