

Bcl7a Cas9-CKO Strategy

Designer:

Jinling Wang

Design Date:

2019-9-30

Project Overview

Project Name

Bcl7a

Project type

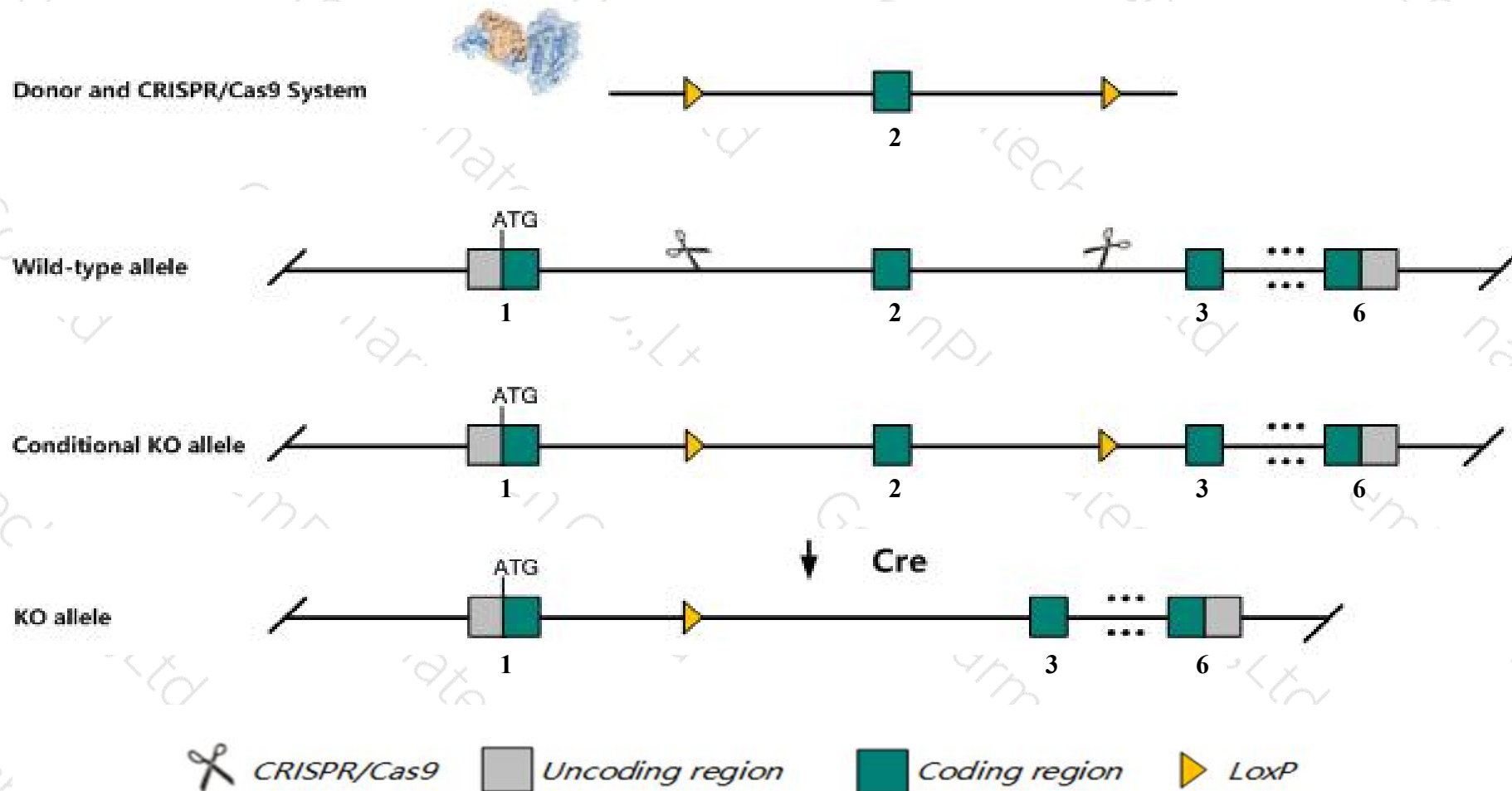
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Bcl7a* gene. The schematic diagram is as follows:



Technical routes

- The *Bcl7a* gene has 3 transcripts. According to the structure of *Bcl7a* gene, exon2 of *Bcl7a-201* (ENSMUST00000031391.8) transcript is recommended as the knockout region. The region contains 82bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Bcl7a* gene. The brief process is as follows: CRISPR/Cas9 system and Donor 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.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

Notice

- The *Bcl7a* gene is located on the Chr5. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)

Bcl7a B cell CLL/lymphoma 7A [Mus musculus (house mouse)]

Gene ID: 77045, updated on 31-Jan-2019

Summary



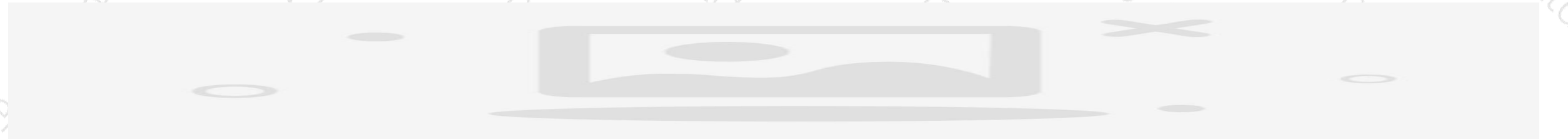
Official Symbol	Bcl7a provided by MGI
Official Full Name	B cell CLL/lymphoma 7A provided by MGI
Primary source	MGI:MGI:1924295
See related	Ensembl:ENSMUSG00000029438
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	4432415N06Rik, A1448316
Expression	Ubiquitous expression in CNS E14 (RPKM 21.1), CNS E11.5 (RPKM 20.9) and 26 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

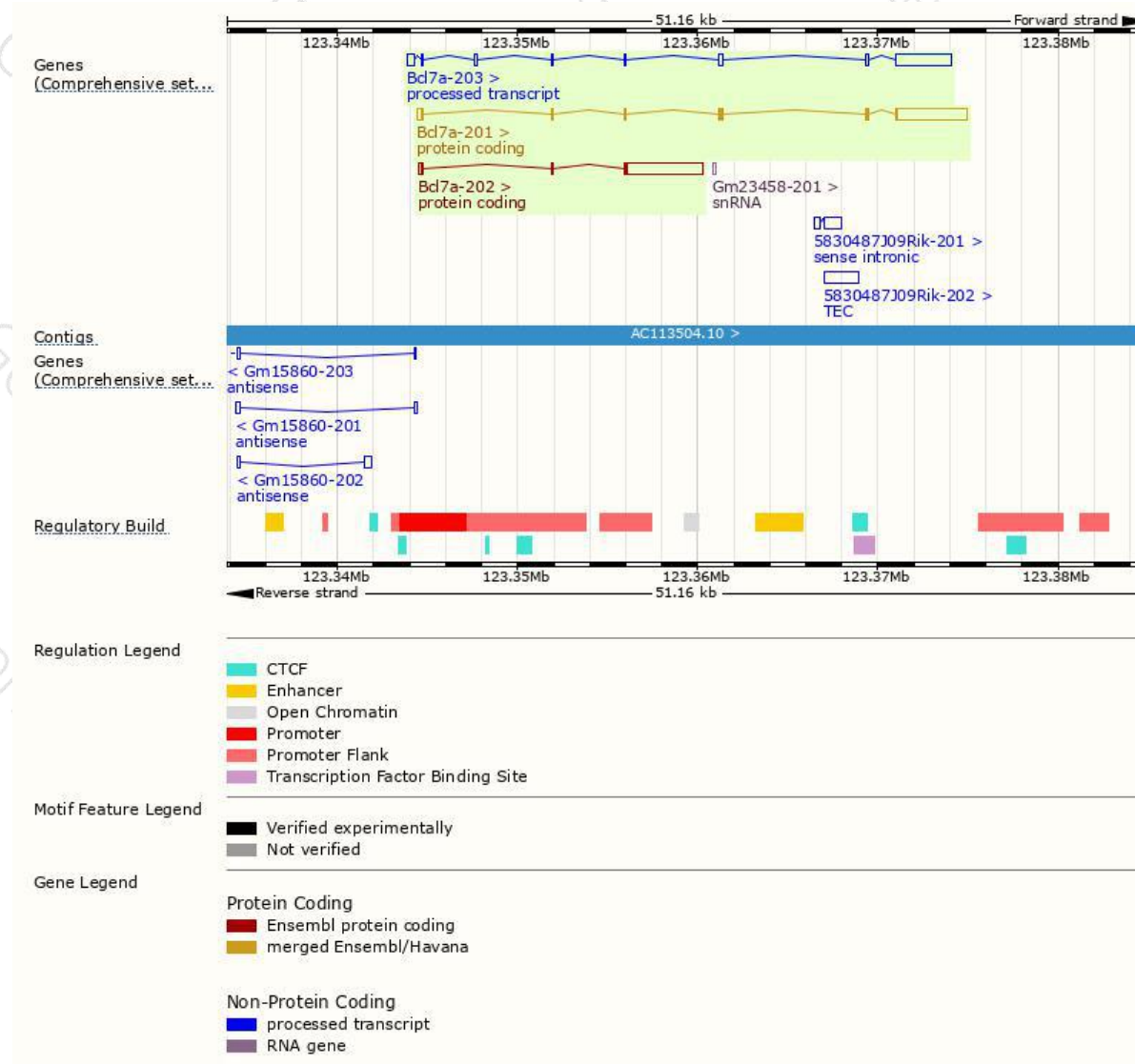
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Bcl7a-201	ENSMUST00000031391.8	4718	210aa	Protein coding	CCDS19661	Q9CXE2	TSL:1 GENCODE basic APPRIS P1
Bcl7a-202	ENSMUST00000117971.1	4642	112aa	Protein coding	-	Q9CXE2	TSL:1 GENCODE basic
Bcl7a-203	ENSMUST00000120593.4	4161	No protein	Processed transcript	-	-	TSL:5

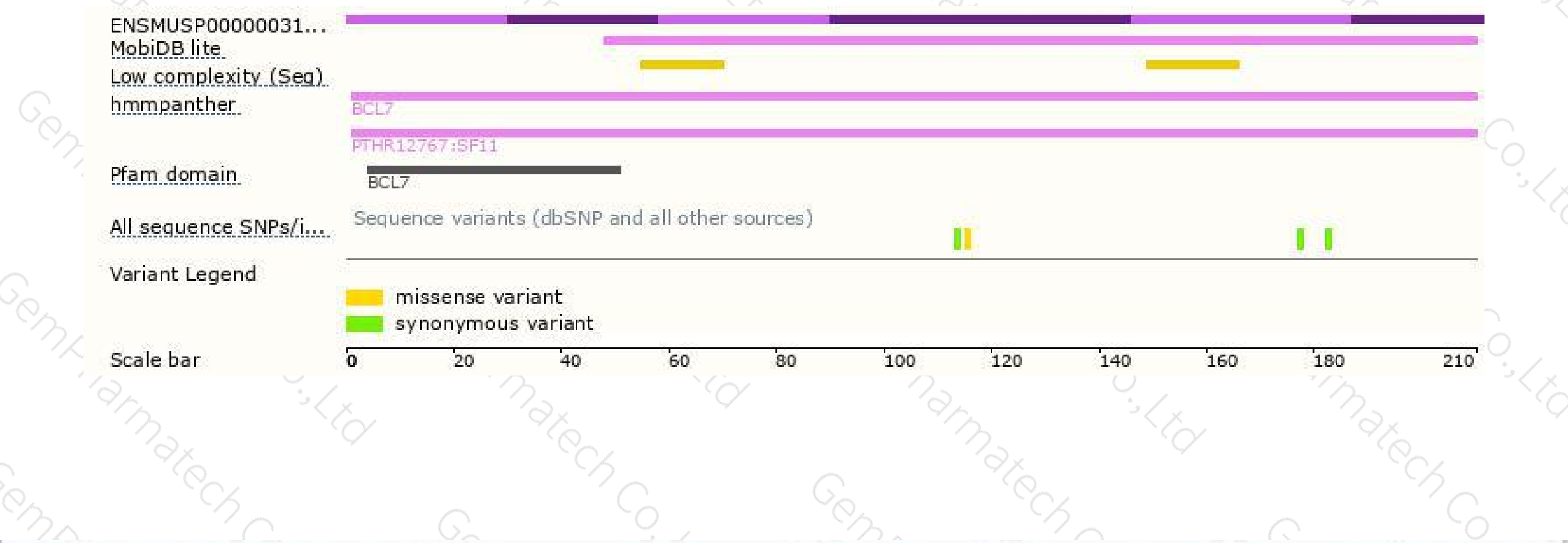
The strategy is based on the design of *Bcl7a-201* transcript,The transcription is shown below



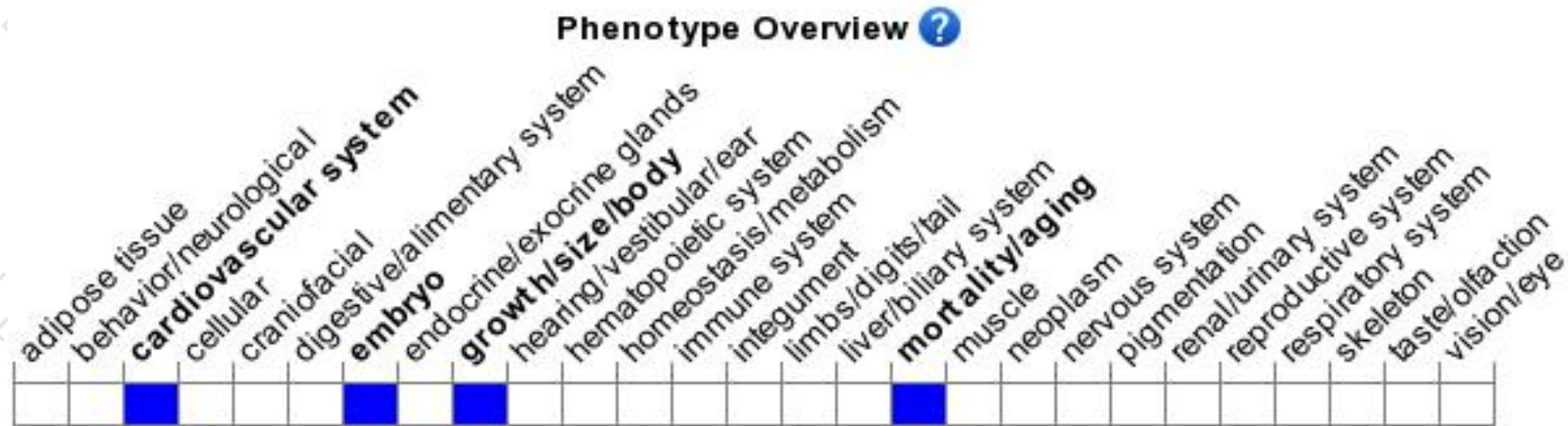
Genomic location distribution



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/>).

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

Tel: 400-9660890

