Cebpe Cas9-KO Strategy

Designer: Yang Zeng
Reviewer: Daohua Xu

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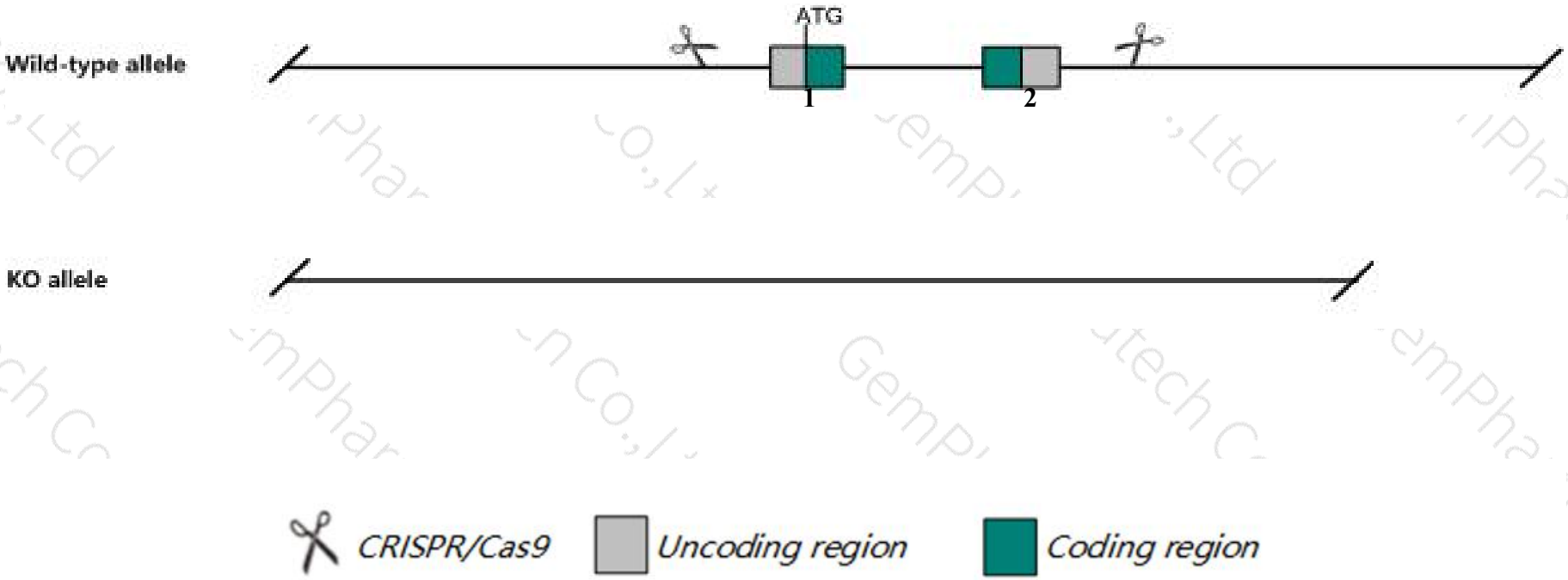
Project Overview

Project Name	***Cebpe***
Project type	**Cas9-KO**
Strain background	**C57BL/6JGpt**

Knockout strategy

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



Technical routes

➢ The *Cebpe* gene has 1 transcript. According to the structure of *Cebpe* gene, exon1-exon2 of *Cebpe-201* (ENSMUST00000064290.7) transcript is recommended as the knockout region.The region contains all of the coding sequence.Knock out the region will result in disruption of protein function.

➢ In this project we use CRISPR/Cas9 technology to modify *Cebpe* gene. The brief process is as follows: CRISPR/Cas9 system

Notice

- According to the existing MGI data,Homozygous mutation of this gene results in impaired neutrophil and eosinophil development and myelodysplasia. Mutant animals are susceptible to secondary bacterial infections such as conjuntivitis, rhinitis, and pneumonia, and become moribund between 2-5 months of age.

- The *Cebpe* gene is located on the Chr14. 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 gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information （NCBI）

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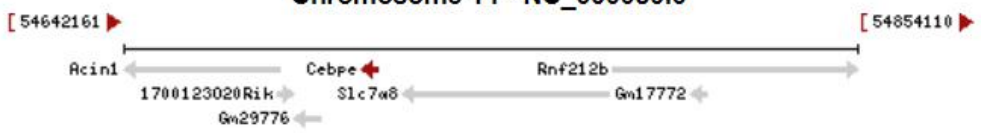
Cebpe CCAAT/enhancer binding protein (C/EBP), epsilon [*Mus musculus* (house mouse)]

Gene ID: 110794, updated on 10-Oct-2019

Summary

Official Symbol	Cebpe provided by MGI
Official Full Name	CCAAT/enhancer binding protein (C/EBP), epsilon provided by MGI
Primary source	MGI:MGI:103572
See related	Ensembl:ENSMUSG00000052435
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	CRP1; Gm294; C/EBPe; C/EBPepsilon
Expression	Biased expression in liver E18 (RPKM 15.2), liver adult (RPKM 4.7) and 9 other tissues See more
Orthologs	human all

Chromosome 14 - NC_000080.6

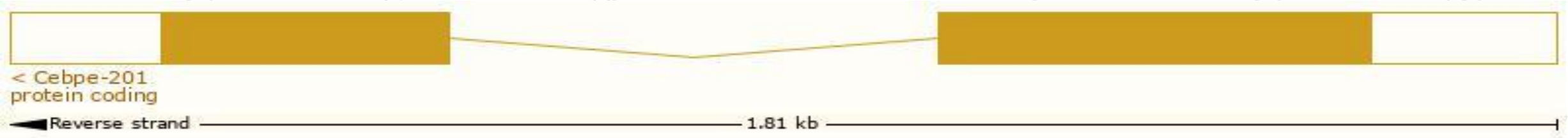


Transcript information（Ensembl）

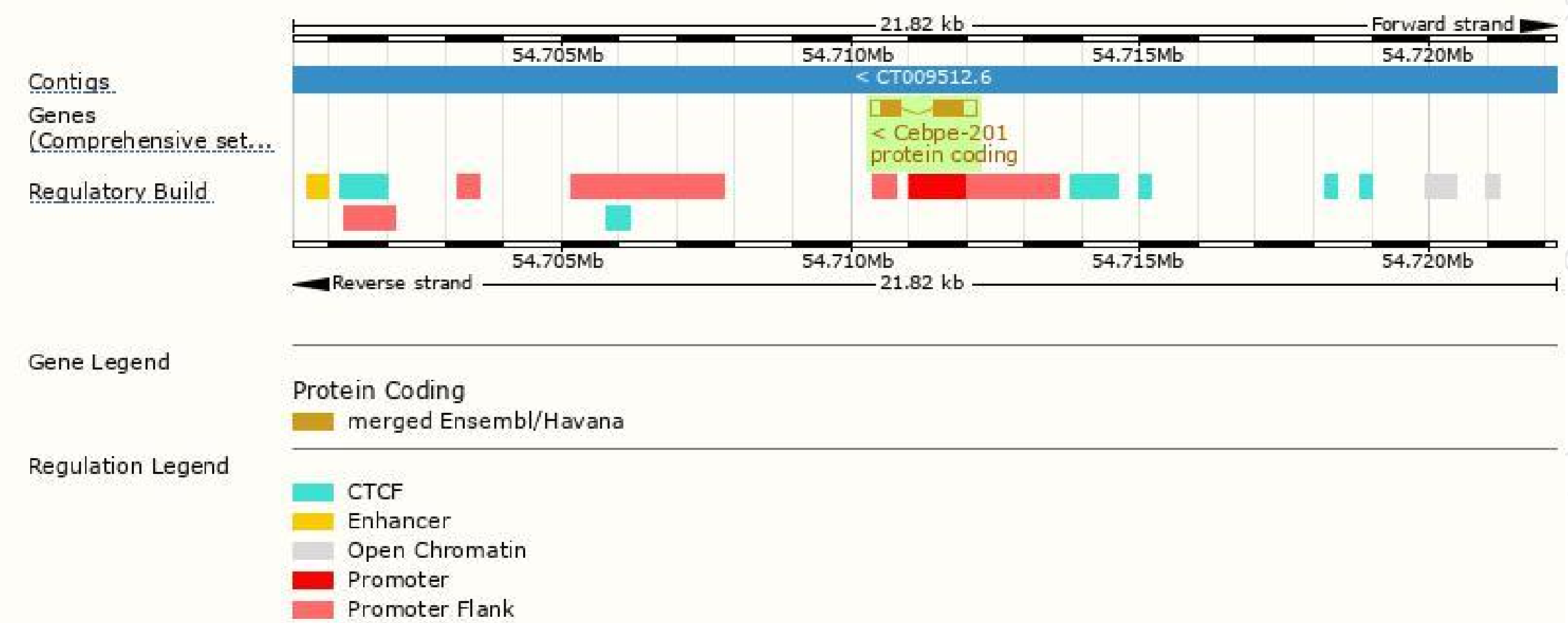
The gene has 1 transcript,and the transcript is shown below:

Name	Transcript ID	bp	Protein	Translation ID	Biotype	CCDS	UniProt	Flags
Cebpe-201	ENSMUST00000064290.7	1241	281aa	ENSMUSP00000068927.6	Protein coding	CCDS27100	Q6PZD9	TSL:1 GENCODE basic APPRIS P1

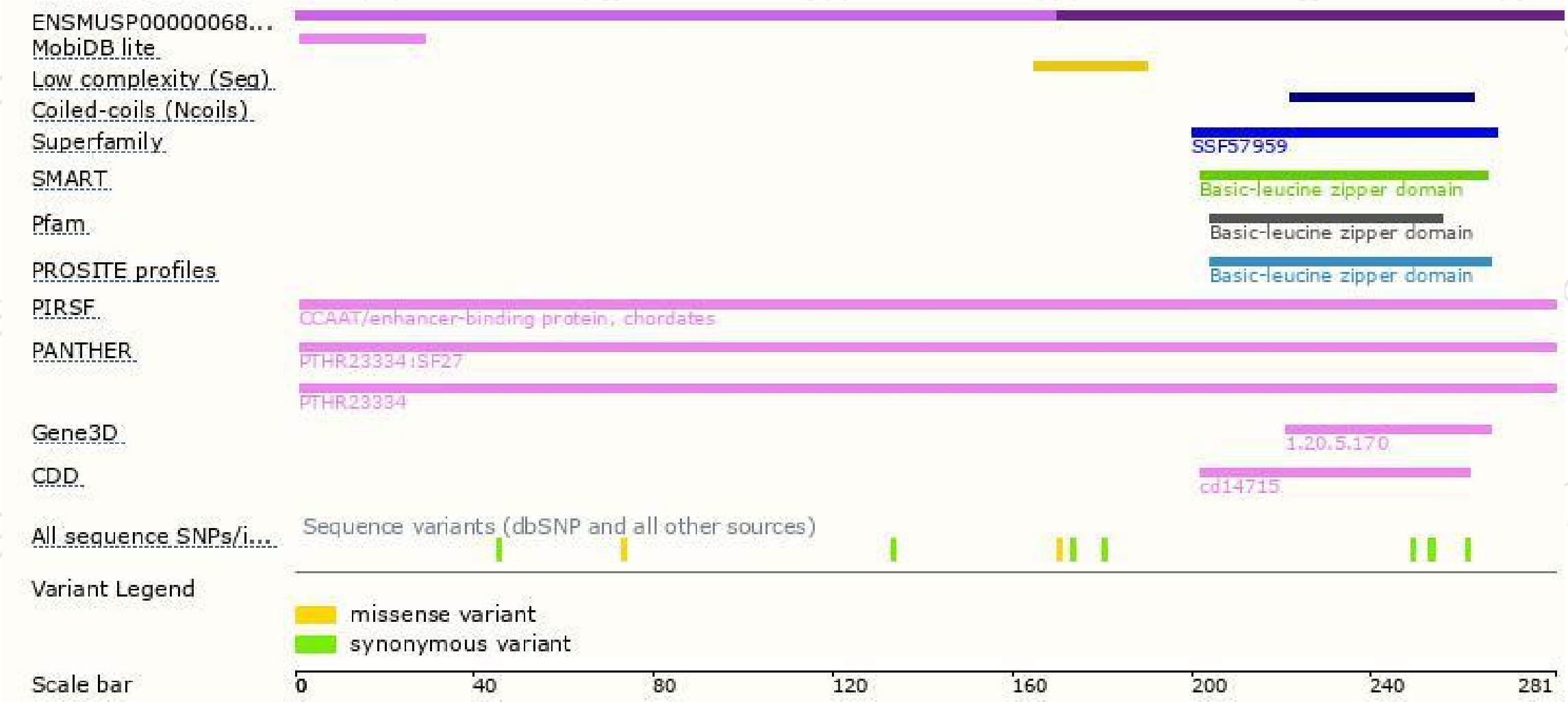
The strategy is based on the design of *Cebpe-201* transcript,The transcription is shown below



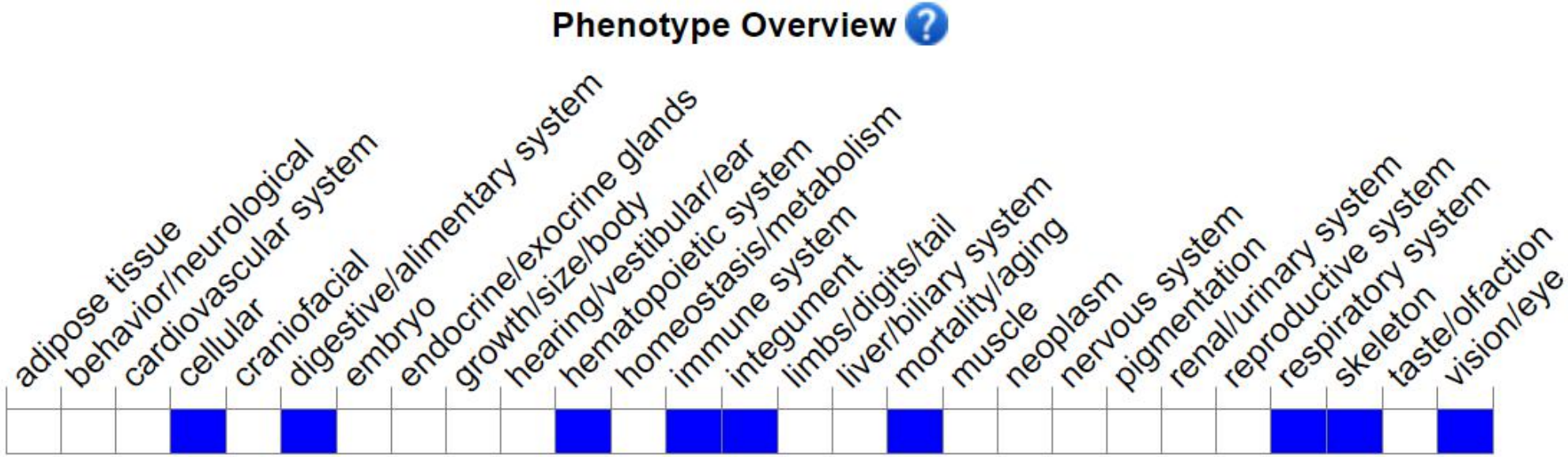
Genomic location distribution



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/).

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集萃药康
GemPharmatech

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

Tel: 400-9660890

集萃药康生物科技
GemPharmatech Co.,Ltd