

Ccl5 Cas9-KO Strategy

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Project Overview



Project Name Ccl5

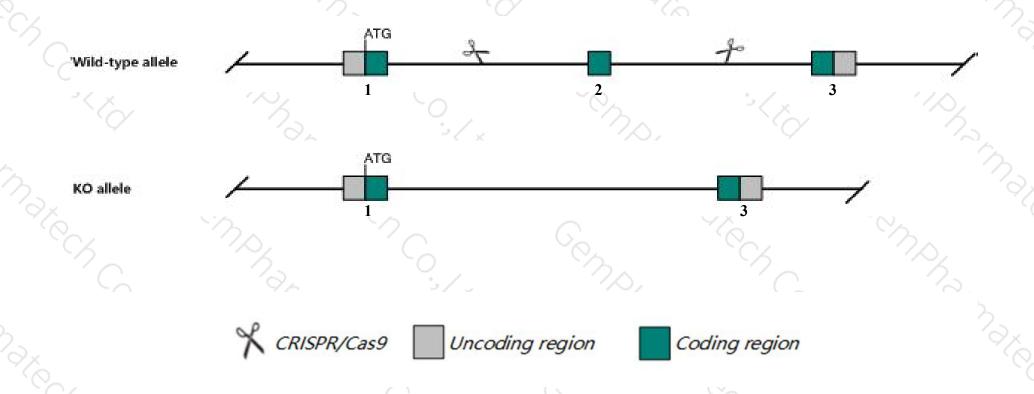
Project type Cas9-KO

Strain background C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Ccl5 gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Ccl5* gene has 2 transcripts. According to the structure of *Ccl5* gene, exon2 of *Ccl5-201*(ENSMUST00000035938.2) transcript is recommended as the knockout region. The region contains 112bp coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Ccl5* gene. The brief process is as follows: CRISPR/Cas9 system v

Notice



- ➤ According to the existing MGI data, Homozygous mutation of this gene results in decreased ear and footpad swelling following injection with DHT. T cell proliferative response is also impaired in mutant mice.
- The *Ccl5* gene is located on the Chr11. 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)



Ccl5 chemokine (C-C motif) ligand 5 [Mus musculus (house mouse)]

Gene ID: 20304, updated on 9-Apr-2019

Summary

☆ ?

Official Symbol Ccl5 provided by MGI

Official Full Name chemokine (C-C motif) ligand 5 provided by MGI

Primary source MGI:MGI:98262

See related Ensembl: ENSMUSG00000035042

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 MuRantes, RANTES, SISd, Scya5, TCP228

Expression Biased expression in small intestine adult (RPKM 266.1), mammary gland adult (RPKM 266.0) and 5 other tissuesSee more

Orthologs <u>human</u> all

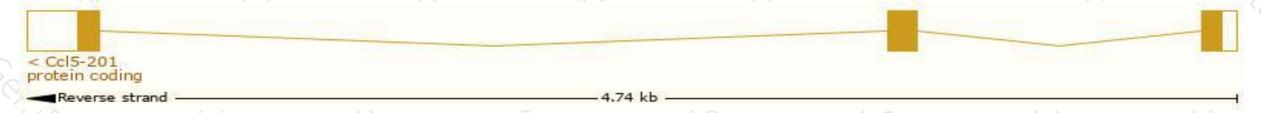
Transcript information (Ensembl)



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

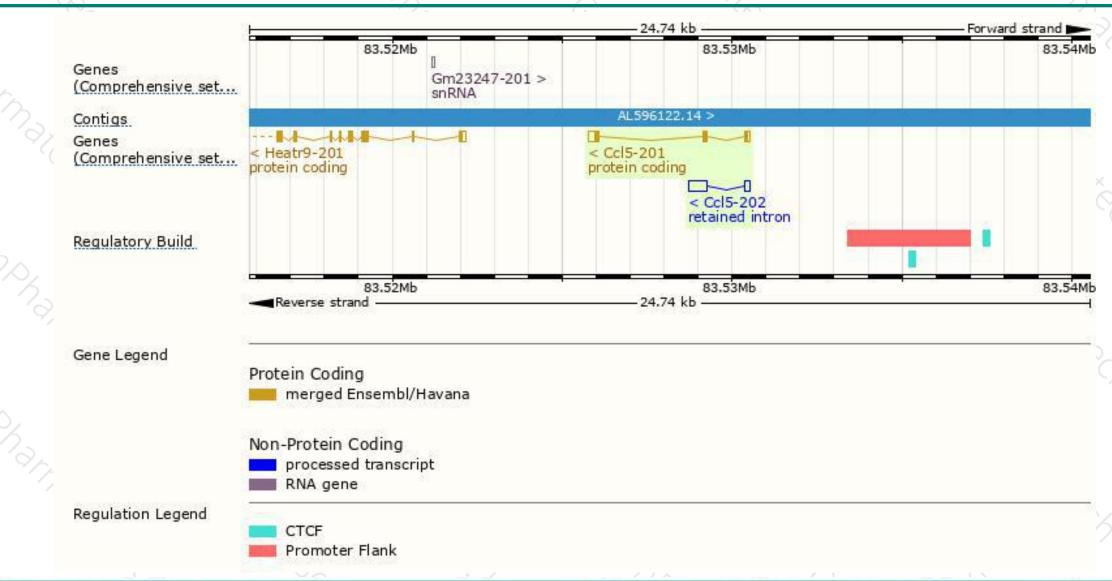
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ccl5-201	ENSMUST00000035938.2	532	<u>91aa</u>	Protein coding	CCDS36254	P30882 Q5XZF2	TSL:1 GENCODE basic APPRIS P1
Ccl5-202	ENSMUST00000125015.1	668	No protein	Retained intron	-		TSL:2

The strategy is based on the design of *Ccl5-201* transcript, The transcription is shown below



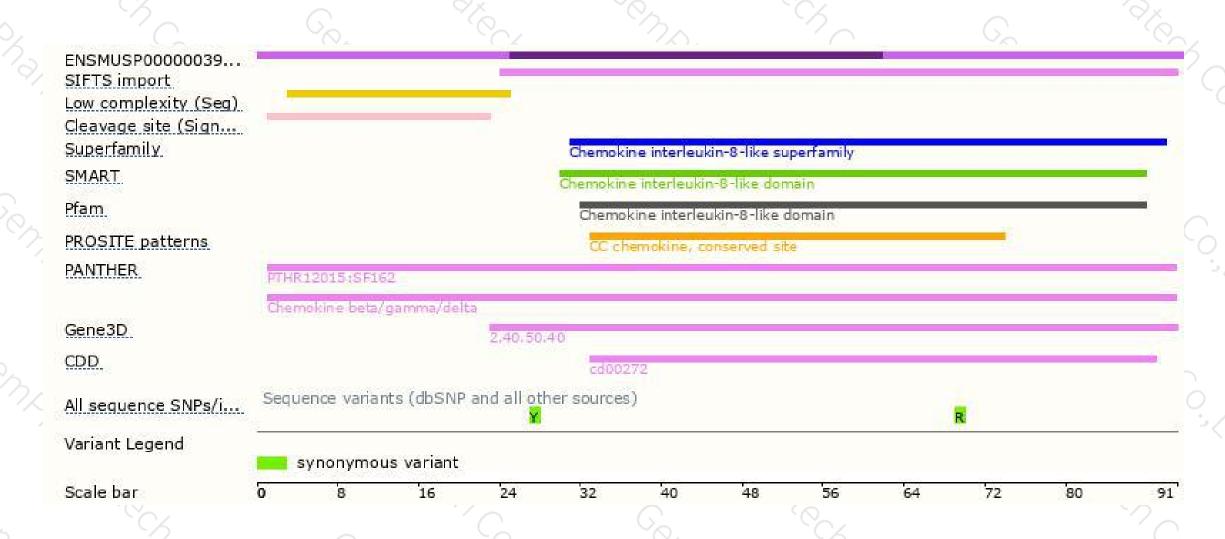
Genomic location distribution





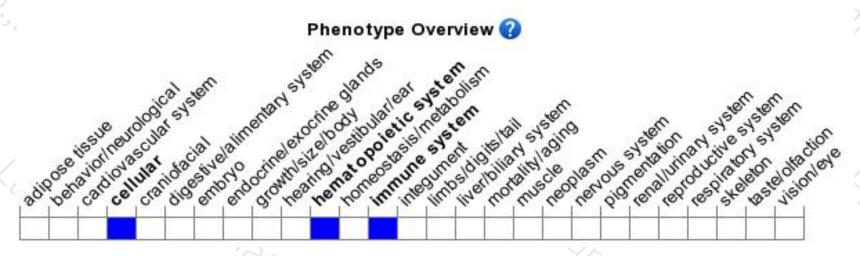
Protein domain





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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If you have any questions, you are welcome to inquire. Tel: 400-9660890





