

Igll1 Cas9-KO Strategy

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Reviewer:

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Project Overview

Project Name

Igll1

Project type

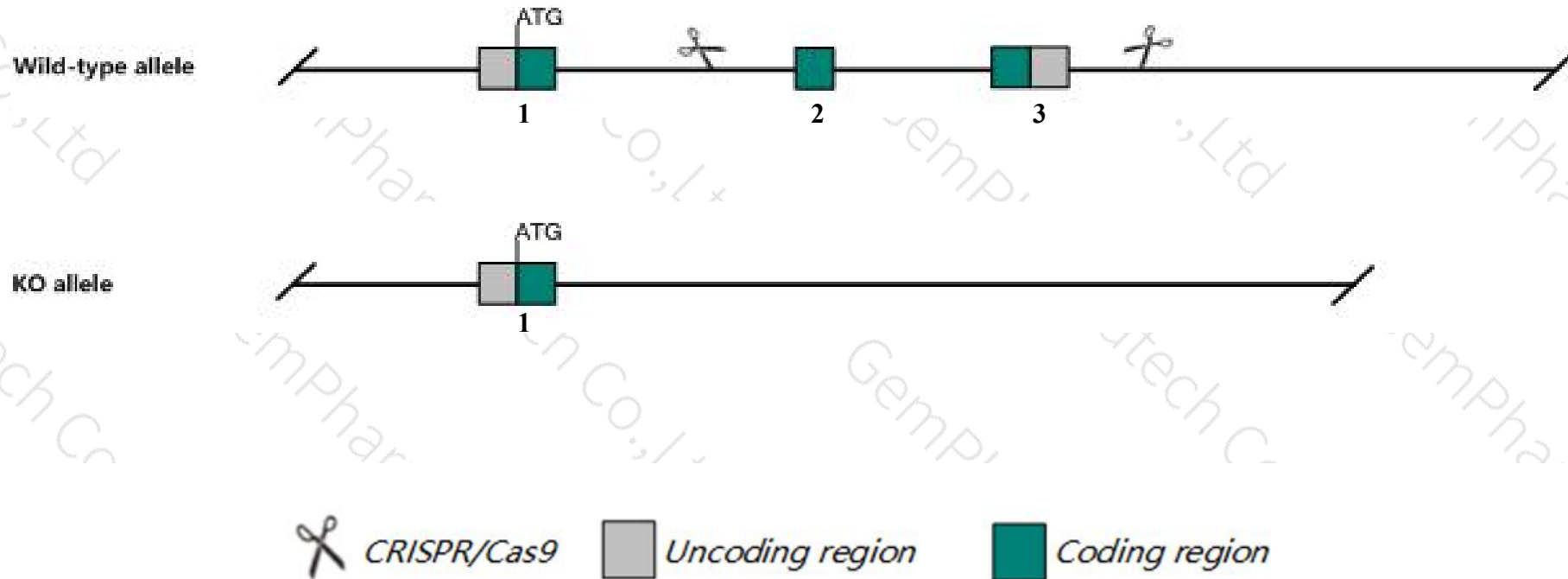
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Igll1* gene. The schematic diagram is as follows:



- The *Igll1* gene has 2 transcripts. According to the structure of *Igll1* gene, exon2-exon3 of *Igll1-201* (ENSMUST00000100136.3) transcript is recommended as the knockout region. The region contains 436bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Igll1* gene. The brief process is as follows: CRISPR/Cas9 system w

- According to the existing MGI data, Mice homozygous for a knock-out allele exhibit spleen hypoplasia, a leaky blockade of B cell development at the pre-B stage, and decreased IgG levels in response to a T-cell dependent antigen.
- The *Igll1* gene is located on the Chr16. 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.

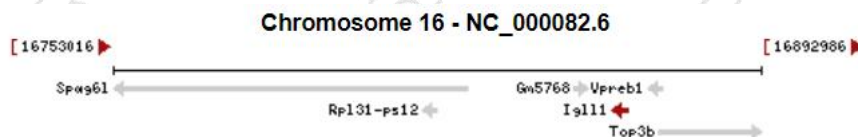
Gene information (NCBI)

Igll1 immunoglobulin lambda-like polypeptide 1 [*Mus musculus* (house mouse)]

Gene ID: 16136, updated on 22-Oct-2019

Summary

Official Symbol	Igll1 provided by MGI
Official Full Name	immunoglobulin lambda-like polypeptide 1 provided by MGI
Primary source	MGI:MGI:96529
See related	Ensembl:ENSMUSG00000075370
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	Igll; Igl-5; Lambda5; BB139905
Expression	Biased expression in liver E18 (RPKM 12.8), liver E14.5 (RPKM 0.9) and 1 other tissue See more
Orthologs	human all

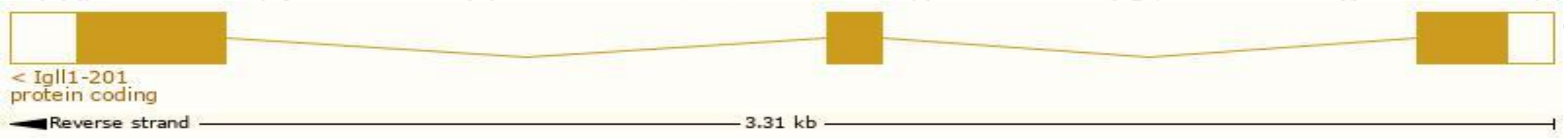


Transcript information (Ensembl)

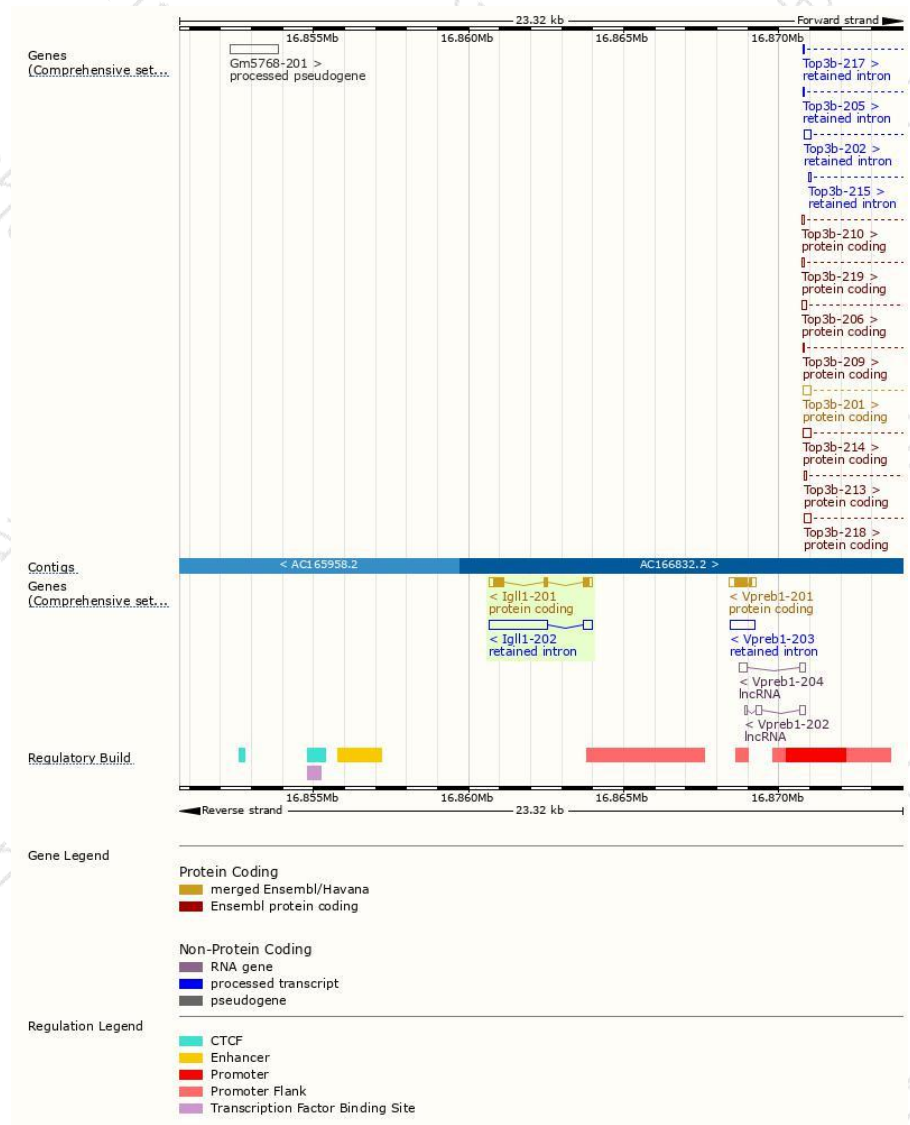
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Igll1-201	ENSMUST00000100136.3	870	209aa	Protein coding	CCDS49775	P20764	TSL:5 GENCODE basic APPRIS P1
Igll1-202	ENSMUST00000231439.1	2161	No protein	Retained intron	-	-	

The strategy is based on the design of *Igll1-201* transcript,The transcription is shown below



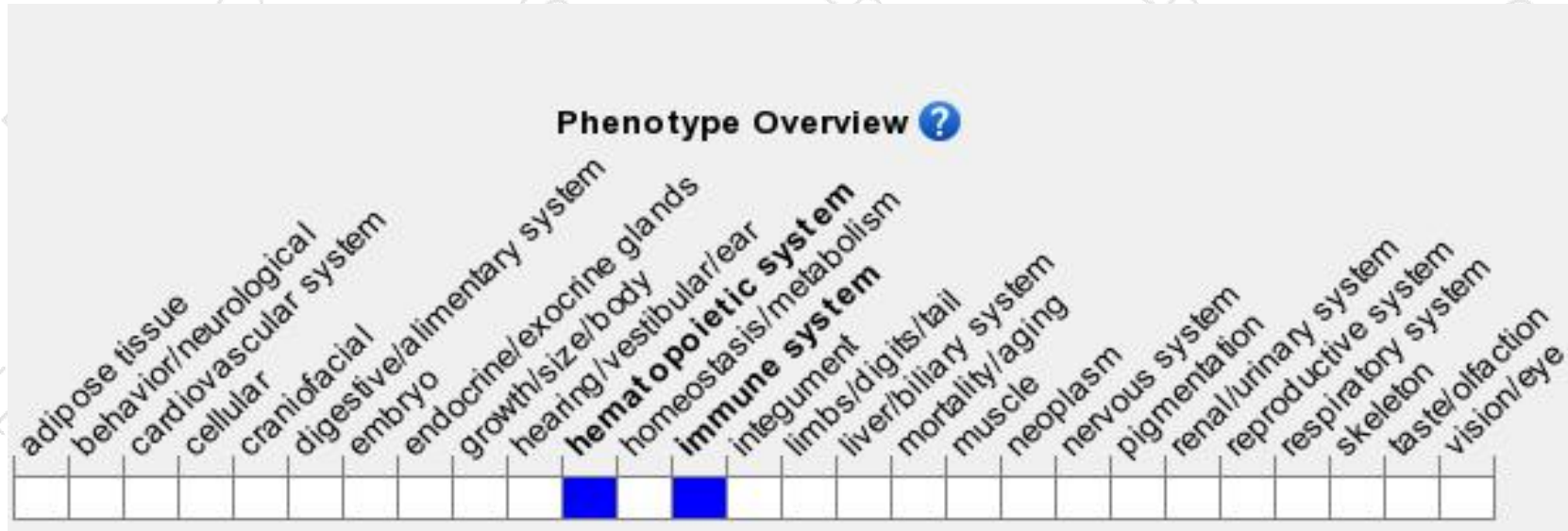
Genomic location distribution



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, Mice homozygous for a knock-out allele exhibit spleen hypoplasia, a leaky blockade of B cell development at the pre-B stage, and decreased IgG levels in response to a T-cell dependent antigen.

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

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