



Isg20l2 Cas9-KO Strategy

Designer: Jia Yu

Reviewer: Xiaojing Li

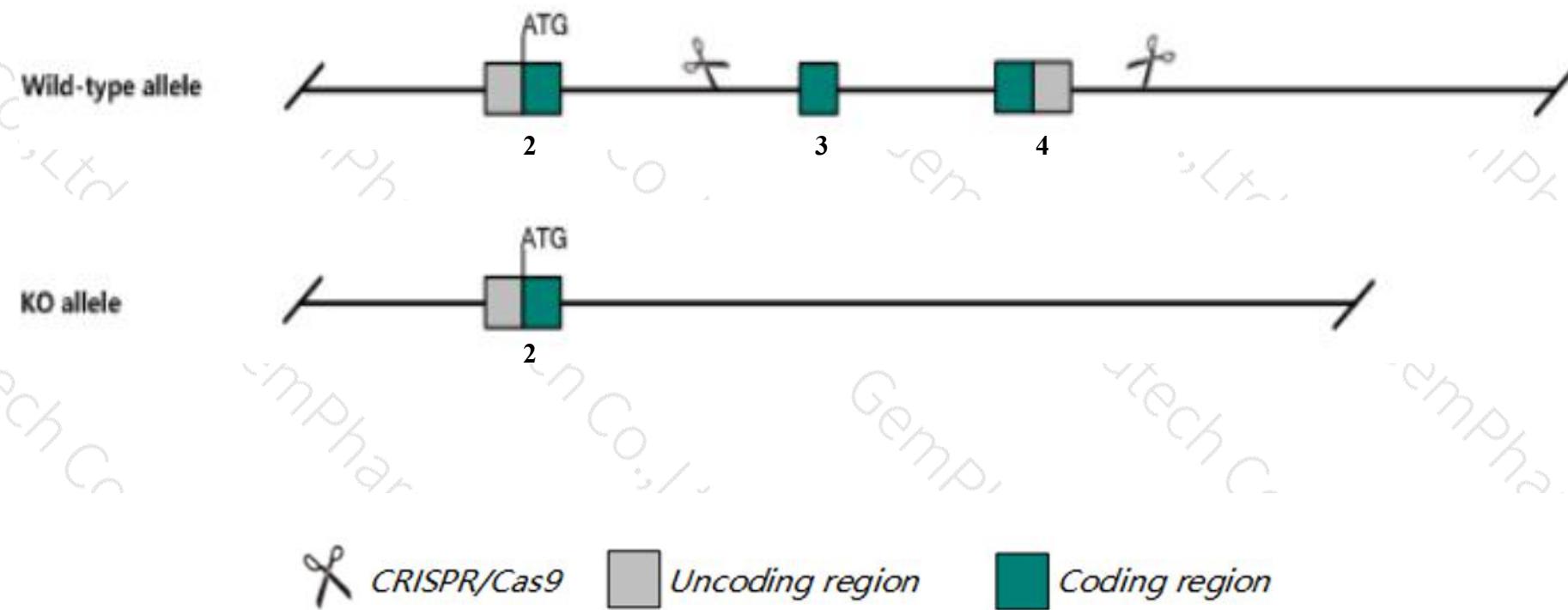
Design Date: 2020-9-25

Project Overview

| | |
|--------------------------|----------------|
| Project Name | <i>Isg20l2</i> |
| Project type | Cas9-KO |
| Strain background | C57BL/6JGpt |

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Isg20l2* gene. The schematic diagram is as follows:



Technical routes

- The *Isg20l2* gene has 1 transcript. According to the structure of *Isg20l2* gene, exon3-exon4 of *Isg20l2-20l*(ENSMUST00000055984.6) transcript is recommended as the knockout region. The region contains 315bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Isg20l2* gene. The brief process is as follows: CRISPR/Cas9 system 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.



集萃药康
GemPharmatech

Notice

- The *Isg20l2* gene is located on the Chr3. 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.



集萃药康
GemPharmatech

Gene information (NCBI)

Isg20l2 interferon stimulated exonuclease gene 20-like 2 [Mus musculus (house mouse)]

Gene ID: 229504, updated on 13-Mar-2020

Summary



Official Symbol Isg20l2 provided by [MGI](#)

Official Full Name interferon stimulated exonuclease gene 20-like 2 provided by [MGI](#)

Primary source [MGI:MGI:2140076](#)

See related [Ensembl:ENSMUSG00000048039](#)

Gene type protein coding

RefSeq status PROVISIONAL

Organism [Mus musculus](#)

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

Also known as 4930429N24, C80587

Expression Ubiquitous expression in testis adult (RPKM 26.3), CNS E11.5 (RPKM 14.8) and 28 other tissues [See more](#)

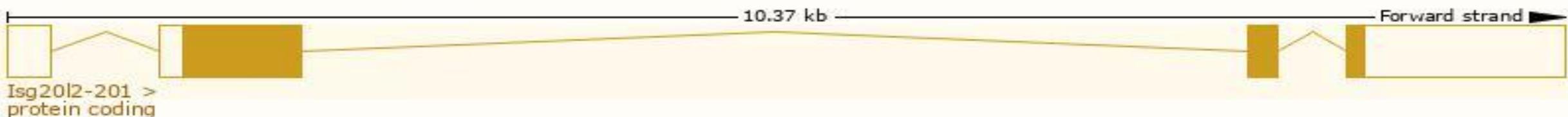
Orthologs [human](#) [all](#)

Transcript information (Ensembl)

The gene has 1 transcript, and the transcript is shown below:

| Name | Transcript ID | bp | Protein | Biotype | CCDS | UniProt | Flags |
|-------------|--------------------------------------|------|-----------------------|----------------|---------------------------|----------------------------|-------------------------------|
| Isg20l2-201 | ENSMUST00000055984.6 | 2893 | 368aa | Protein coding | CCDS17459 | A0A0R4J0R3 | TSL:1 GENCODE basic APPRIS P1 |

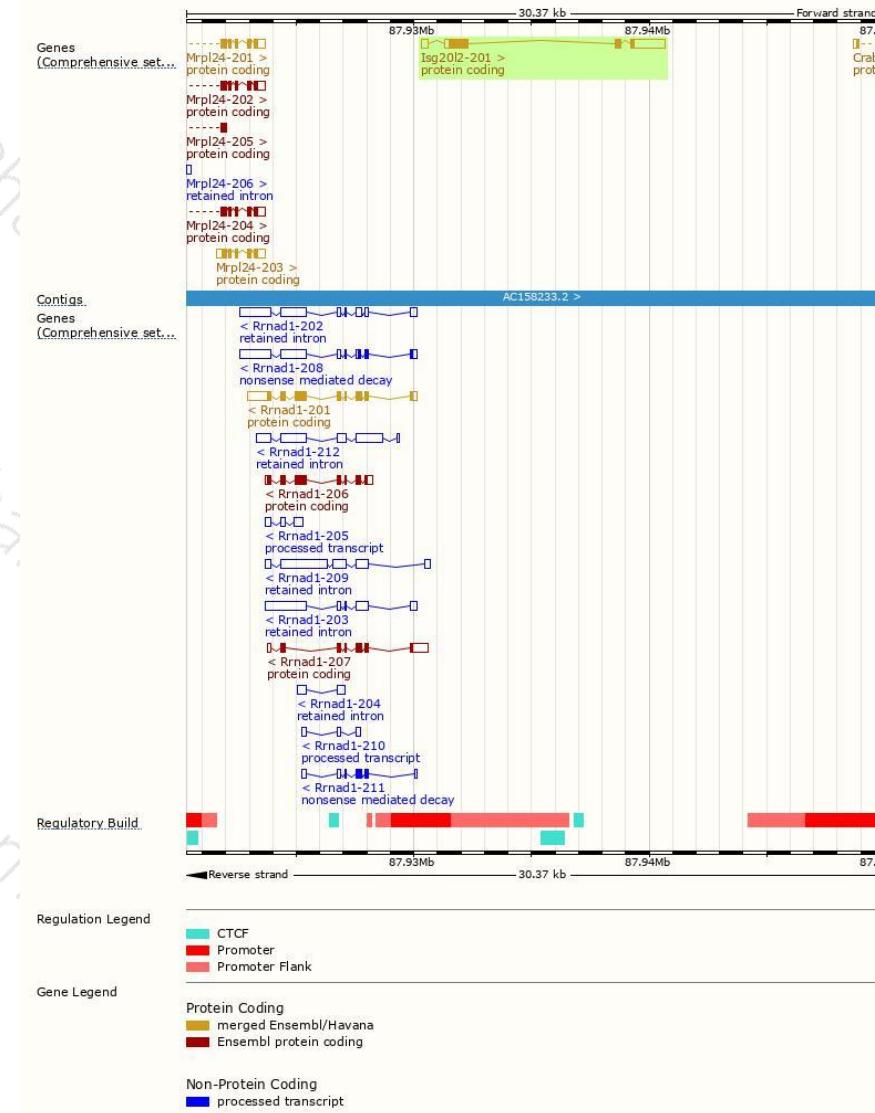
The strategy is based on the design of *Isg20l2-201* transcript, the transcription is shown below:



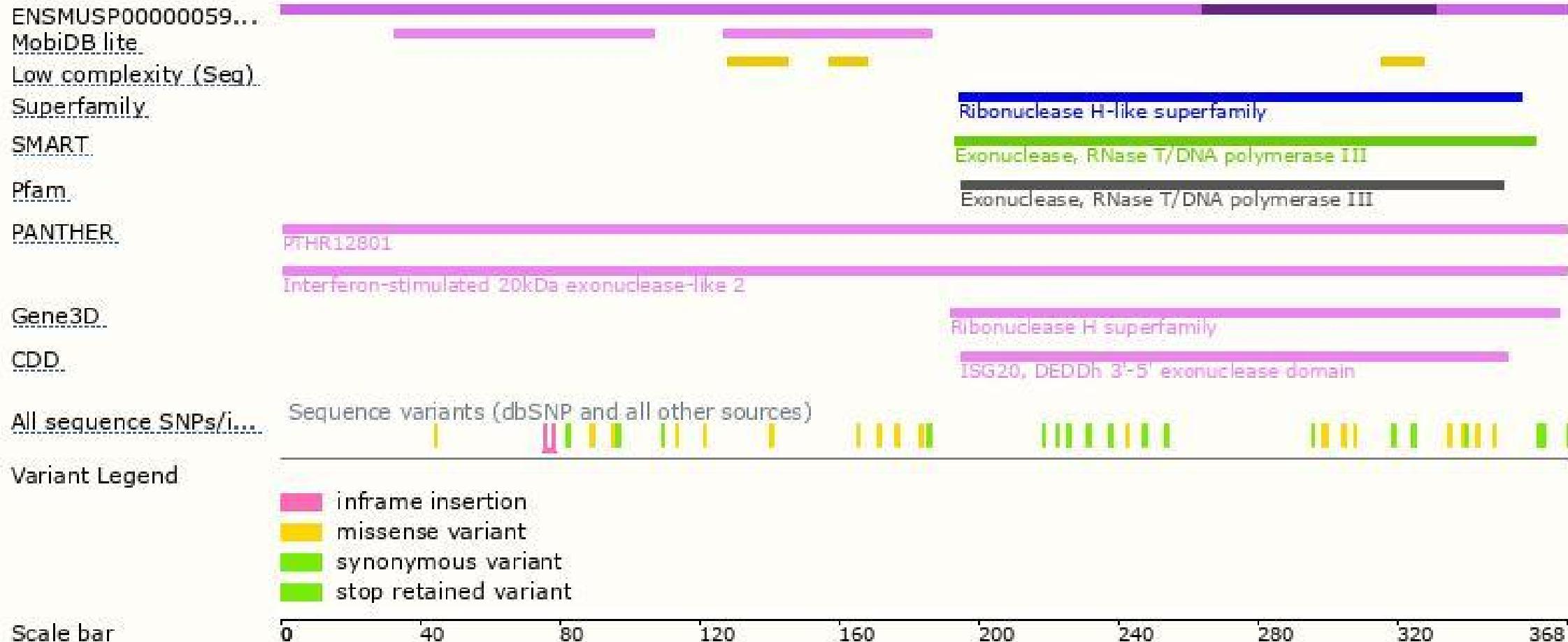


集萃药康
GemPharmatech

Genomic location distribution



Protein domain





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

Tel: 400-9660890



集萃药康生物科技
GemPharmatech Co.,Ltd

