Rictor Cas9-CKO Strategy

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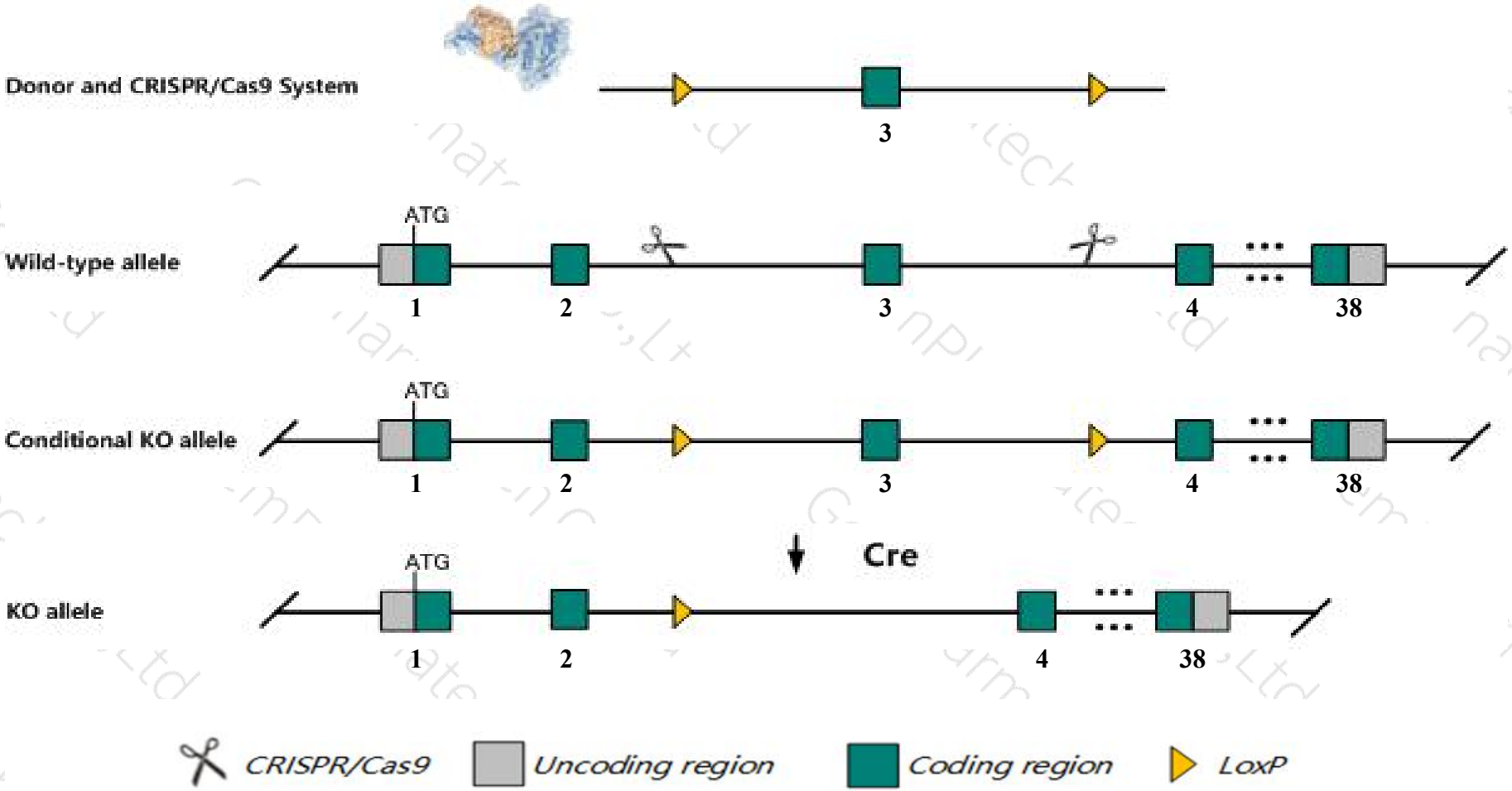
Project Overview

Project Name	***Rictor***
Project type	**Cas9-CKO**
Strain background	**C57BL/6JGpt**

Conditional Knockout strategy

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



Technical routes

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- The *Rictor* gene has 5 transcripts. According to the structure of *Rictor* gene, exon3 of *Rictor-201* (ENSMUST00000061656.7) transcript is recommended as the knockout region.The region contains 98bp coding sequence. Knock out the region will result in disruption of protein function.

- In this project we use CRISPR/Cas9 technology to modify *Rictor* 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

- According to the existing MGI data,Mice homozygous for a null allele exhibit embryonic lethality during organogenesis associated with abnormal placental morphology.

- The *Rictor* gene is located on the Chr15. 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）

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Rictor RPTOR independent companion of MTOR, complex 2 [Mus musculus (house mouse)]

Gene ID: 78757, updated on 9-Apr-2019

Summary

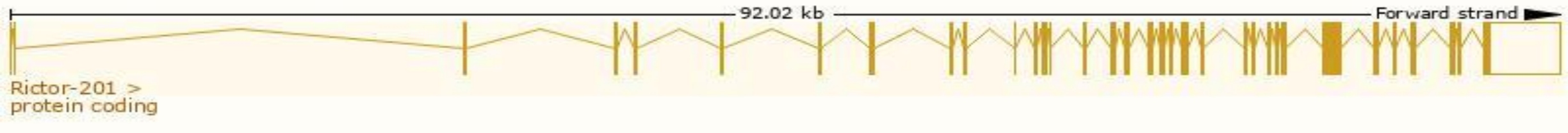
Official Symbol	Rictor provided by MGI
Official Full Name	RPTOR independent companion of MTOR, complex 2 provided by MGI
Primary source	MGI:MGI:1926007
See related	Ensembl:ENSMUSG00000050310
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	4921505C17Rik, 6030405M08Rik, AVO3, AW492497, D530039E11Rik
Expression	Ubiquitous expression in thymus adult (RPKM 3.9), cerebellum adult (RPKM 2.3) and 28 other tissues See more
Orthologs	human all

Transcript information（Ensembl）

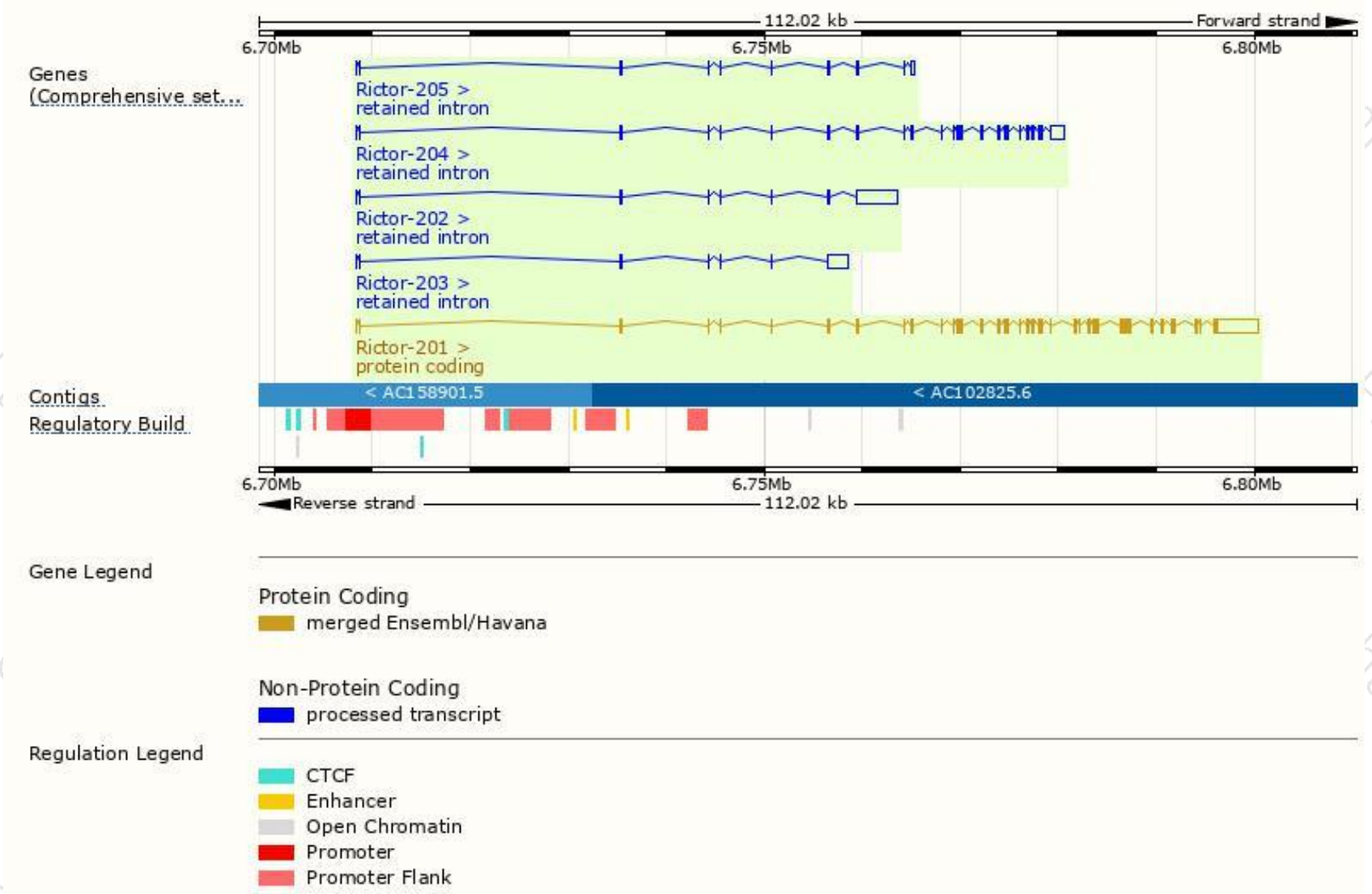
The gene has 5 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Rictor-201	ENSMUST00000061656.7	9312	1708aa	Protein coding	CCDS37032	Q6QI06	TSL:1 GENCODE basic APPRIS P1
Rictor-202	ENSMUST00000226181.1	4695	No protein	Retained intron	-	-	
Rictor-204	ENSMUST00000228266.1	3879	No protein	Retained intron	-	-	
Rictor-203	ENSMUST00000226201.1	2633	No protein	Retained intron	-	-	
Rictor-205	ENSMUST00000228918.1	1211	No protein	Retained intron	-	-	

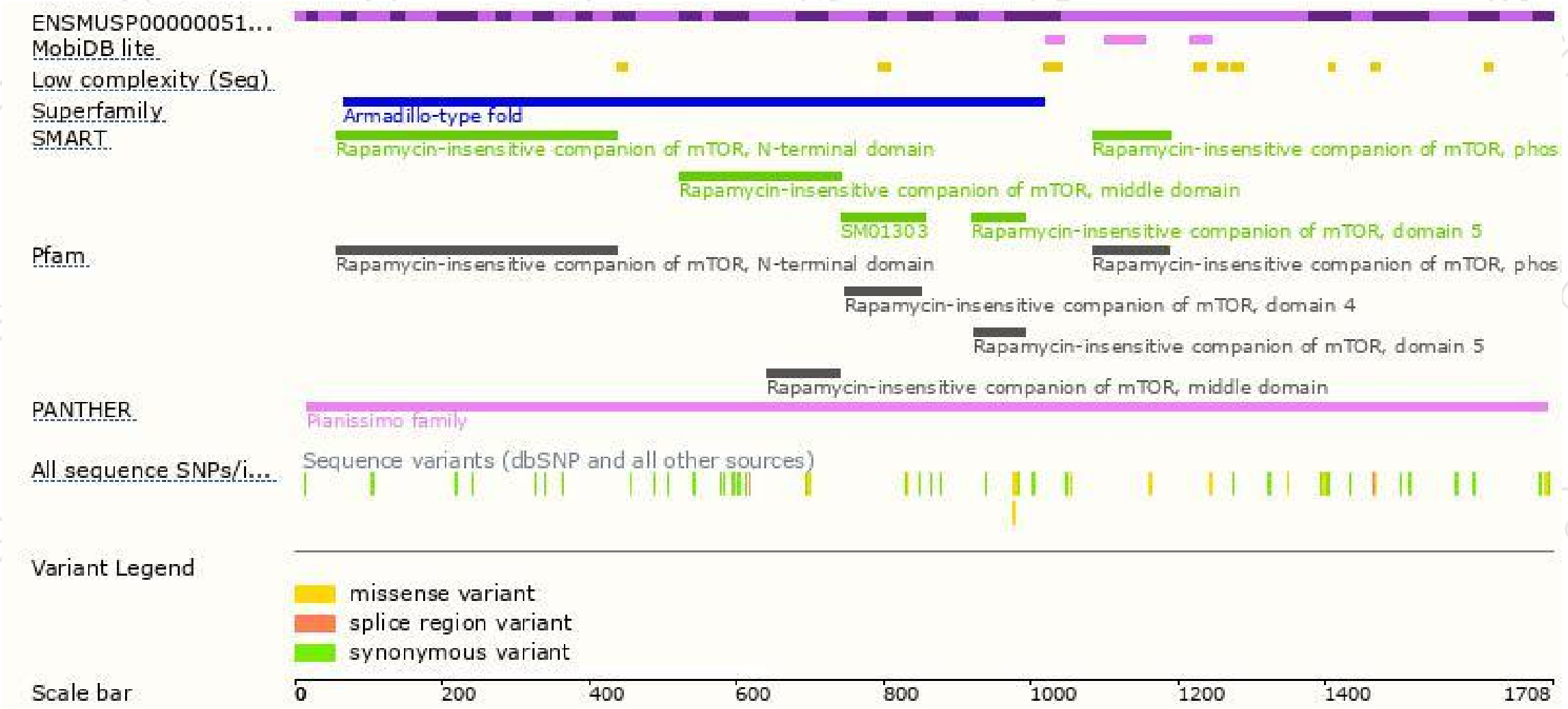
The strategy is based on the design of *Rictor-201* transcript,The transcription is shown below



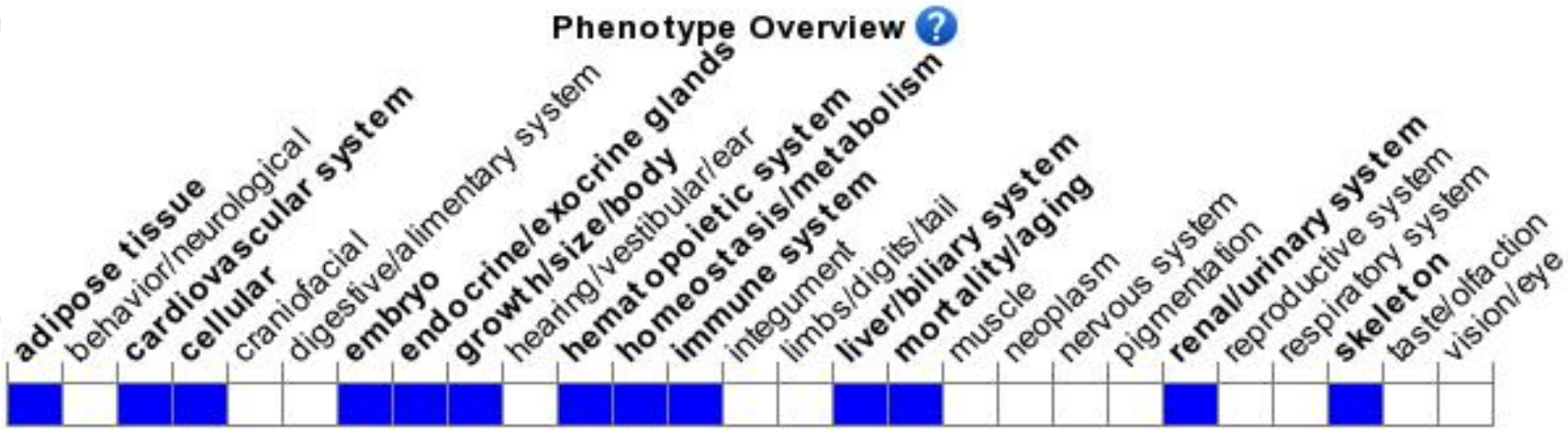
Genomic location distribution



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 null allele exhibit embryonic lethality during organogenesis associated with abnormal placental morphology.

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

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