

Casp14 Cas9-KO Strategy

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Project Overview

Project Name

Casp14

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Casp14* gene. The schematic diagram is as follows:



- The *Casp14* gene has 2 transcripts. According to the structure of *Casp14* gene, exon3-exon4 of *Casp14-201* (ENSMUST00000005488.8) transcript is recommended as the knockout region. The region contains 376bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Casp14* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice homozygous for a null allele exhibit impaired skin barrier function, skin dehydration and increased damage in response to UVB irradiation.
- The *Casp14* gene is located on the Chr10. 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)

Casp14 caspase 14 [*Mus musculus* (house mouse)]

Gene ID: 12365, updated on 13-Aug-2019

Summary



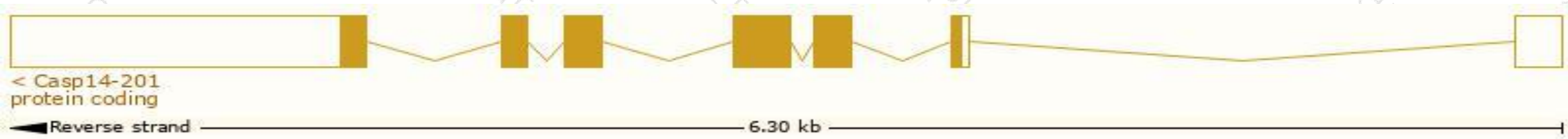
| | |
|--------------------|---|
| Official Symbol | Casp14 provided by MGI |
| Official Full Name | caspase 14 provided by MGI |
| Primary source | MGI:MGI:1335092 |
| See related | Ensembl:ENSMUSG000000005355 |
| Gene type | protein coding |
| RefSeq status | VALIDATED |
| Organism | Mus musculus |
| Lineage | Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus |
| Also known as | MICE; mini-ICE |
| Expression | Biased expression in stomach adult (RPKM 3.3), colon adult (RPKM 1.2) and 2 other tissues See more |
| Orthologs | human all |

Transcript information (Ensembl)

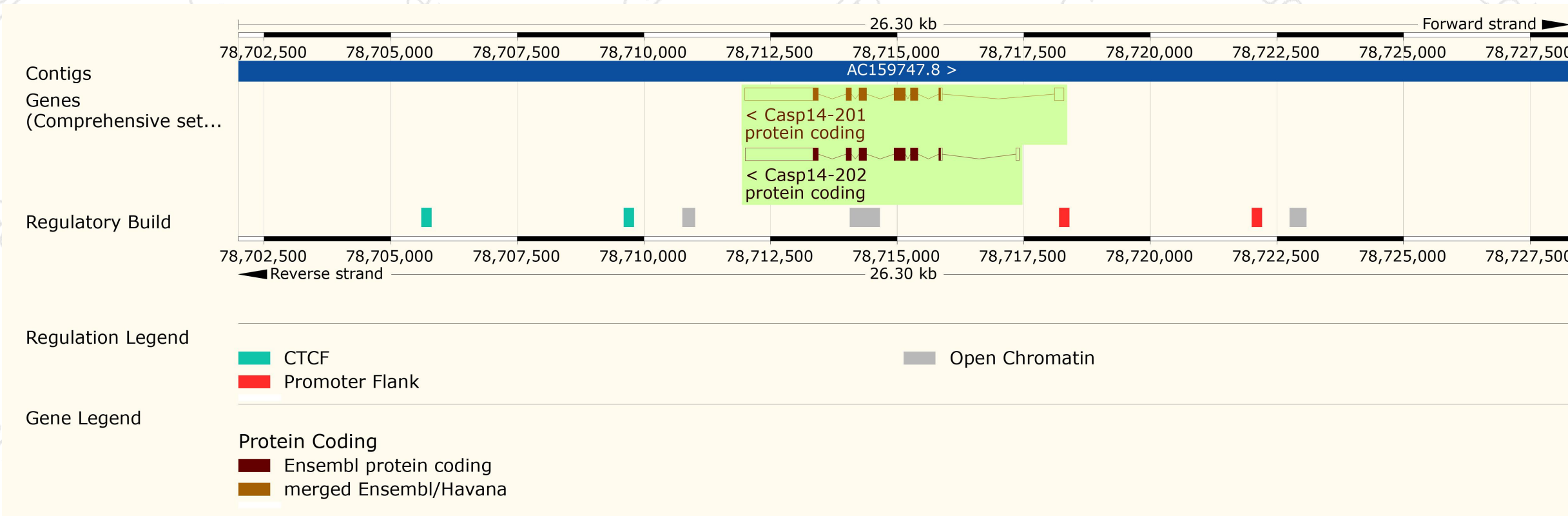
The gene has 2 transcripts,all transcripts are shown below:

| Name | Transcript ID | bp | Protein | Biotype | CCDS | UniProt | Flags |
|------------|--------------------------------------|------|-----------------------|----------------|---------------------------|-------------------------------|-------------------------------|
| Casp14-201 | ENSMUST00000005488.8 | 2336 | 257aa | Protein coding | CCDS23969 | O89094 Q542Q1 | TSL:1 GENCODE basic APPRIS P1 |
| Casp14-202 | ENSMUST00000219237.1 | 2200 | 257aa | Protein coding | CCDS23969 | O89094 Q542Q1 | TSL:5 GENCODE basic APPRIS P1 |

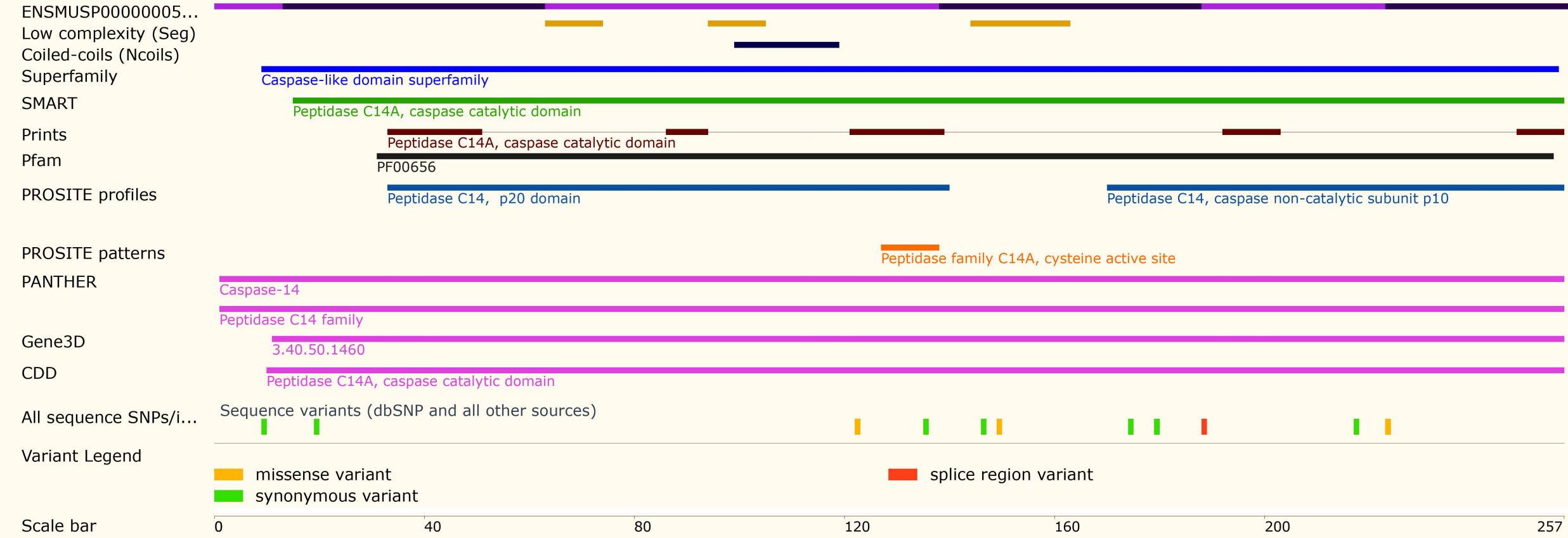
The strategy is based on the design of *Casp14-201* transcript,The transcription is shown below



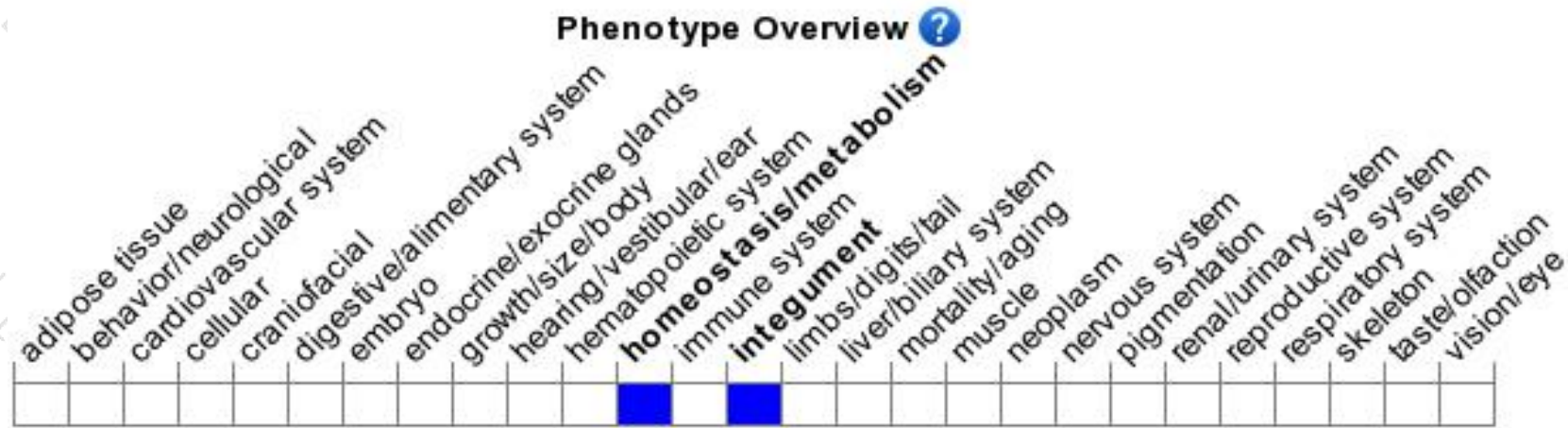
Genomic location distribution



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 impaired skin barrier function, skin dehydration and increased damage in response to UVB irradiation.

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

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