

Rasgrf2 Cas9-CKO Strategy

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Date:2020-02-05

Project Overview



Project Name

Rasgrf2

Project type

Cas9-CKO

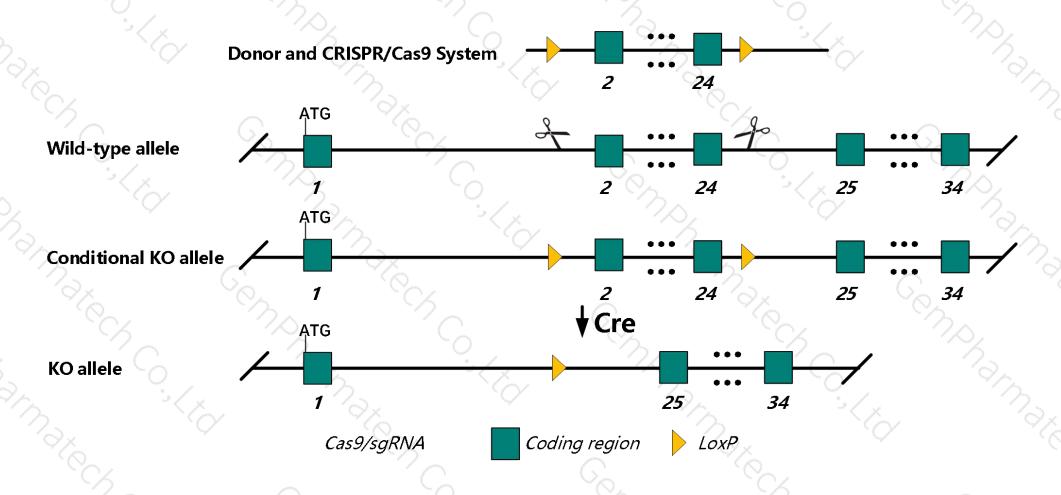
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Rasgrf2 gene. The schematic diagram is as follows:



Technical routes



- The *Rasgrf2* gene has 6 transcripts. According to the structure of *Rasgrf2* gene, exon2-exon24 of *Rasgrf2-201* (ENSMUST00000099326.9) transcript is recommended as the knockout region. The region contains 2254bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Rasgrf2* gene. The brief process is as follows:CRISPR/Cas9 system and Donor were microinjected into the fertilized eggs of C57BL/6JGpt mice. Fertilized eggs were transplanted to obtain positive F0 mice which were confirmed by PCR and sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6JGpt mice.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

Notice



- ➤ According to the existing MGI data, Mice homozygous for a targeted null mutation exhibit decreased Il2 and TNF-alpha production in stimulated T cells. Mice homozygous for mutations in both *Rasgrf1* and *Rasgrf2* exhibit no additional abnormalities than those observed in the *Rasgrf1* mutant mice.
- ➤ The effect on transcript *Rasgrf2*-202&203&204&205 is unknown.
- The *Rasgrf2* 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)



Rasgrf2 RAS protein-specific guanine nucleotide-releasing factor 2 [Mus musculus (house mouse)]

Gene ID: 19418, updated on 5-Nov-2019

Summary

Official Full Name RAS protein-specific guanine nucleotide-releasing factor 2 provided by MGI

Primary source MGI:MGI:109137

Official Symbol Rasgrf2 provided by MGI

See related Ensembl:ENSMUSG00000021708

Gene type protein coding
RefSeq status VALIDATED
Organism <u>Mus musculus</u>

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae;

Murinae; Mus; Mus

Also known as Grf2; AW048350; Ras-GRF2; 6330417G04Rik

Annotation information Annotation category: suggests misassembly

Annotation category: partial on reference assembly

Expression Biased expression in frontal lobe adult (RPKM 11.9), cortex adult (RPKM 11.9) and 14 other tissues See more

Orthologs human all

Genomic context

☆ ?

Location: 13 C3; 13 47.43 cM

See Rasgrf2 in Genome Data Viewer

Exon count: 26

| Annotation release | Status | Assembly | Chr | Location | |
|--------------------|-------------------|------------------------------|-----|---|--|
| 108 | current | GRCm38.p6 (GCF_000001635.26) | 13 | NC_000079.6 (9188040792131828, complement) | |
| Build 37.2 | previous assembly | MGSCv37 (GCF_000001635.18) | 13 | NC_000079.5 (9279269592901449, complement) , (9202001292127647, complement) | |

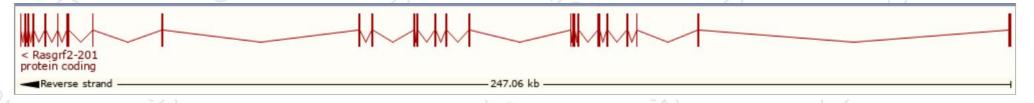
Transcript information (Ensembl)



The gene has 6 transcripts, all transcripts are shown below:

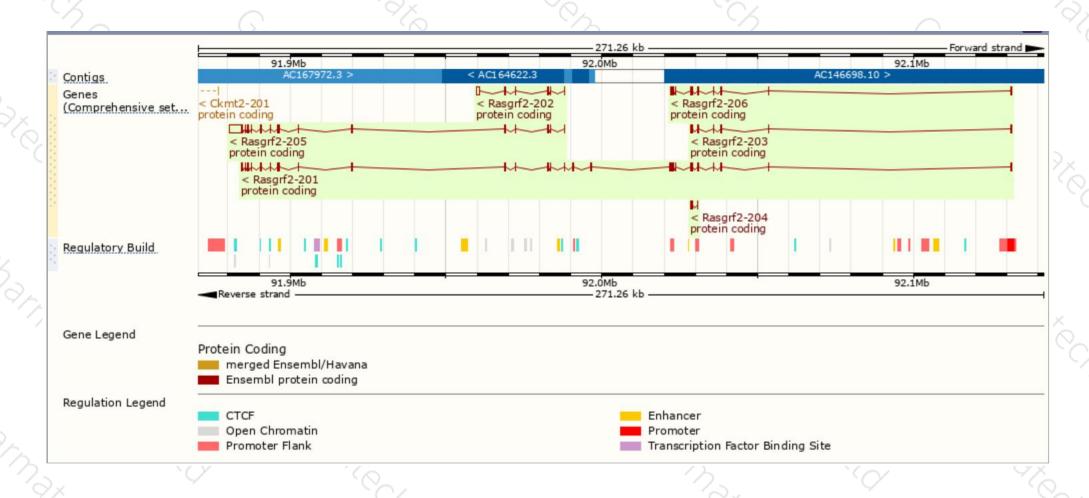
| Name 🍦 | Transcript ID 🗼 | bp 🍦 | Protein | Biotype | CCDS | UniProt 🖕 | Flags |
|-------------|----------------------|------|---------------|----------------|------------------|-------------|--------------------------------|
| Rasgrf2-205 | ENSMUST00000151408.7 | 5797 | <u>588aa</u> | Protein coding | - | F7B9R2函 | CDS 5' incomplete TSL:1 |
| Rasgrf2-201 | ENSMUST00000099326.9 | 3567 | <u>1188aa</u> | Protein coding | | D3Z6K8配 | TSL:5 GENCODE basic APPRIS P1 |
| Rasgrf2-206 | ENSMUST00000216219.1 | 2115 | <u>505aa</u> | Protein coding | 1.5 | A0A1L1SS23译 | TSL:5 GENCODE basic |
| Rasgrf2-202 | ENSMUST00000142378.1 | 1789 | 252aa | Protein coding | - | F6TYF8译 | CDS 5' incomplete TSL:1 |
| Rasgrf2-203 | ENSMUST00000146492.2 | 1323 | 387aa | Protein coding | 18 | D3Z685怪 | CDS 3' incomplete TSL:5 |
| Rasgrf2-204 | ENSMUST00000149630.7 | 502 | <u>168aa</u> | Protein coding | 5 - 5 | F6TCD3₽ | CDS 5' and 3' incomplete TSL:5 |

The strategy is based on the design of Rasgrf2-201 transcript, The transcription is shown below



Genomic location distribution





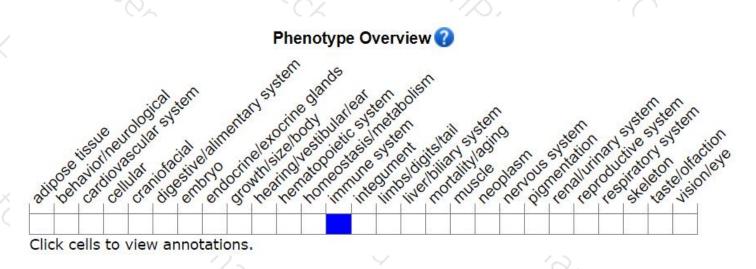
Protein domain





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 targeted null mutation exhibit decreased Il2 and TNF-alpha production in stimulated T cells. Mice homozygous for mutations in both Rasgrf1 and Rasgrf2 exhibit no additional abnormalities than those observed in the Rasgrf1 mutant mice.



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





