

Arfgef3 Cas9-CKO Strategy

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Reviewer:

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Project Overview

Project Name

Arfgef3

Project type

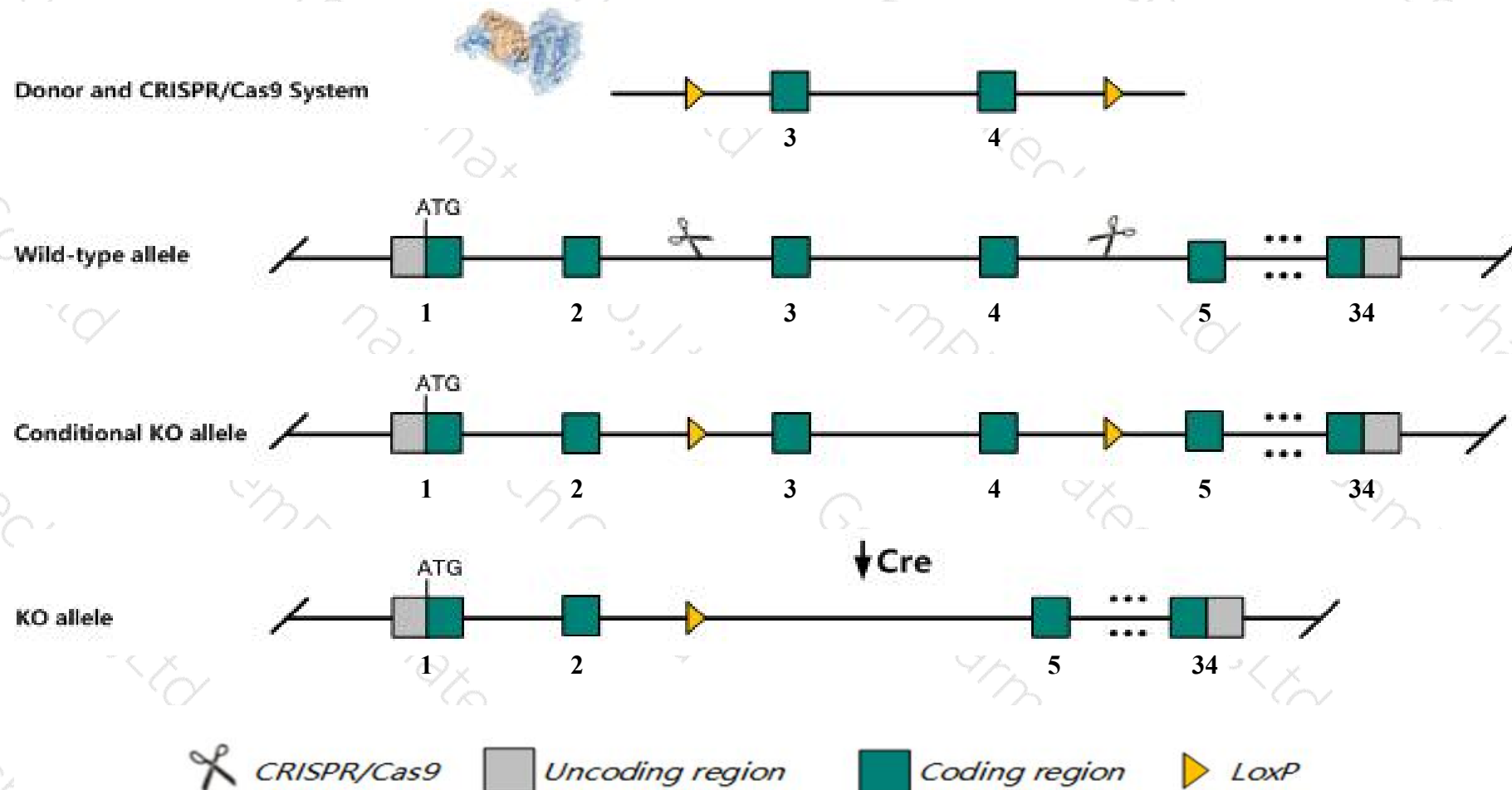
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Arfgef3* gene. The schematic diagram is as follows:



- The *Arfgef3* gene has 2 transcripts. According to the structure of *Arfgef3* gene, exon3-exon4 of *Arfgef3*-202 (ENSMUST00000215836.1) transcript is recommended as the knockout region. The region contains 214bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Arfgef3* 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, Mice homozygous for a knock-out allele exhibit increased insulin granule biogenesis and insulin secretion.
- The *Arfgef3* gene is located on the Chr10. 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)

Arfgef3 ARFGEF family member 3 [*Mus musculus* (house mouse)]

Gene ID: 215821, updated on 13-Mar-2020

Summary

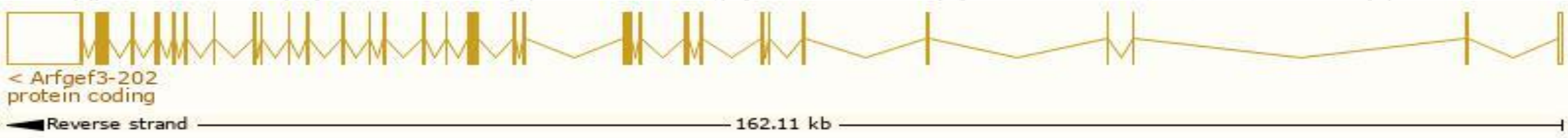
Official Symbol	Arfgef3 provided by MGI
Official Full Name	ARFGEF family member 3 provided by MGI
Primary source	MGI:MGI:106387
See related	Ensembl:ENSMUSG00000019852
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	<i>Mus musculus</i>
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	BIG3; mKIAA1244; 9630004D11; D10Bwg1379e; B930094H20Rik
Expression	Broad expression in cortex adult (RPKM 3.5), colon adult (RPKM 3.2) and 19 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

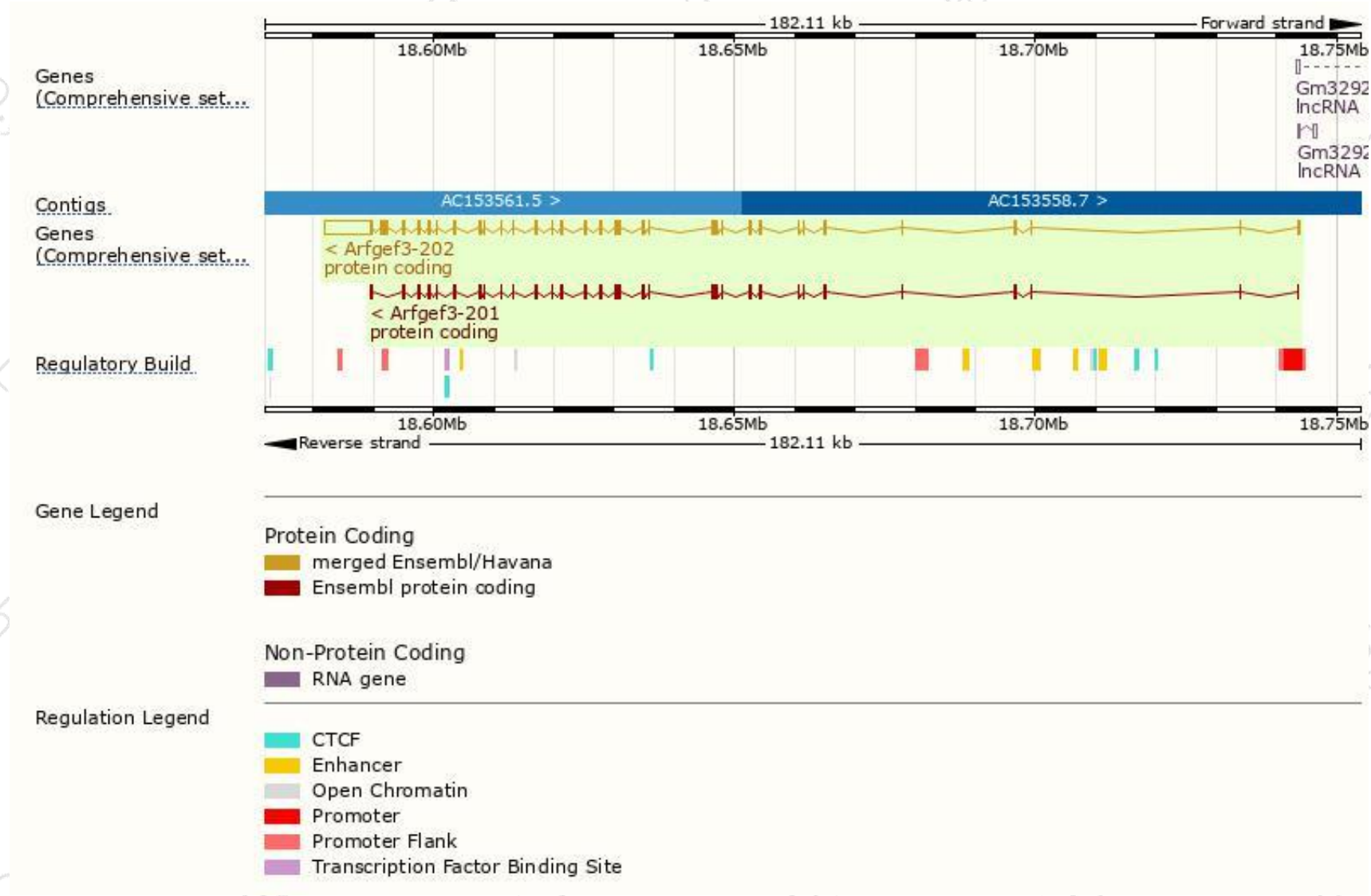
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Arfgef3-202	ENSMUST00000215836.1	14579	2170aa	Protein coding	CCDS23712	Q3UGY8	TSL:1 GENCODE basic APPRIS P2
Arfgef3-201	ENSMUST00000019999.6	5606	1709aa	Protein coding	-	A0A1N9M5I8	TSL:1 GENCODE basic APPRIS ALT2

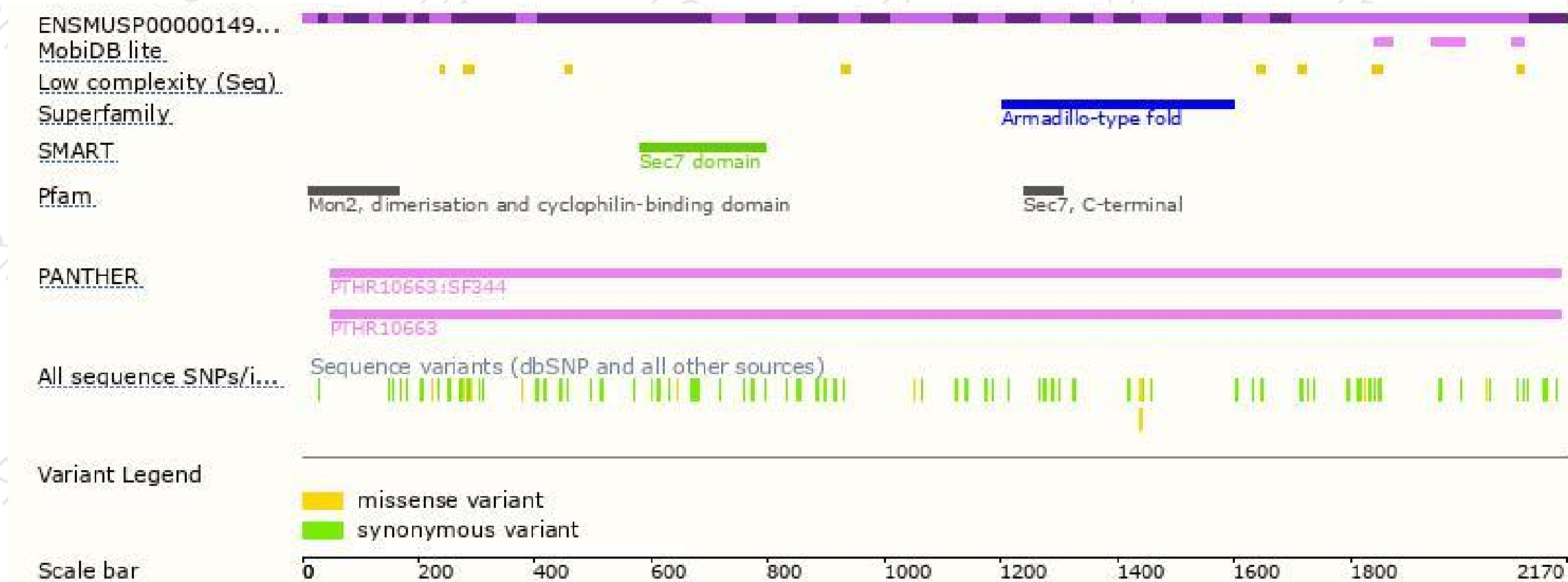
The strategy is based on the design of *Arfgef3-202* transcript,The transcription is shown below



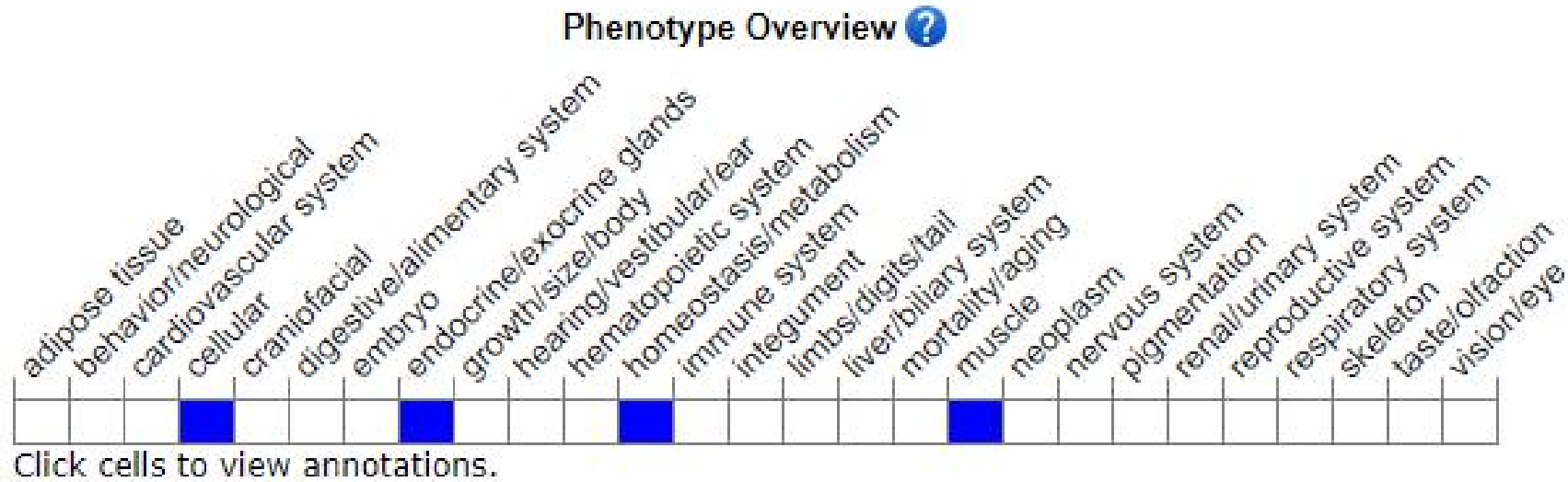
Genomic location distribution



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 increased insulin granule biogenesis and insulin secretion.

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

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