

Cebpg Cas9-KO Strategy

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Project Overview

Project Name

Cebpg

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

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



- The *Cebpg* gene has 4 transcripts. According to the structure of *Cebpg* gene, exon2 of *Cebpg-202* (ENSMUST00000130491.2) 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 *Cebpg* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Homozygous mutation of this gene results in impaired natural killer cell cytotoxic activity and IFN-gamma production.
- The *Cebpg* gene is located on the Chr7. 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)

Cebpg CCAAT/enhancer binding protein (C/EBP), gamma [Mus musculus (house mouse)]

Gene ID: 12611, updated on 31-Jan-2019

Summary



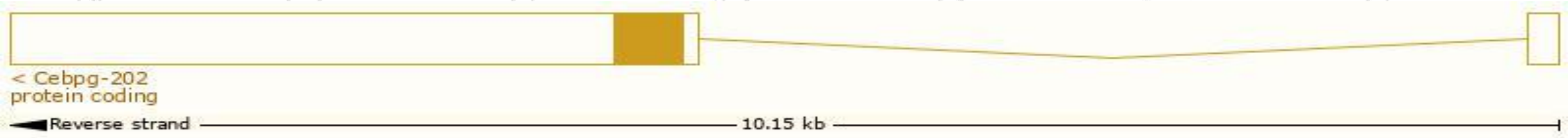
Official Symbol	Cebpg provided by MGI
Official Full Name	CCAAT/enhancer binding protein (C/EBP), gamma provided by MGI
Primary source	MGI:MGI:104982
See related	Ensembl:ENSMUSG00000056216
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	C/EBP[g], C/EBPgamma, GPE1-BP, Gpe1bp, Ig/EBP
Expression	Ubiquitous expression in liver E18 (RPKM 10.2), CNS E11.5 (RPKM 8.7) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

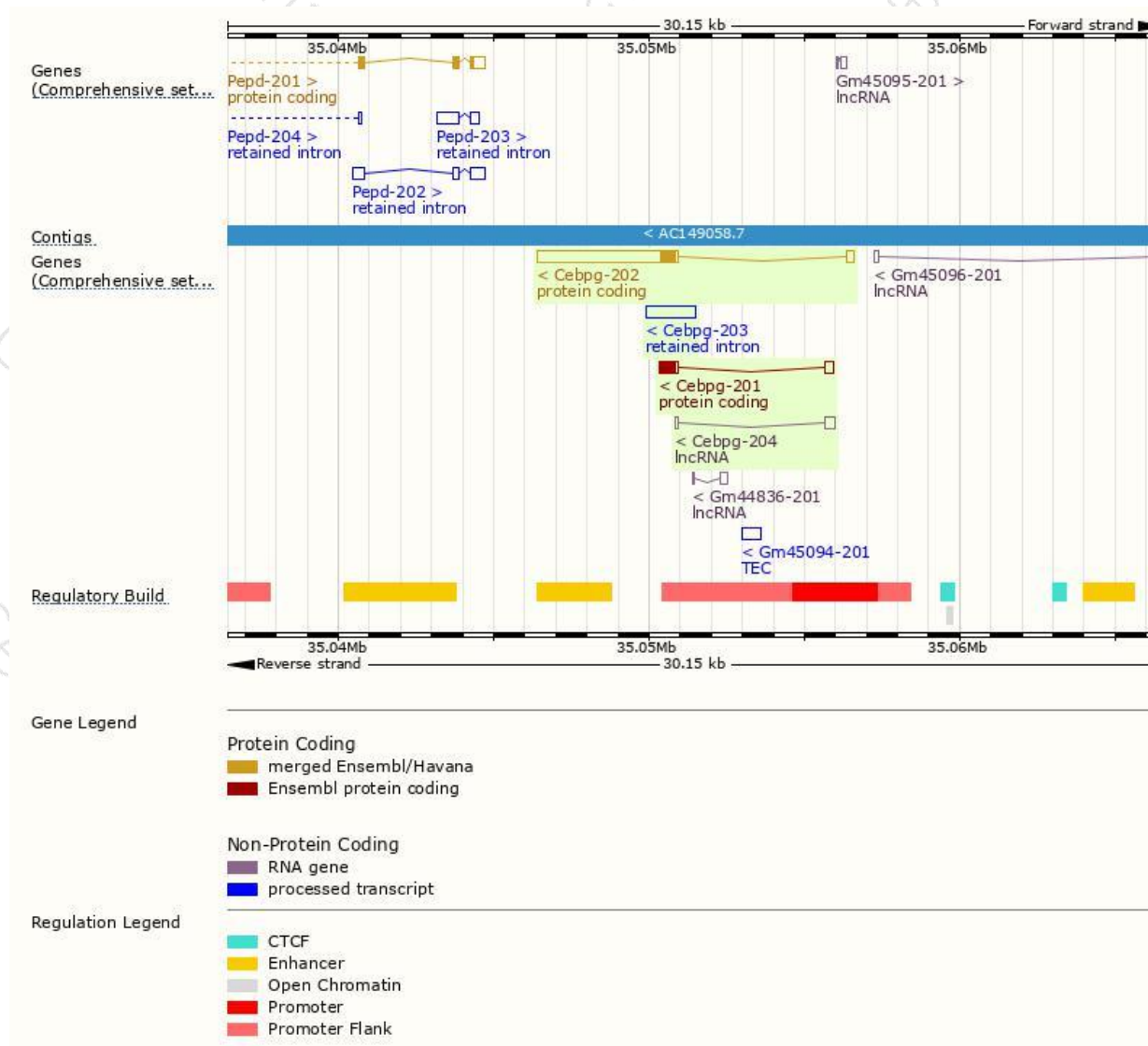
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Cebpg-202	ENSMUST00000130491.2	4709	150aa	Protein coding	CCDS21144	P53568 Q54AJ1	TSL:1 GENCODE basic APPRIS P1
Cebpg-201	ENSMUST00000070191.9	914	150aa	Protein coding	CCDS21144	P53568 Q54AJ1	TSL:2 GENCODE basic APPRIS P1
Cebpg-203	ENSMUST00000205752.1	1610	No protein	Retained intron	-	-	TSL:NA
Cebpg-204	ENSMUST00000206505.1	399	No protein	lncRNA	-	-	TSL:3

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



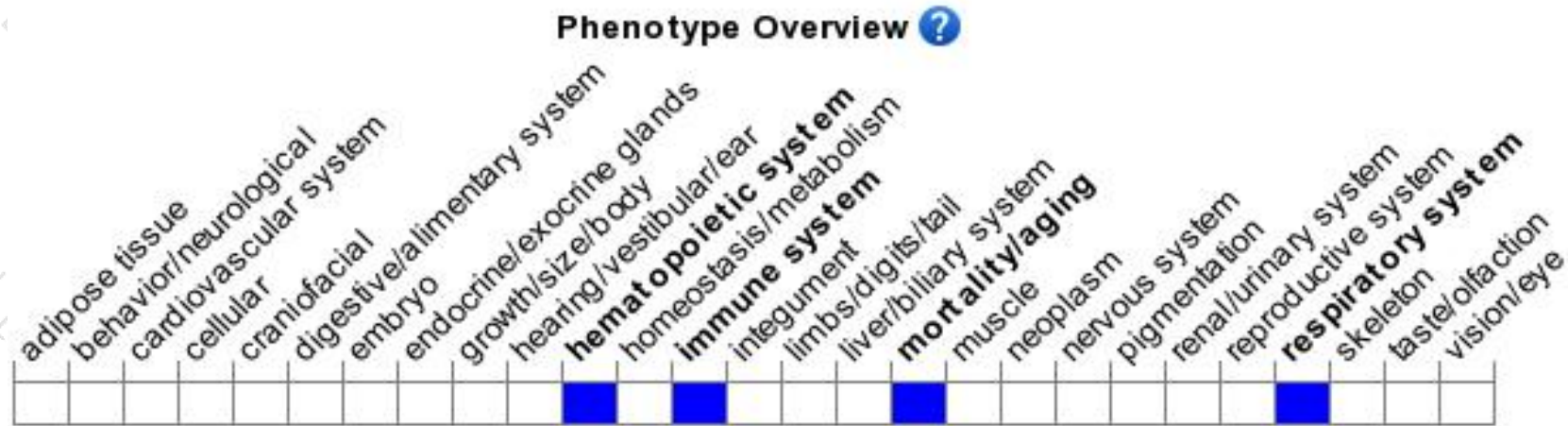
Genomic location distribution



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, Homozygous mutation of this gene results in impaired natural killer cell cytotoxic activity and IFN-gamma production.

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

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