

Adgrg7 Cas9-KO Strategy

Designer:

Huan Wang

Design Date:

2019-7-22

Project Overview

Project Name

Adgrg7

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Adgrg7* gene. The schematic diagram is as follows:



- The *Adgrg7* gene has 1 transcript. According to the structure of *Adgrg7* gene, exon2-exon4 of *Adgrg7-201* (ENSMUST00000023437.4) transcript is recommended as the knockout region. The region contains 332bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Adgrg7* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice homozygous for a knock-out allele exhibit reduced body weight gain and increased frequency of peristalsis and slow wave potential in the small intestine.
- The *Adgrg7* gene is located on the Chr16. 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)

Adgrg7 adhesion G protein-coupled receptor G7 [Mus musculus (house mouse)]

Gene ID: 239853, updated on 24-Feb-2019

Summary



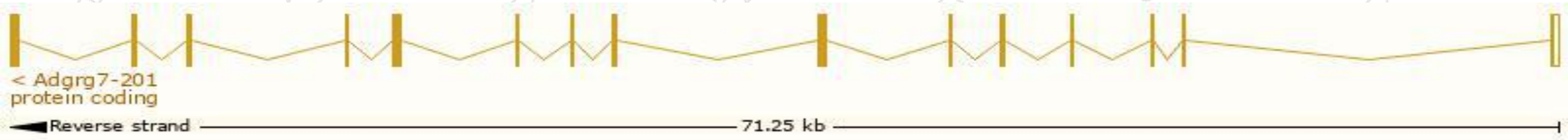
Official Symbol	Adgrg7 provided by MGI
Official Full Name	adhesion G protein-coupled receptor G7 provided by MGI
Primary source	MGI:MGI:2441732
See related	Ensembl:ENSMUSG00000022755
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	9130020O16Rik, Gpr128
Expression	Biased expression in large intestine adult (RPKM 13.6), colon adult (RPKM 8.1) and 2 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
Adgrg7-201	ENSMUST00000023437.4	2698	785aa	Protein coding	CCDS28225	Q8BM96	TSL:1 GENCODE basic APPRIS P1

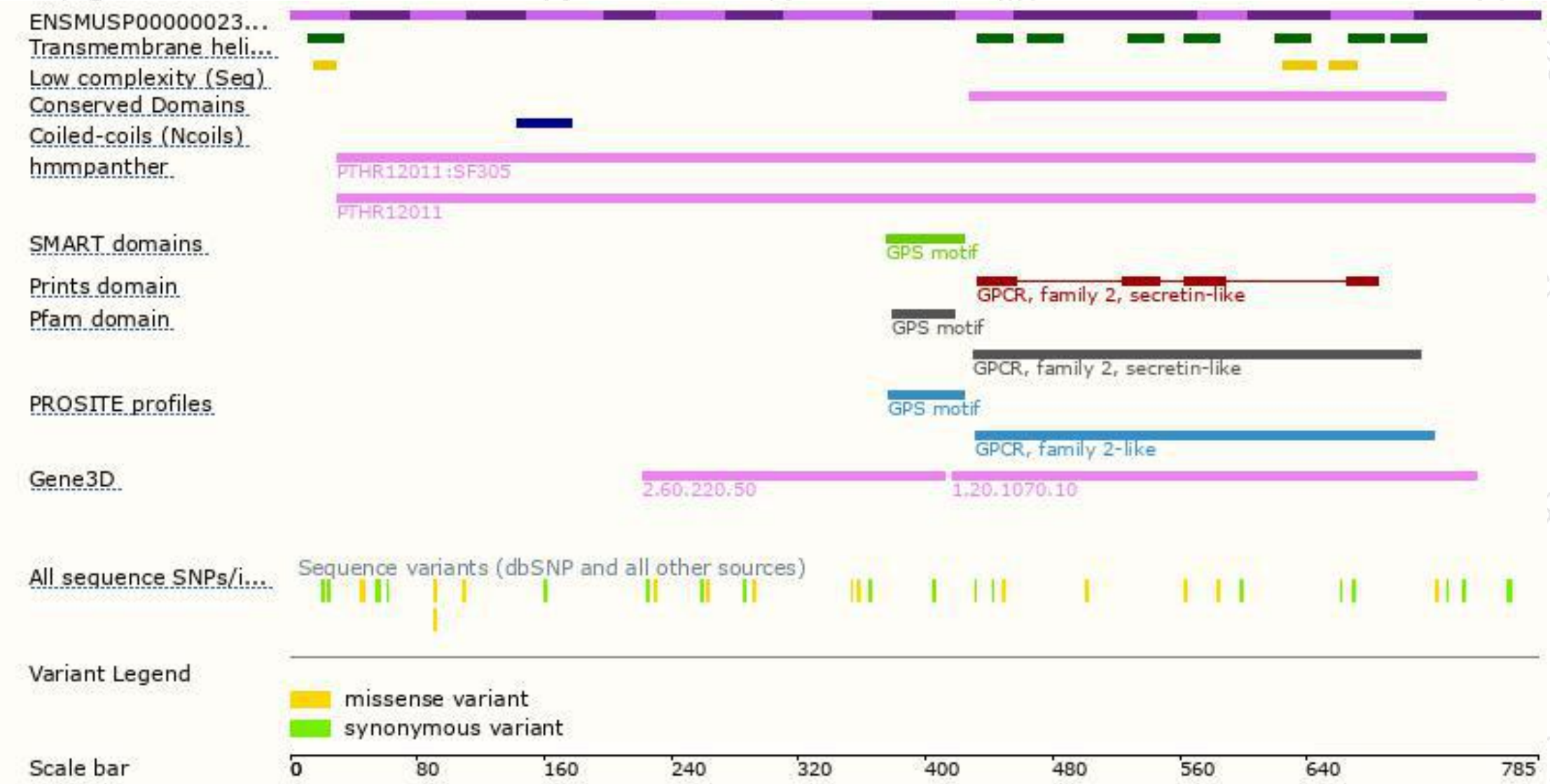
The strategy is based on the design of *Adgrg7-201* transcript, The transcription is shown below



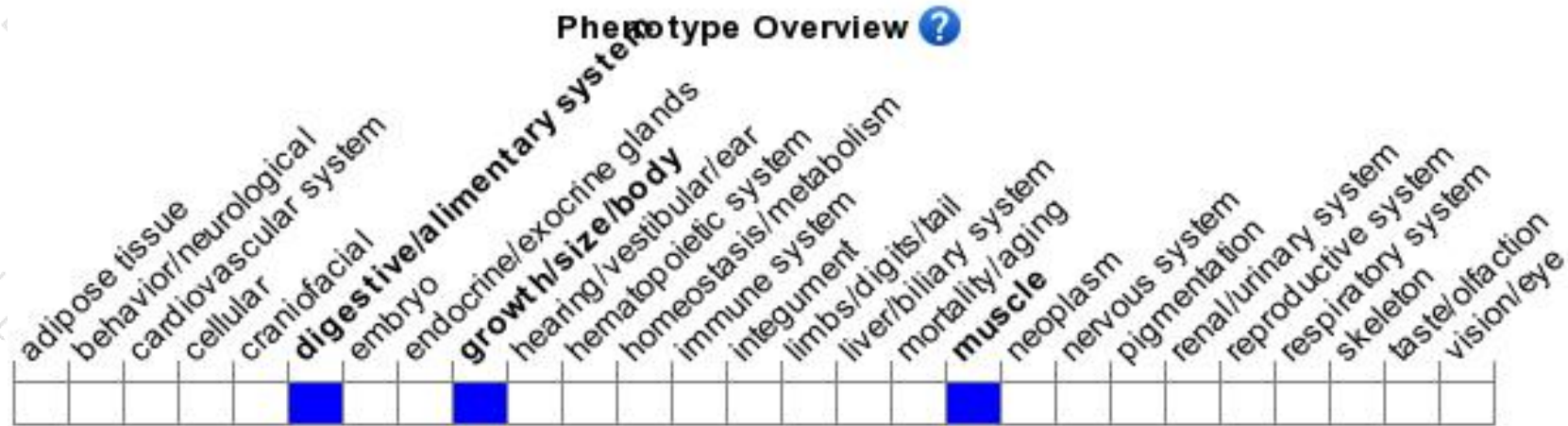
Genomic location distribution



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 knock-out allele exhibit reduced body weight gain and increased frequency of peristalsis and slow wave potential in the small intestine.

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

Tel: 400-9660890

