

Cxcl14 Cas9-KO Strategy

Designer:

Daohua Xu

Reviewer:

Huimin Su

Design Date:

2019-9-16

Project Overview

Project Name

Cxcl14

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Cxcl14* gene. The schematic diagram is as follows:



- The *Cxcl14* gene has 2 transcripts. According to the structure of *Cxcl14* gene, exon1-exon4 of *Cxcl14-201* (ENSMUST00000021970.10) 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 *Cxcl14* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Homozygous null mutant cause partial lethality before weaning, however surviving null are small but fertile and show insulin-sensitive phenotype in female under high fat diet feeding condition.
- The *Cxcl14* gene is located on the Chr13. 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)

Cxcl14 chemokine (C-X-C motif) ligand 14 [Mus musculus (house mouse)]

Gene ID: 57266, updated on 19-Feb-2019

Summary



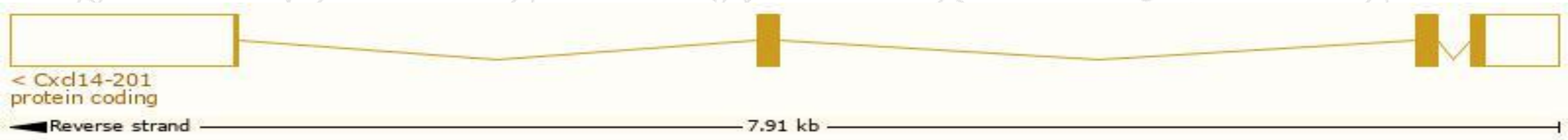
Official Symbol	Cxcl14 provided by MGI
Official Full Name	chemokine (C-X-C motif) ligand 14 provided by MGI
Primary source	MGI:MGI:1888514
See related	Ensembl:ENSMUSG000000021508
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	1110031L23Rik, 1200006I23Rik, AI414372, BMAC, BRAK, KS1, Kec, MIP-2g, MIP2gamma, NJAC, Scyb14, bolekin
Expression	Broad expression in limb E14.5 (RPKM 108.6), bladder adult (RPKM 100.8) and 19 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

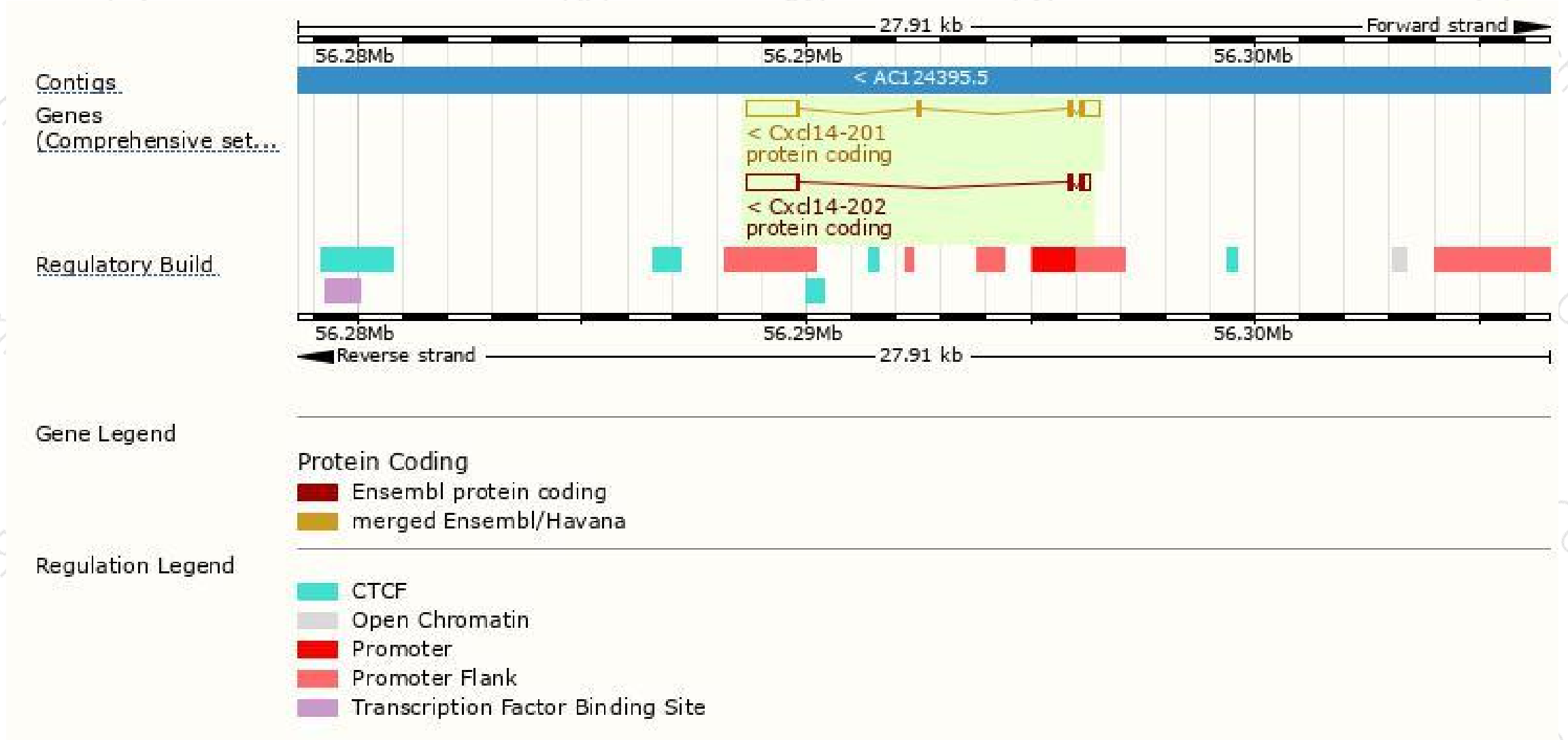
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Cxcl14-201	ENSMUST00000021970.10	1827	99aa	Protein coding	CCDS36679	Q9WUQ5	TSL:1 GENCODE basic APPRIS P1
Cxcl14-202	ENSMUST00000224801.1	1467	61aa	Protein coding	-	Q6AXC2	GENCODE basic

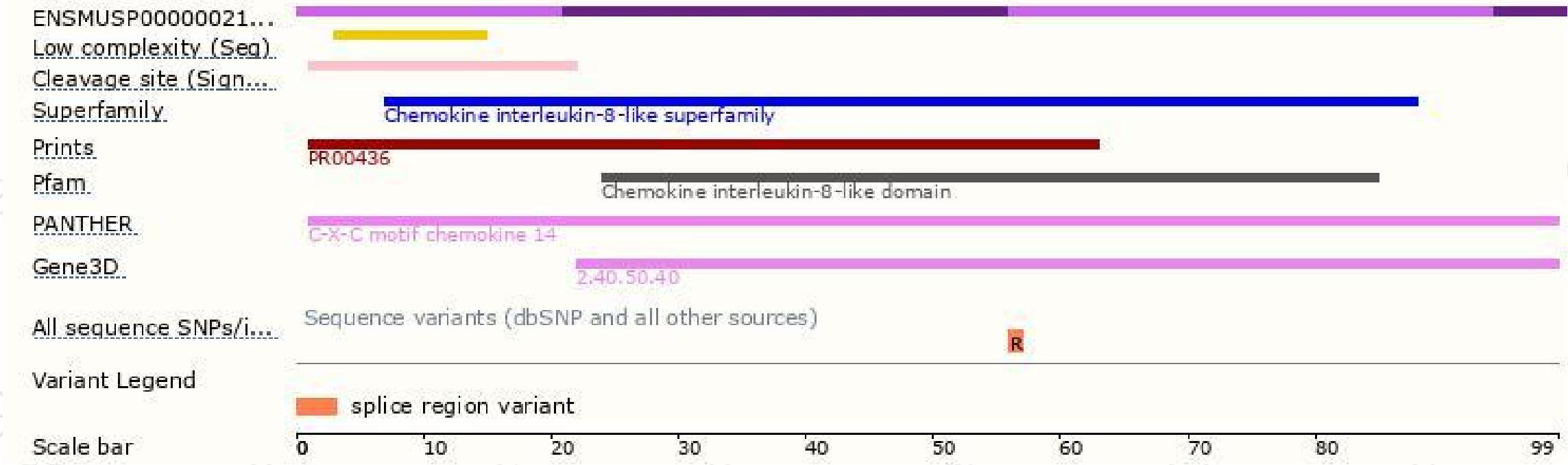
The strategy is based on the design of *Cxcl14-201* transcript,The transcription is shown below



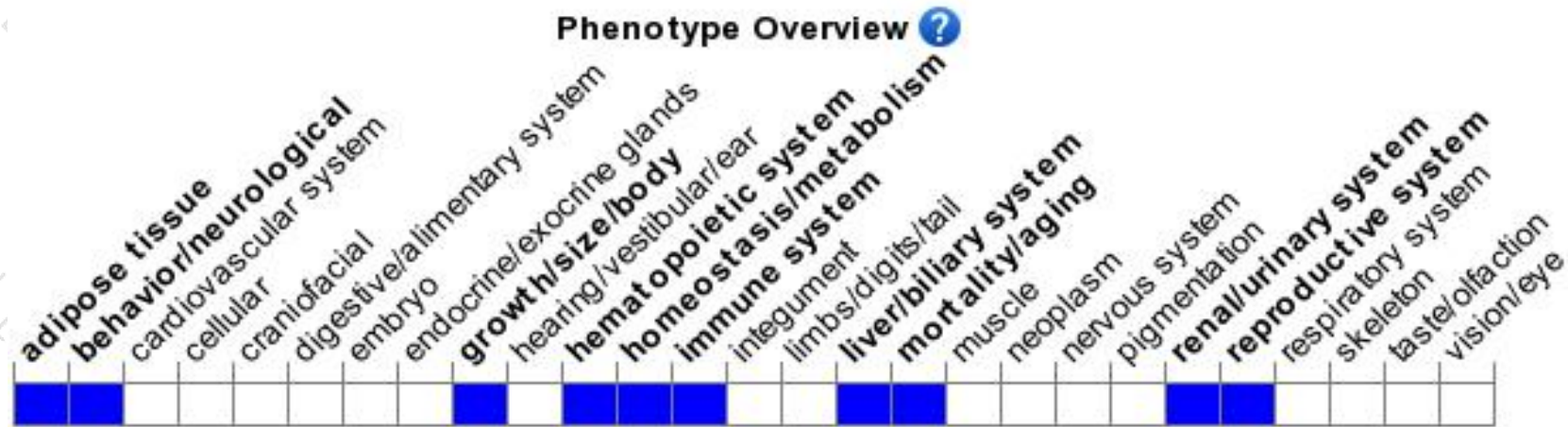
Genomic location distribution



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, Homozygous null mutant cause partial lethality before weaning, however surviving null are small but fertile and show insulin-sensitive phenotype in female under high fat diet feeding condition.

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

Tel: 400-9660890

