

Wdr24 Cas9-CKO Strategy

Designer:

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Project Overview



Project Name

Wdr24

Project type

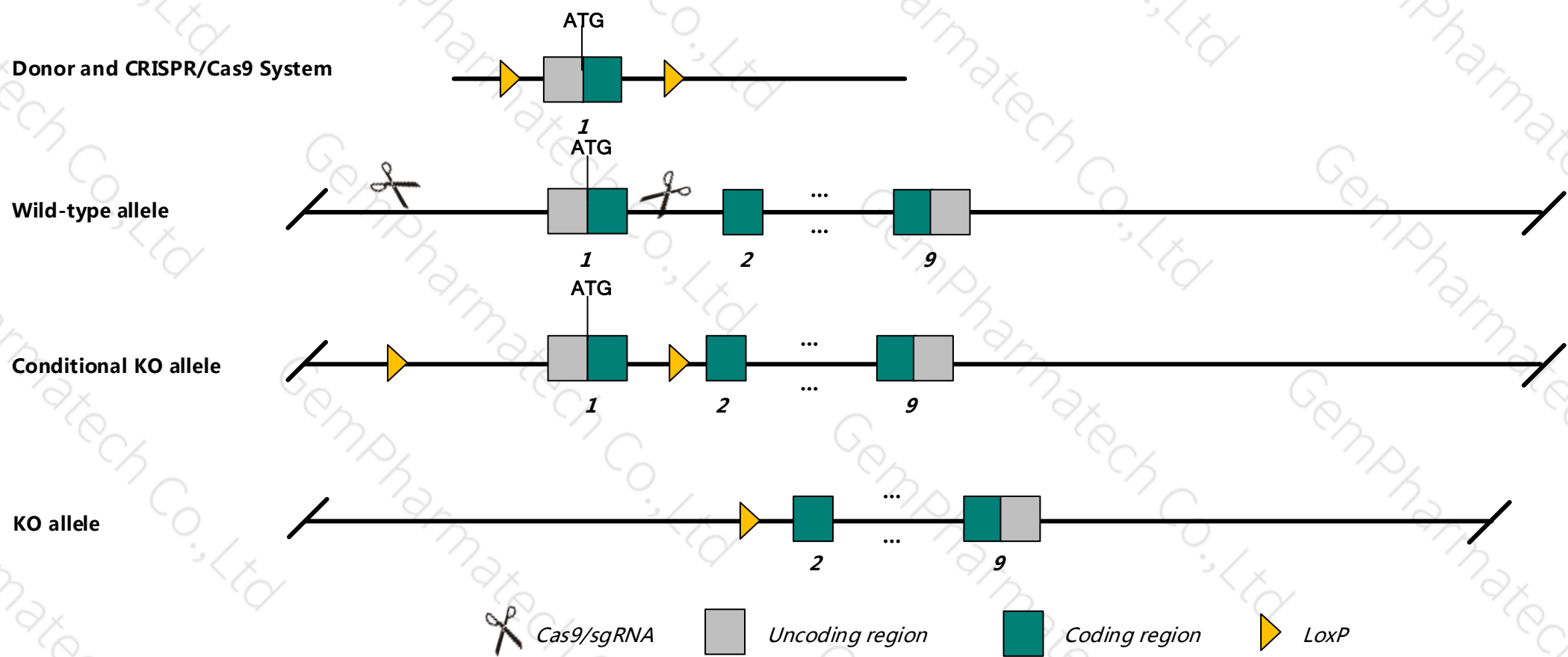
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Wdr24* gene. The schematic diagram is as follows:



- The *Wdr24* gene has 4 transcripts. According to the structure of *Wdr24* gene, exon1 of *Wdr24-201* (ENSMUST00000026833.5) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Wdr24* gene. The brief process is as follows: sgRNA was transcribed in vitro, donor vector was constructed. Cas9, sgRNA 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 was knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues or cell types.

- The KO region contains functional region of the *Fbxl16* and *Gm26694* gene. Knockout the region may affect the function of *Fbxl16* and *Gm26694* gene.
- The *Wdr24* gene is located on the Chr17. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)

Wdr24 WD repeat domain 24 [*Mus musculus* (house mouse)]

Gene ID: 268933, updated on 5-Aug-2018

Summary

Official Symbol Wdr24 provided by MGI

Official Full Name WD repeat domain 24 provided by MGI

Primary source [MGI:MGI:2446285](#)

See related [Ensembl:ENSMUSG00000025737](#) [Vega:OTTMUSG00000029399](#)

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 BC037651

Expression Ubiquitous expression in thymus adult (RPKM 26.9), ovary adult (RPKM 24.9) and 28 other tissues [See more](#)

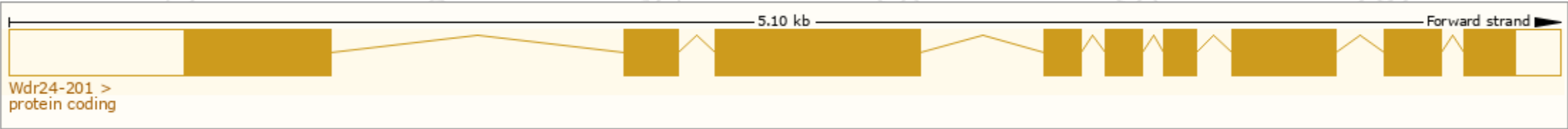
Orthologs [human](#) [all](#)

Transcript information (Ensembl)

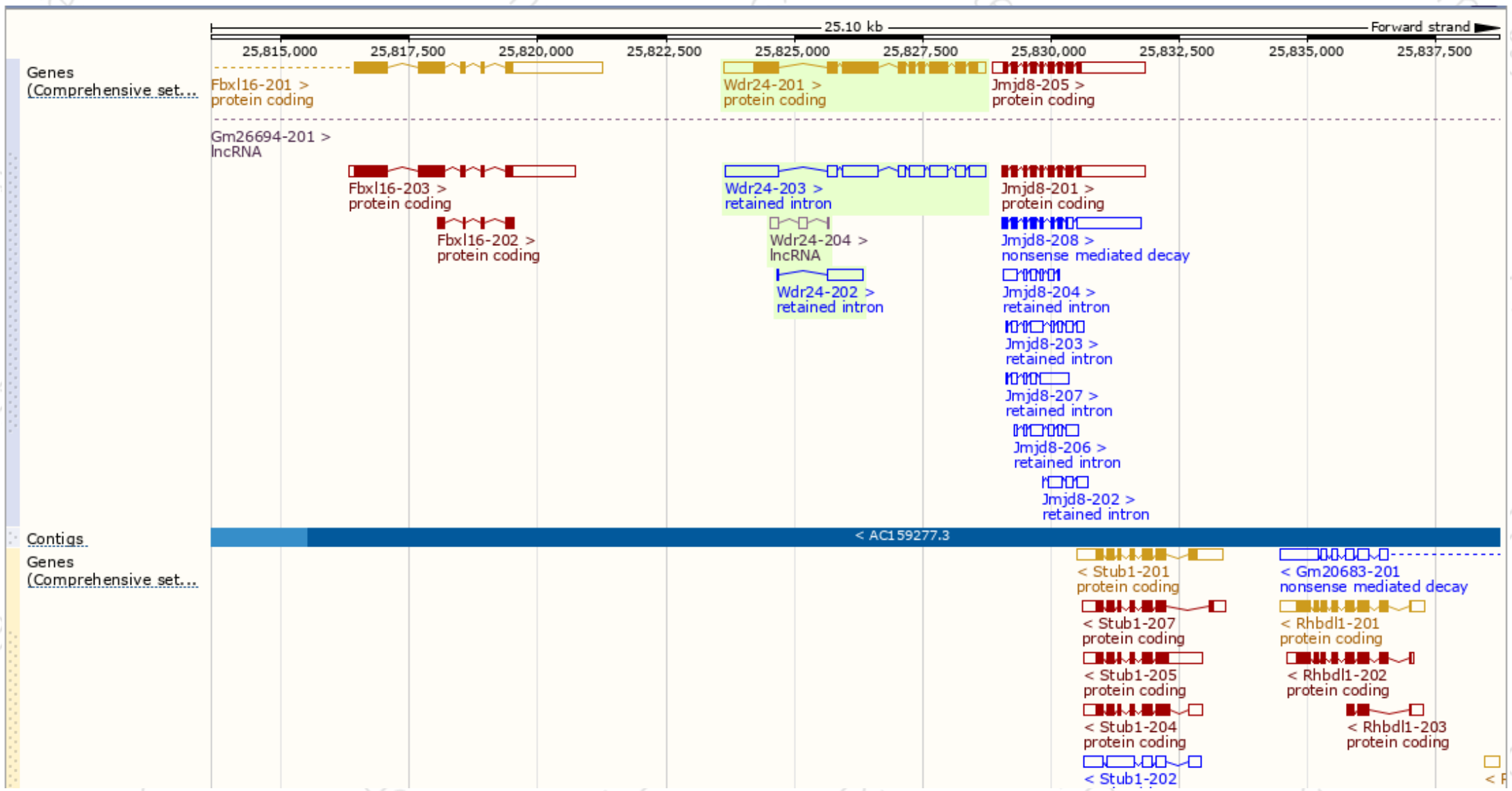
The gene has 4 transcripts, and all transcripts are shown below:

Show/hide columns (1 hidden)							Filter	
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags	
Wdr24-201	ENSMUST00000026833.5	3099	790aa	Protein coding	CCDS28531	Q8CFJ9	TSL:1	GENCODE basic APPRIS P1
Wdr24-203	ENSMUST00000160349.1	3139	No protein	Retained intron	-	-	TSL:1	
Wdr24-202	ENSMUST00000160275.1	716	No protein	Retained intron	-	-	TSL:3	
Wdr24-204	ENSMUST00000160829.1	317	No protein	lncRNA	-	-	TSL:5	

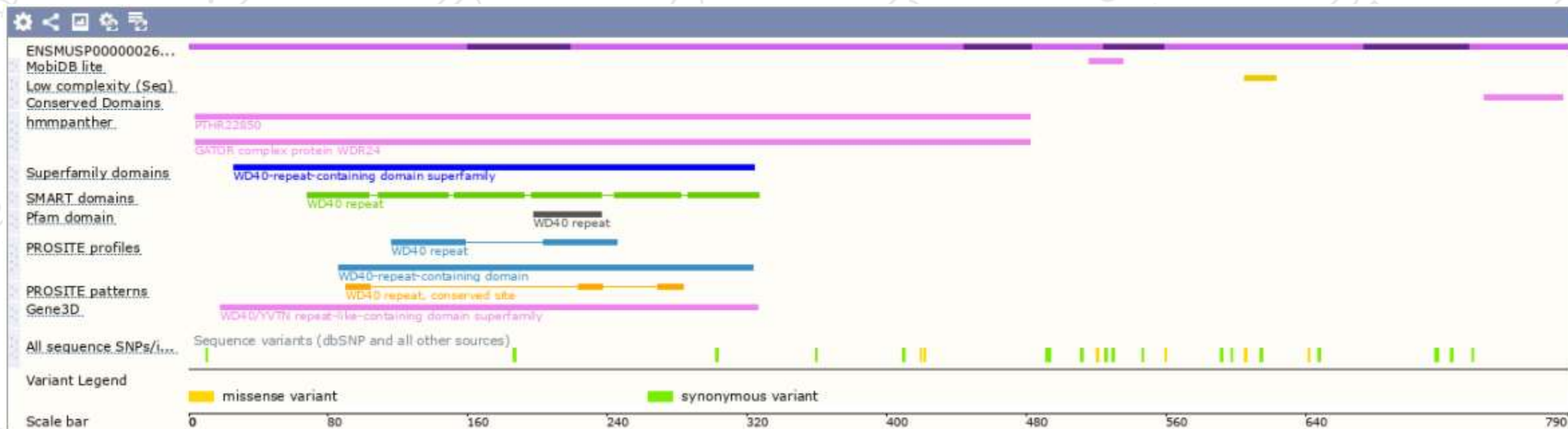
The strategy is based on the design of Wdr24-201 transcript, The transcription is shown below



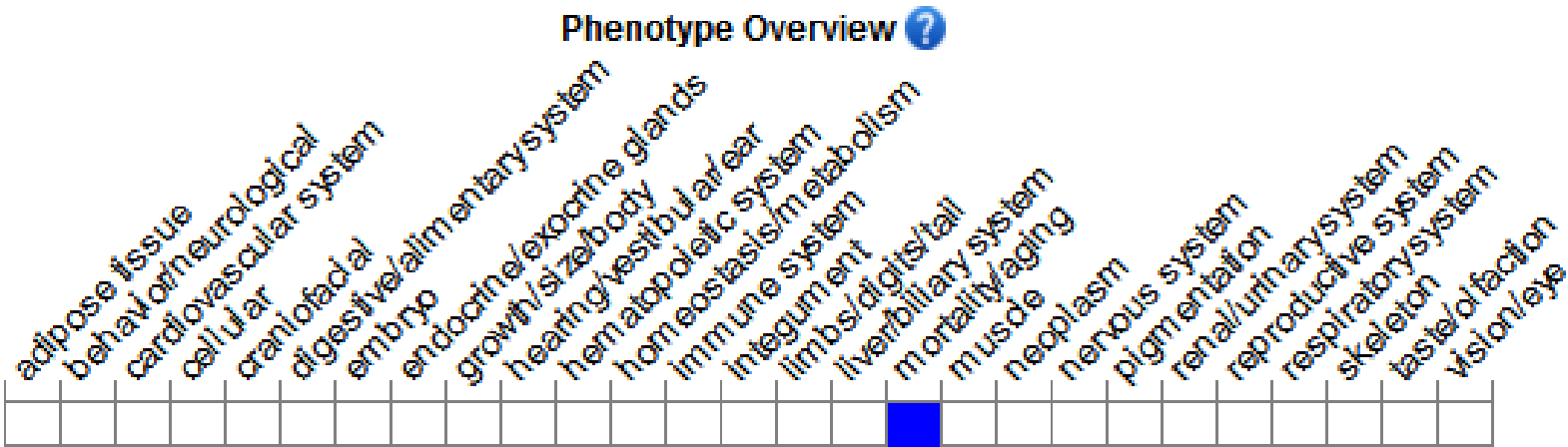
Genomic location distribution



Protein domain



Mouse phenotype description(MGI)



Click cells to view annotations.

Phenotypes affected by the gene are marked in blue.Data quoted from MGI database(<http://www.informatics.jax.org/>) .

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