

Btk Cas9-KO Strategy

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Project Overview



Project Name Btk

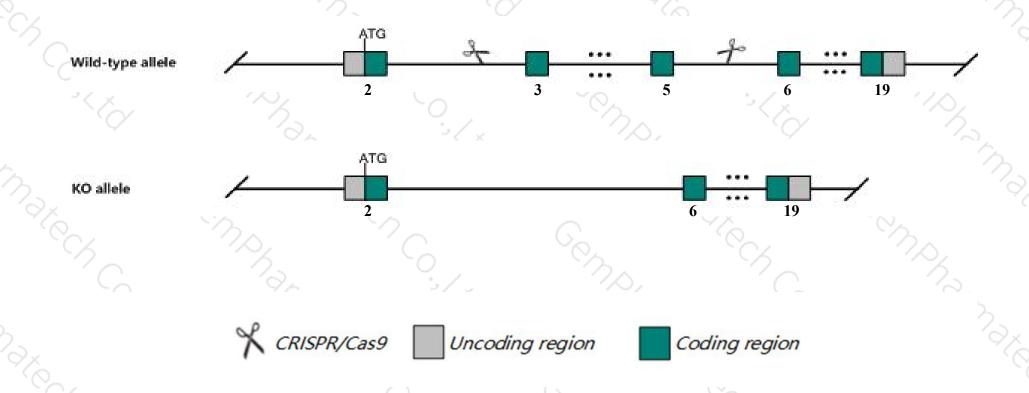
Project type Cas9-KO

Strain background C57BL/6JGpt

Knockout strategy



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



Technical routes



- ➤ The *Btk* gene has 5 transcripts. According to the structure of *Btk* gene, exon3-exon5 of *Btk-201*(ENSMUST00000033617.12) transcript is recommended as the knockout region. The region contains 250bp coding sequence Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Btk* gene. The brief process is as follows: CRISPR/Cas9 system we

Notice



- ➤ According to the existing MGI data, Mutants have immune defects including reduced B cell numbers, serum immunoglobulin deficiencies, and defective responses to B cell activators and thymus-independent antigens. B-1 B cells are absent in these mice.
- > The *Btk* 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)



Btk Bruton agammaglobulinemia tyrosine kinase [Mus musculus (house mouse)]

Gene ID: 12229, updated on 31-Jan-2019

Summary

☆ ?

Official Symbol Btk provided by MGI

Official Full Name Bruton agammaglobulinemia tyrosine kinase provided byMGI

Primary source MGI:MGI:88216

See related Ensembl: ENSMUSG00000031264

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 Al528679, xid

Expression Biased expression in liver E14 (RPKM 7.3), liver E14.5 (RPKM 5.9) and 11 other tissuesSee more

Orthologs <u>human</u> all

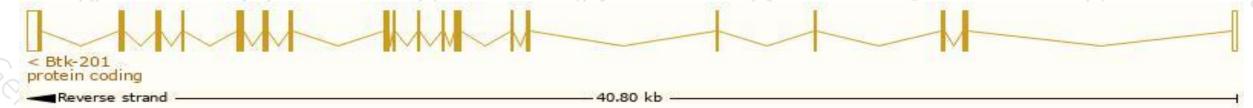
Transcript information (Ensembl)



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

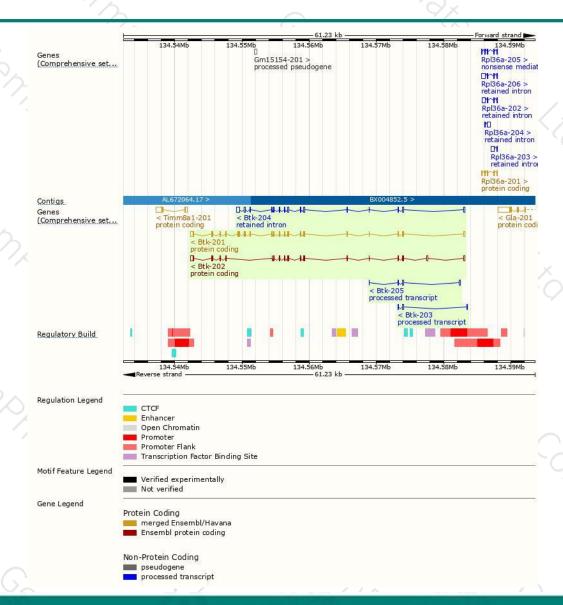
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Btk-201	ENSMUST00000033617.12	2540	659aa	Protein coding	CCDS30396	P35991	TSL:1 GENCODE basic APPRIS P1
Btk-202	ENSMUST00000113213.1	2121	<u>483aa</u>	Protein coding	-	A2BDW0	TSL:5 GENCODE basic
Btk-205	ENSMUST00000150245.7	481	No protein	Processed transcript		20	TSL:2
Btk-203	ENSMUST00000128333.1	388	No protein	Processed transcript	22	20	TSL:3
Btk-204	ENSMUST00000132664.7	1920	No protein	Retained intron	-	-	TSL:5

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



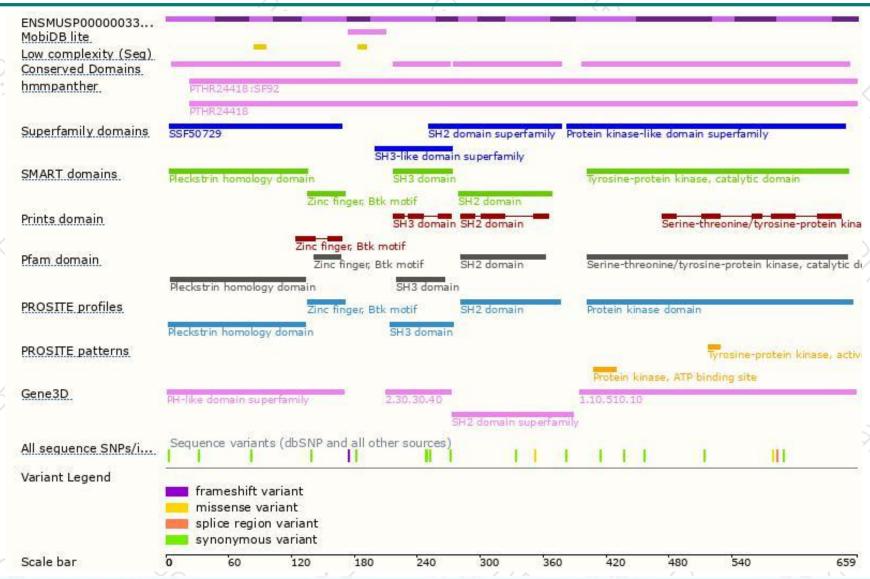
Genomic location distribution





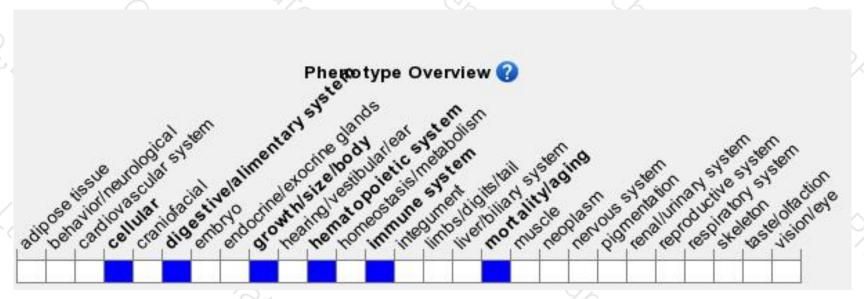
Protein domain





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, Mutants have immune defects including reduced B cell numbers, serum immunoglobulin deficiencies, and defective responses to B cell activators and thymus-independent antigens. B-1 B cells are absent in these mice.



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





