



Dact1 Cas9-CKO Strategy

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Reviewer:

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Design Date:

2019-11-18

Project Overview

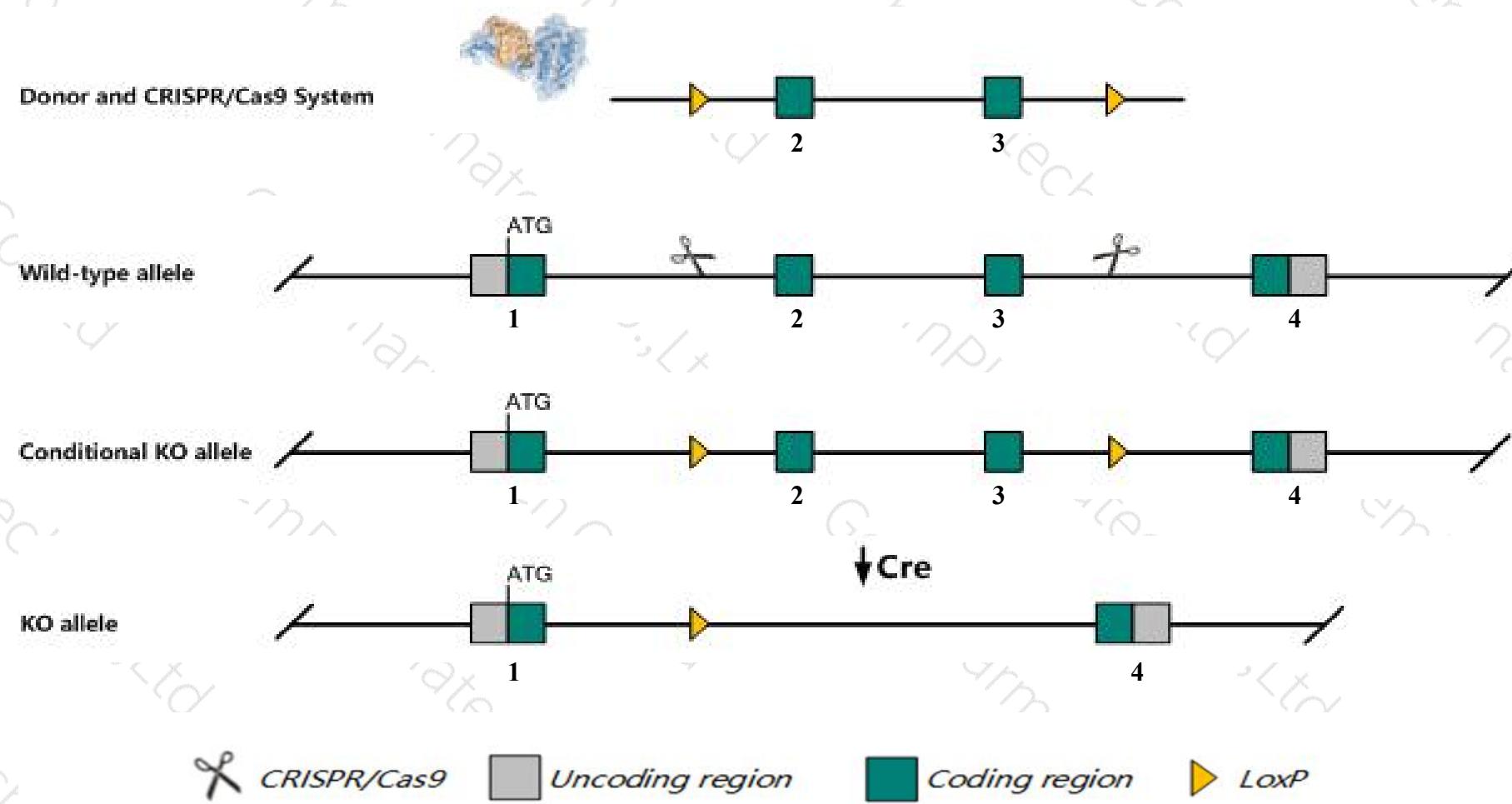
Project Name**Dact1**

Project type**Cas9-CKO**

Strain background**C57BL/6JGpt**

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Dact1* gene. The schematic diagram is as follows:



Technical routes

- The *Dactl* gene has 2 transcripts. According to the structure of *Dactl* gene, exon2-exon3 of *Dactl*-201 (ENSMUST00000061273.11) transcript is recommended as the knockout region. The region contains 289bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Dactl* 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.



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Notice

- According to the existing MGI data, Mice homozygous for a knock-out allele exhibit neonatal lethality, abnormal embryogenesis, blind-ended colons, and abnormal renal/urinary system.
- The *Dact1* gene is located on the Chr12. 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.



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Gene information (NCBI)

Dact1 dishevelled-binding antagonist of beta-catenin 1 [Mus musculus (house mouse)]

Gene ID: 59036, updated on 31-Jan-2019

Summary



Official Symbol Dact1 provided by [MGI](#)

Official Full Name dishevelled-binding antagonist of beta-catenin 1 provided by [MGI](#)

Primary source [MGI:MGI:1891740](#)

See related [Ensembl:ENSMUSG00000044548](#)

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 4921528D17Rik, AI115603, DAPPER, DAPPER1, FRODO, Frd1, Frodo1, MDpr1, MTNG3, THYEX3

Expression Broad expression in CNS E14 (RPKM 15.4), whole brain E14.5 (RPKM 14.9) and 24 other tissues [See more](#)

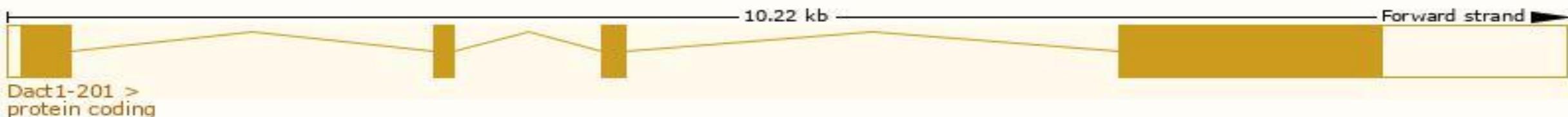
Orthologs [human](#) [all](#)

Transcript information (Ensembl)

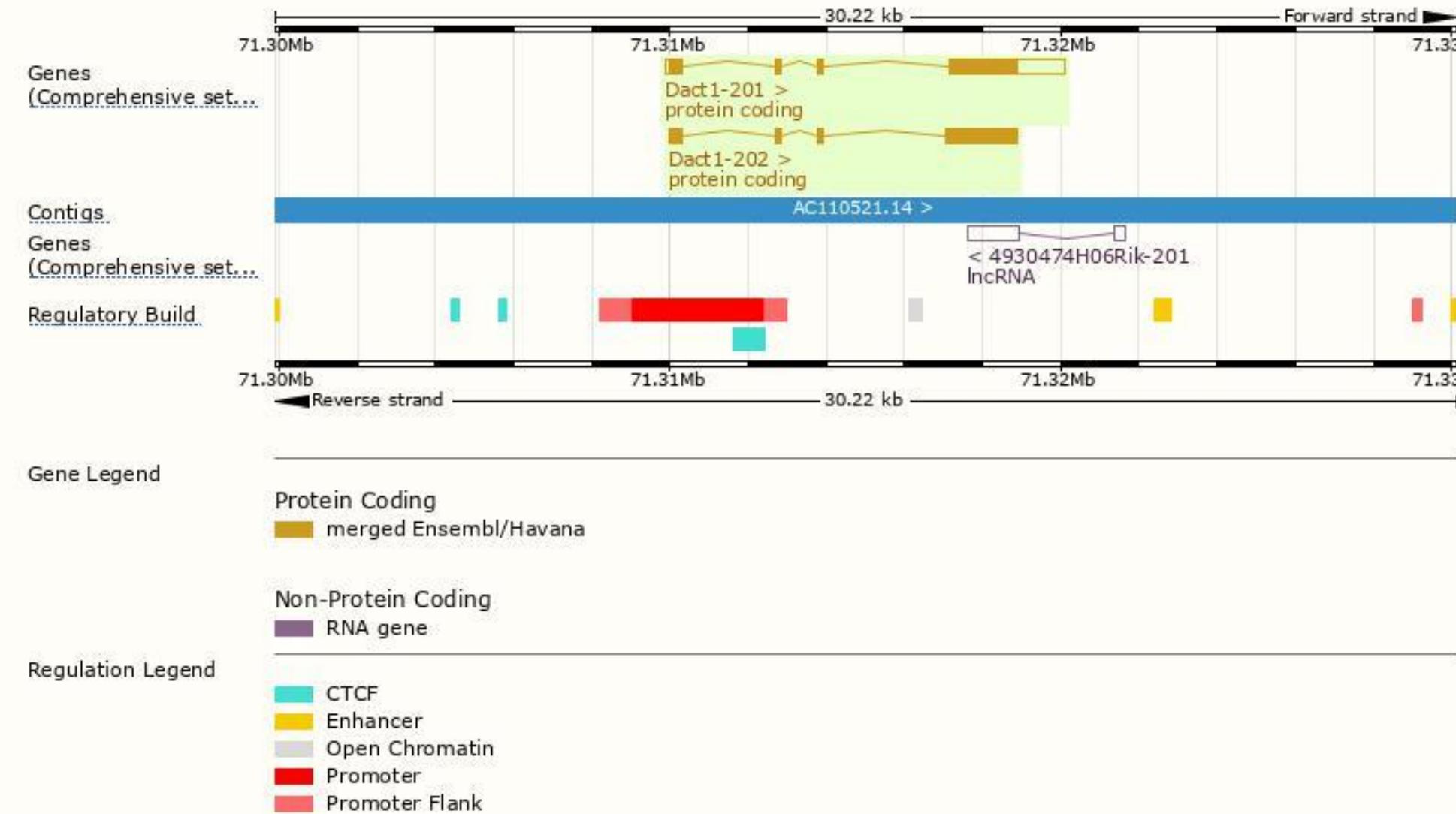
The gene has 2 transcripts, all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Dact1-201	ENSMUST00000061273.11	3650	778aa	Protein coding	CCDS25963	Q8R4A3	TSL:1 GENCODE basic
Dact1-202	ENSMUST00000150639.1	2448	815aa	Protein coding	CCDS49083	D3Z5V0	TSL:3 GENCODE basic APPRIS P1

The strategy is based on the design of *Dact1-201* transcript, The transcription is shown below



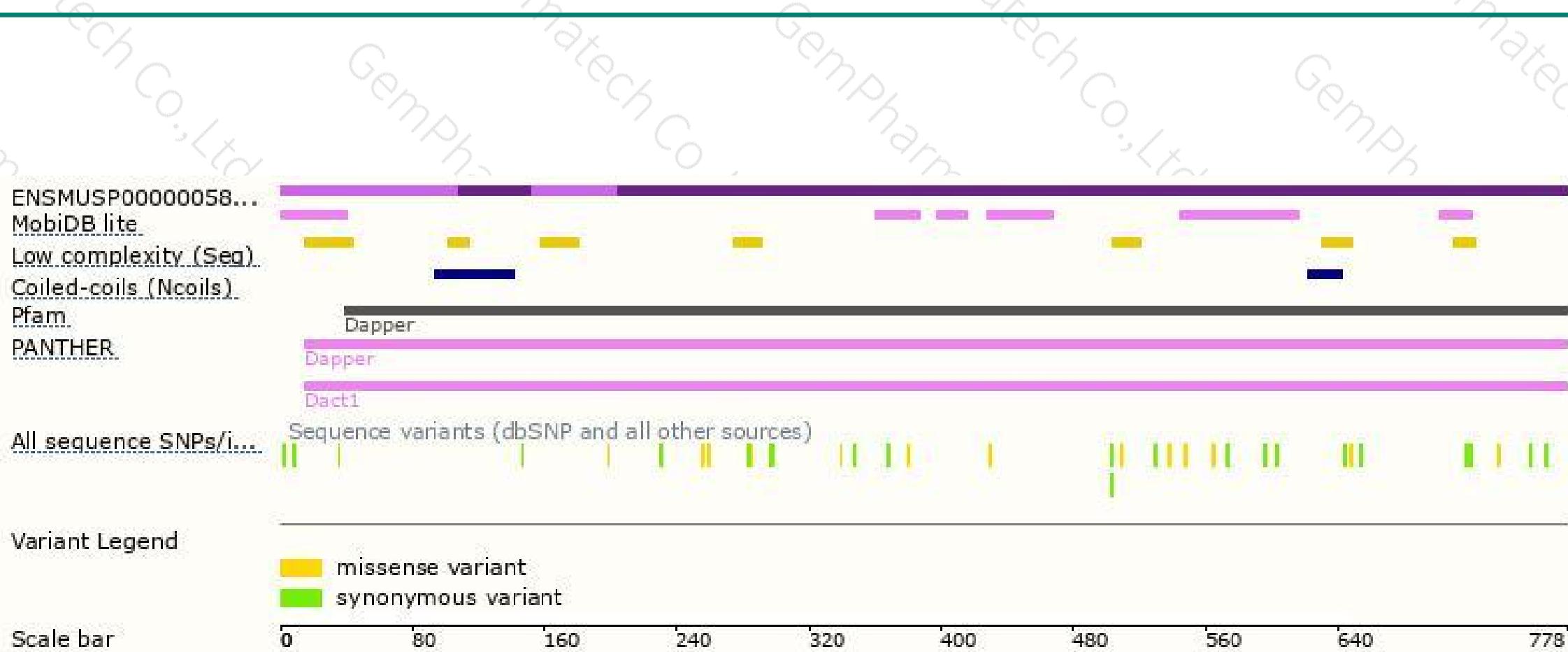
Genomic location distribution





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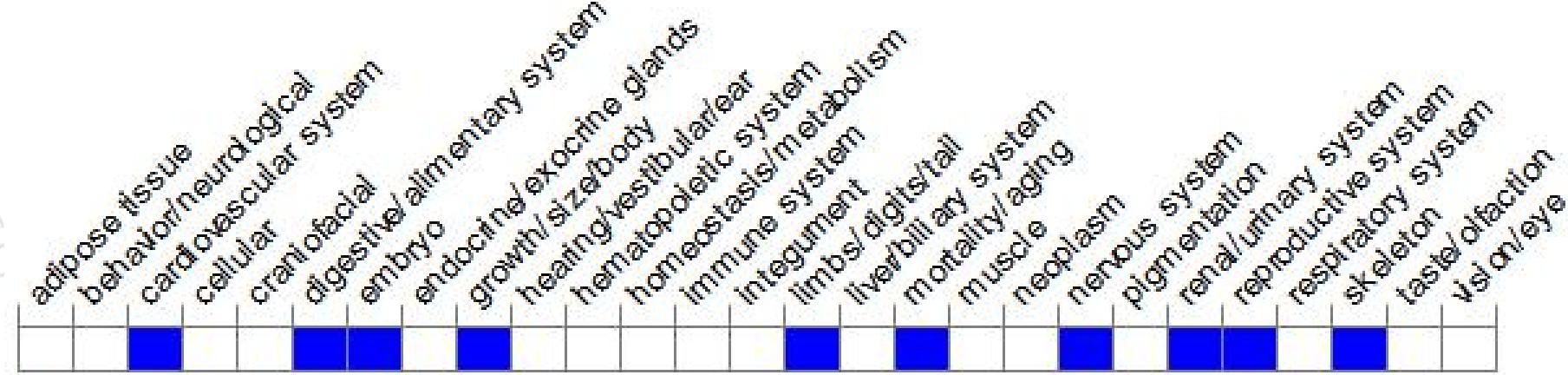
Protein domain



Mouse phenotype description(MGI)



Phenotype Overview



Click cells to view annotations

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 neonatal lethality, abnormal embryogenesis, blind-ended colons, and abnormal renal/urinary system.



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

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