

Cdh17 Cas9-CKO Strategy

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Design Date:2020-2-11

Project Overview

Project Name

Cdh17

Project type

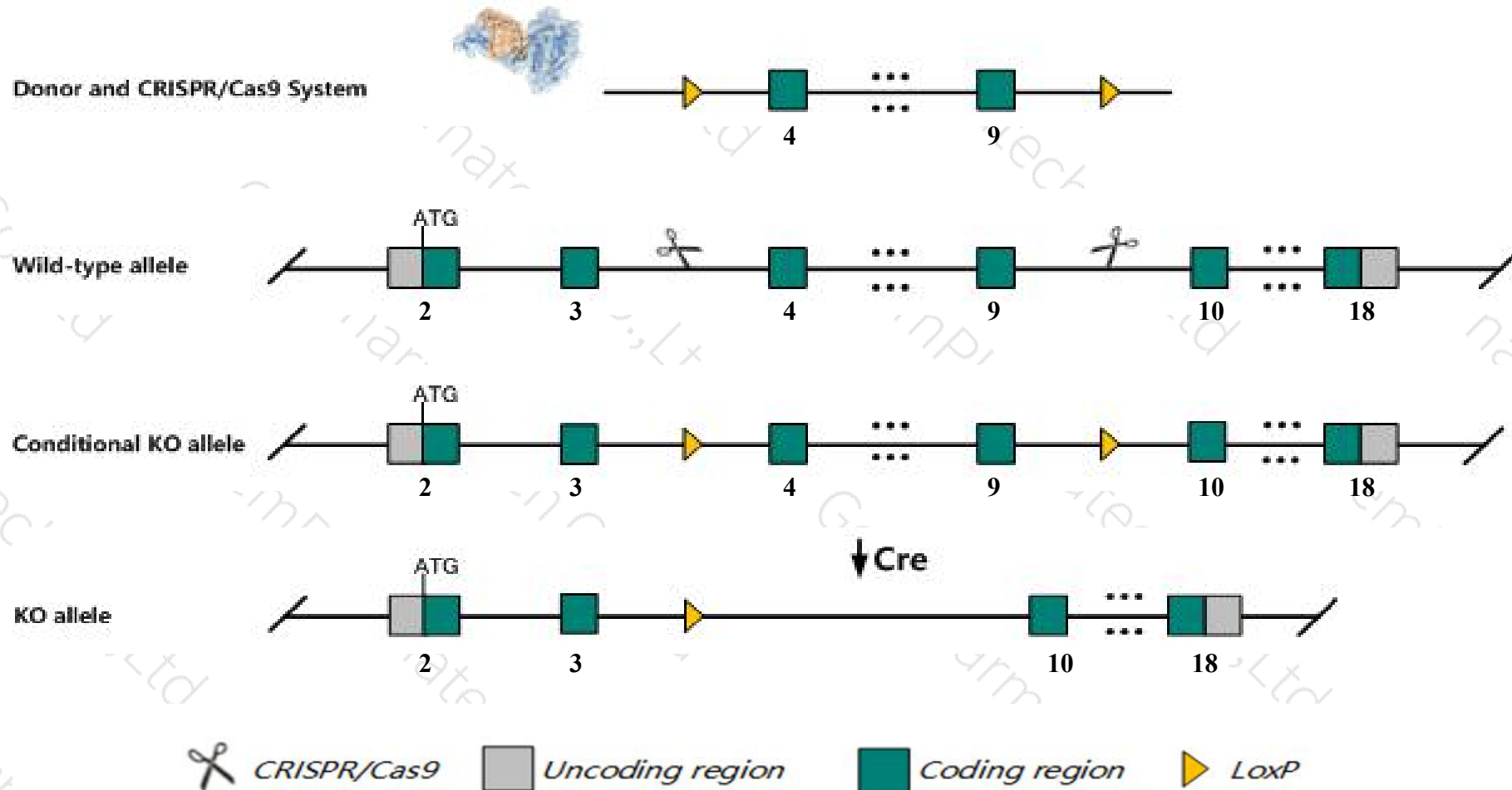
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Cdh17* gene. The schematic diagram is as follows:



- The *Cdh17* gene has 2 transcripts. According to the structure of *Cdh17* gene, exon4-exon9 of *Cdh17-201* (ENSMUST00000029871.11) transcript is recommended as the knockout region. The region contains 916bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Cdh17* 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 mutant mice exhibit impaired B lymphocyte development and impaired IgG1 and IgG3 antibody response to T-independent antigen.
- The *Cdh17* gene is located on the Chr4. 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)

Cdh17 cadherin 17 [*Mus musculus* (house mouse)]

Gene ID: 12557, updated on 26-Oct-2019

Summary

Official Symbol	Cdh17 provided by MGI
Official Full Name	cadherin 17 provided by MGI
Primary source	MGI:MGI:1095414
See related	Ensembl:ENSMUSG00000028217
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	HPT-1; HPT-1/LI
Expression	Biased expression in large intestine adult (RPKM 105.6), small intestine adult (RPKM 27.0) and 2 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

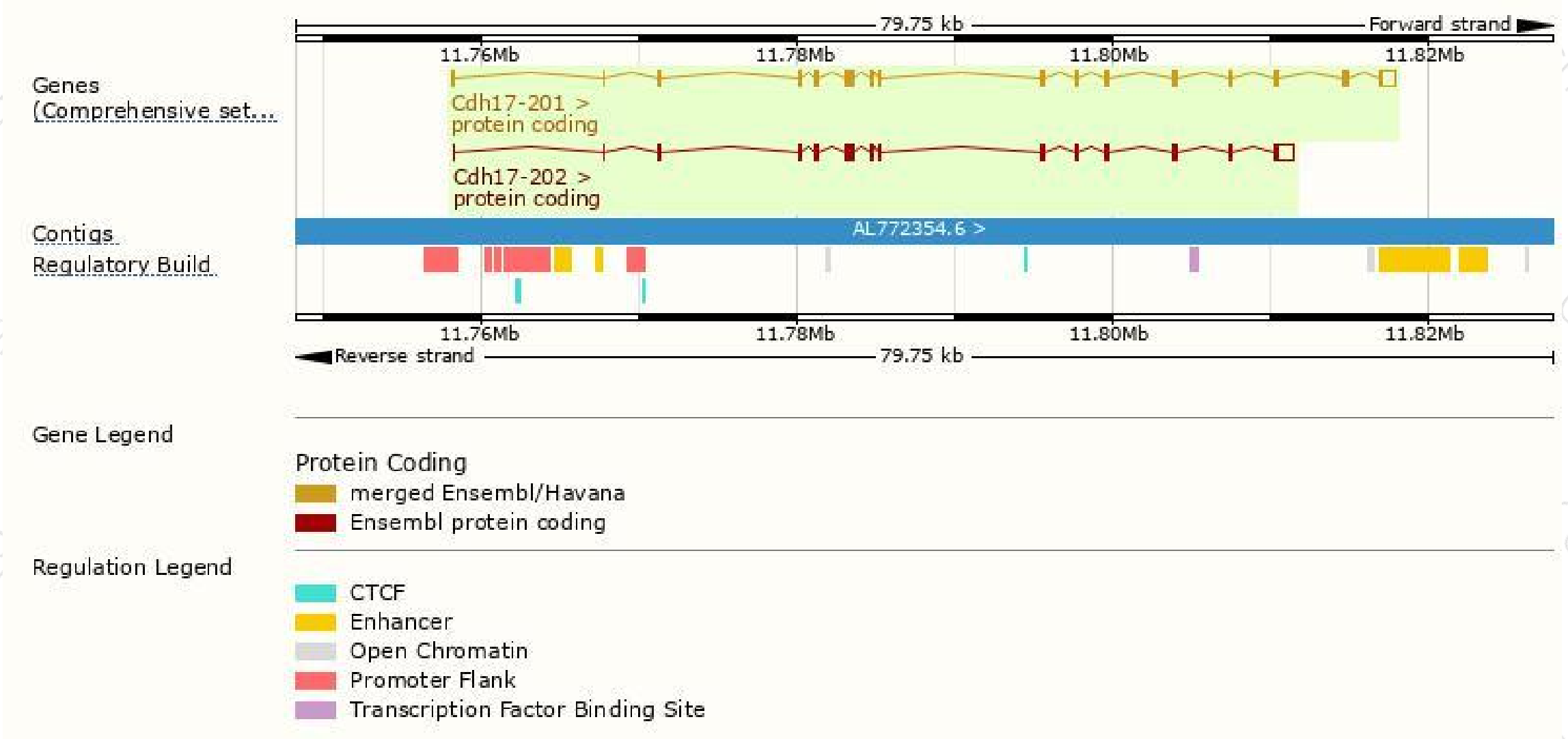
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Cdh17-201	ENSMUST00000029871.11	3462	827aa	Protein coding	CCDS17972	Q9R100	TSL:1 GENCODE basic APPRIS P1
Cdh17-202	ENSMUST00000108303.1	3287	728aa	Protein coding	-	A2AKS7	TSL:2 GENCODE basic

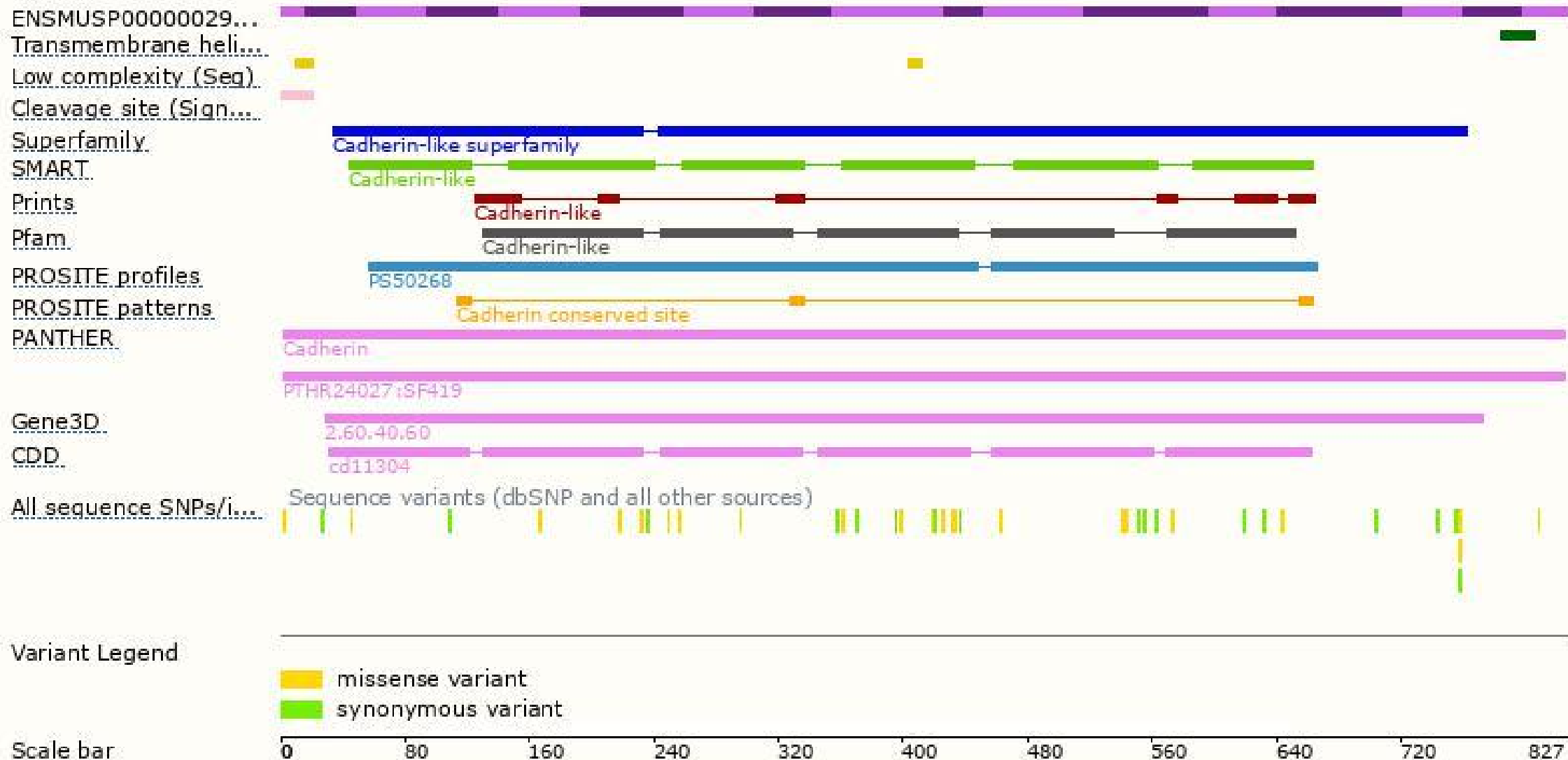
The strategy is based on the design of *Cdh17-201* transcript,The transcription is shown below



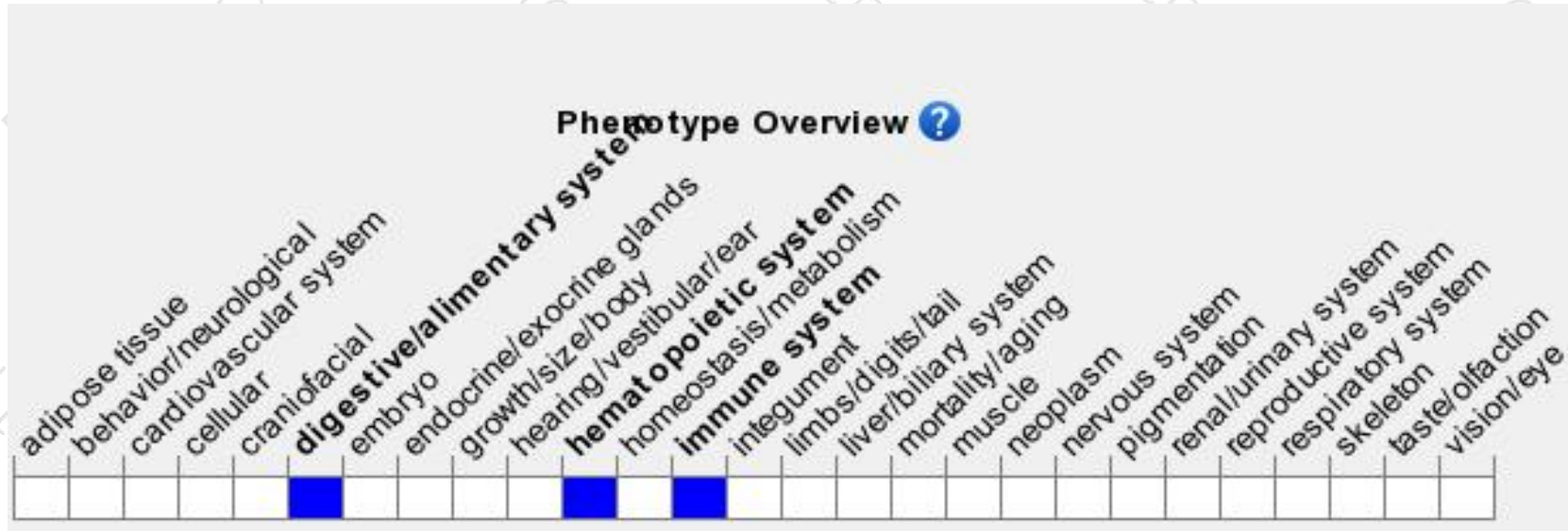
Genomic location distribution



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 mutant mice exhibit impaired B lymphocyte development and impaired IgG and IgG3 antibody response to T-independent antigen.

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

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