

Btn1a1 Cas9-CKO Strategy

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Design Date:

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Project Overview

Project Name

Btn1a1

Project type

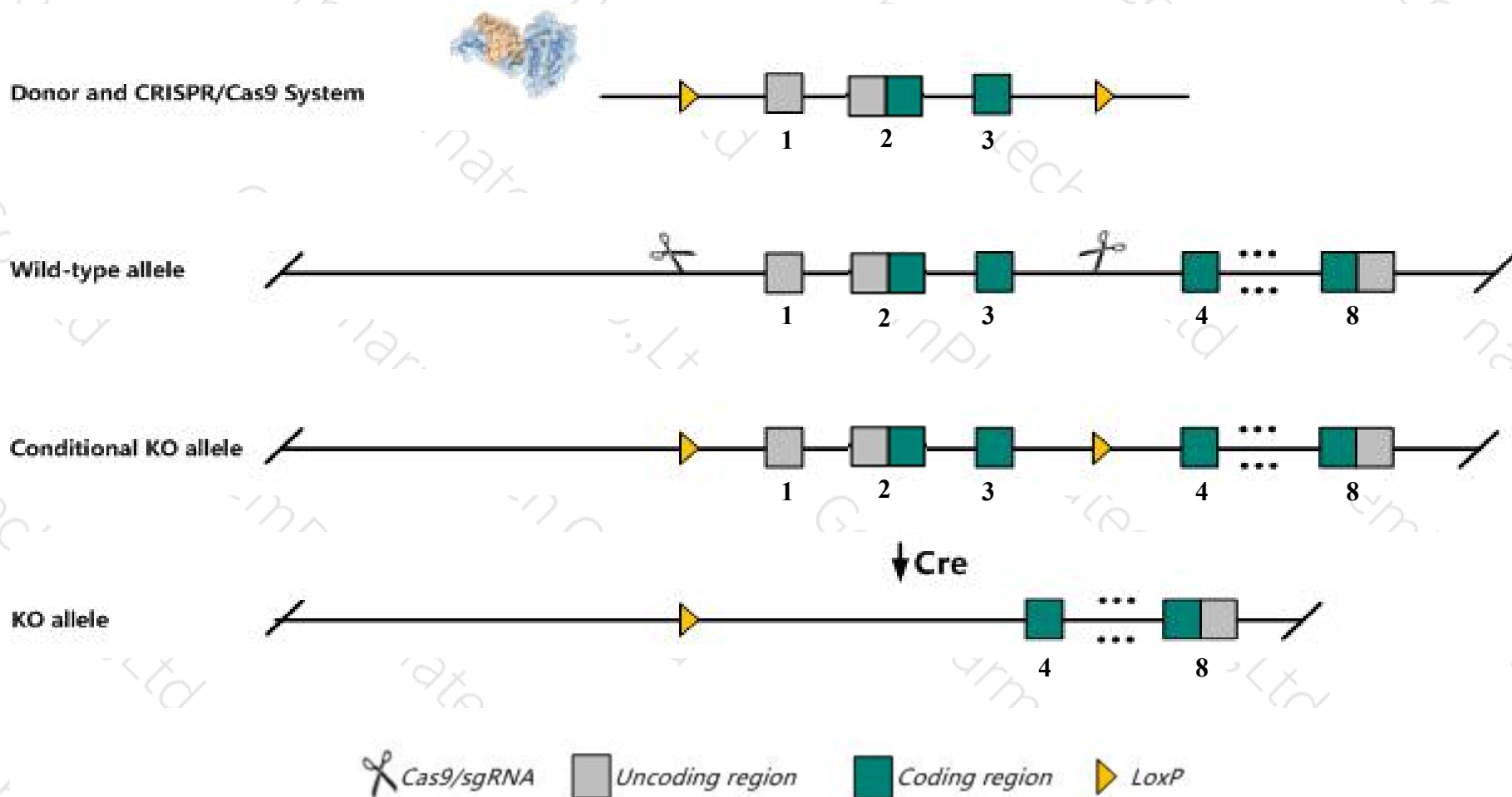
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Btn1a1* gene. The schematic diagram is as follows:



Technical routes

- The *Btn1a1* gene has 3 transcripts. According to the structure of *Btn1a1* gene, exon1-exon3 of *Btn1a1-201* (ENSMUST00000041674.13) transcript is recommended as the knockout region. The region contains the translational start codon ATG. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Btn1a1* 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.

- According to the existing MGI data, Homozygous null mice display impaired lactation and abnormal lipid accumulation in the mammary gland.
- The *Btn1a1* gene is located on the Chr13. 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.

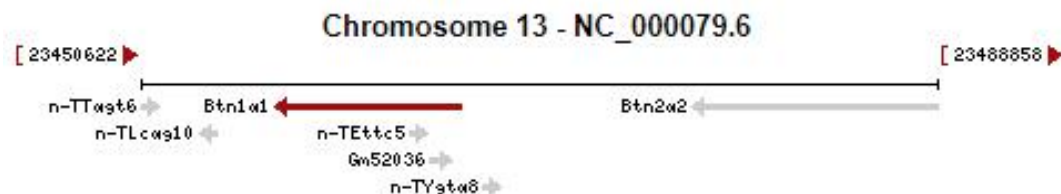
Gene information (NCBI)

Btn1a1 butyrophilin, subfamily 1, member A1 [*Mus musculus* (house mouse)]

Gene ID: 12231, updated on 12-Aug-2019

Summary

Official Symbol	Btn1a1 provided by MGI
Official Full Name	butyrophilin, subfamily 1, member A1 provided by MGI
Primary source	MGI:MGI:103118
See related	Ensembl:ENSMUSG000000000706
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	Btn
Expression	Biased expression in mammary gland adult (RPKM 3.3), liver E14.5 (RPKM 1.1) and 3 other tissues See more
Orthologs	human all

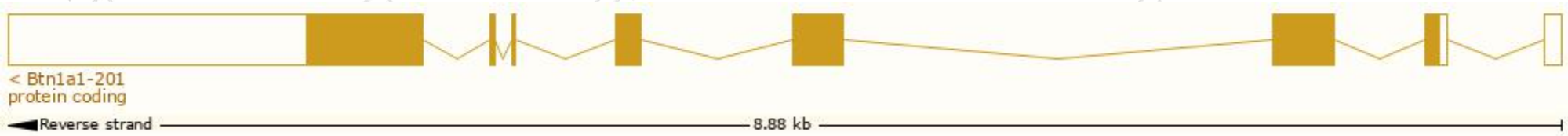


Transcript information (Ensembl)

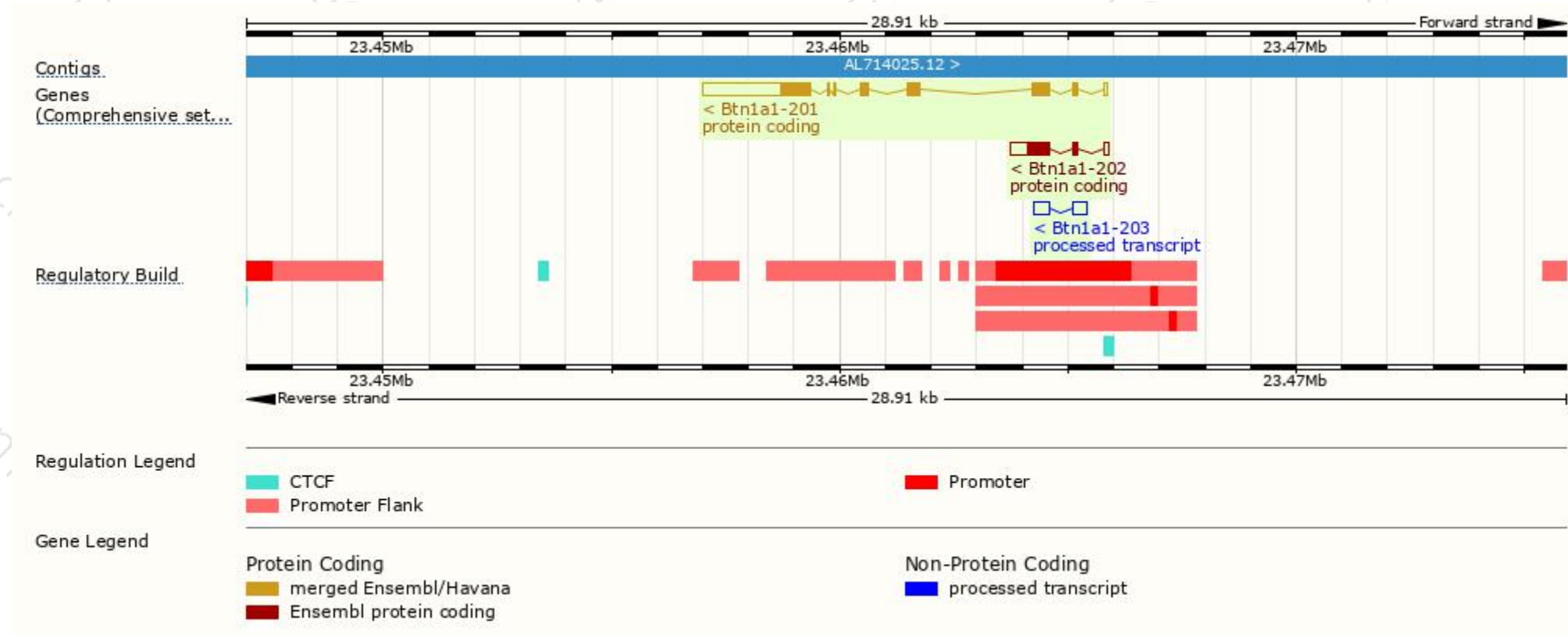
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Btn1a1-201	ENSMUST00000041674.13	3423	524aa	Protein coding	CCDS26339	Q3UQC0 Q62556	TSL:1 GENCODE basic APPRIS P1
Btn1a1-202	ENSMUST00000110434.1	1077	179aa	Protein coding	-	Q921X1	TSL:1 GENCODE basic
Btn1a1-203	ENSMUST00000225831.1	647	No protein	Processed transcript	-	-	-

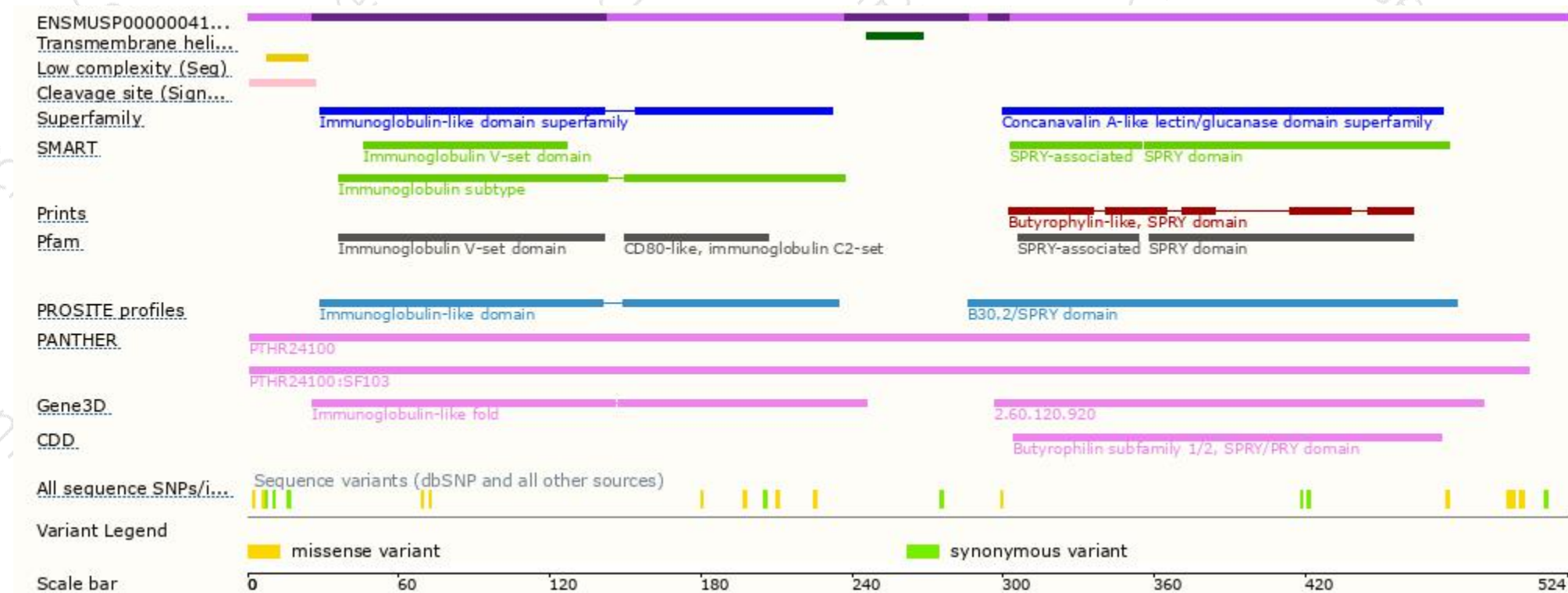
The strategy is based on the design of *Btn1a1-201* transcript,The transcription is shown below



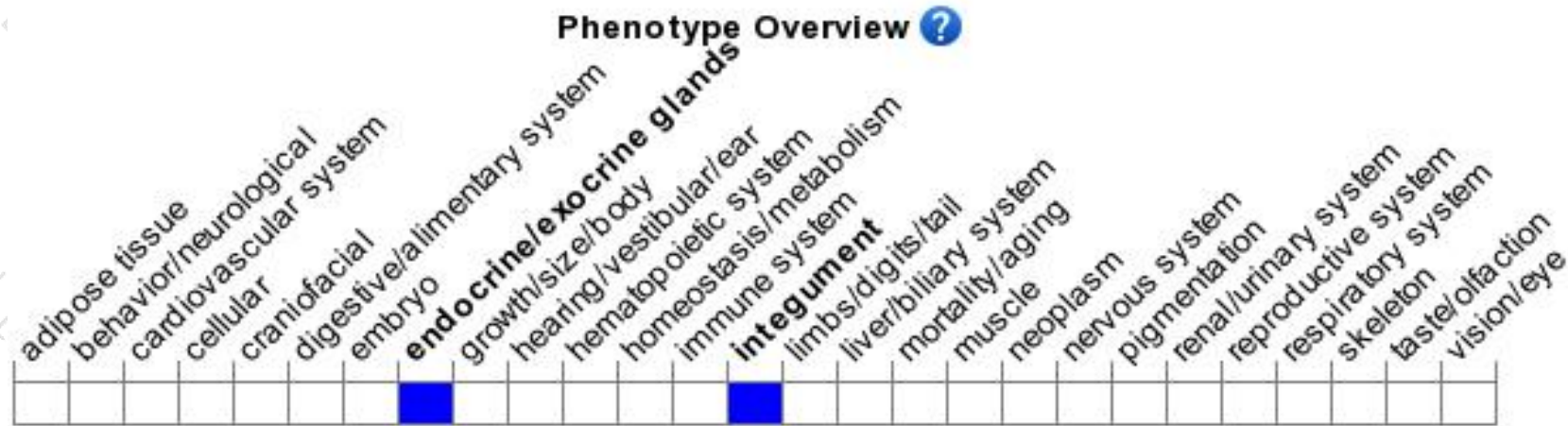
Genomic location distribution



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, Homozygous null mice display impaired lactation and abnormal lipid accumulation in the mammary gland.

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

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