

***Golga5* Cas9-KO Strategy**

Designer: JiaYu

Reviewer: Xiaojing Li

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Project Overview

Project Name

Golga5

Project type

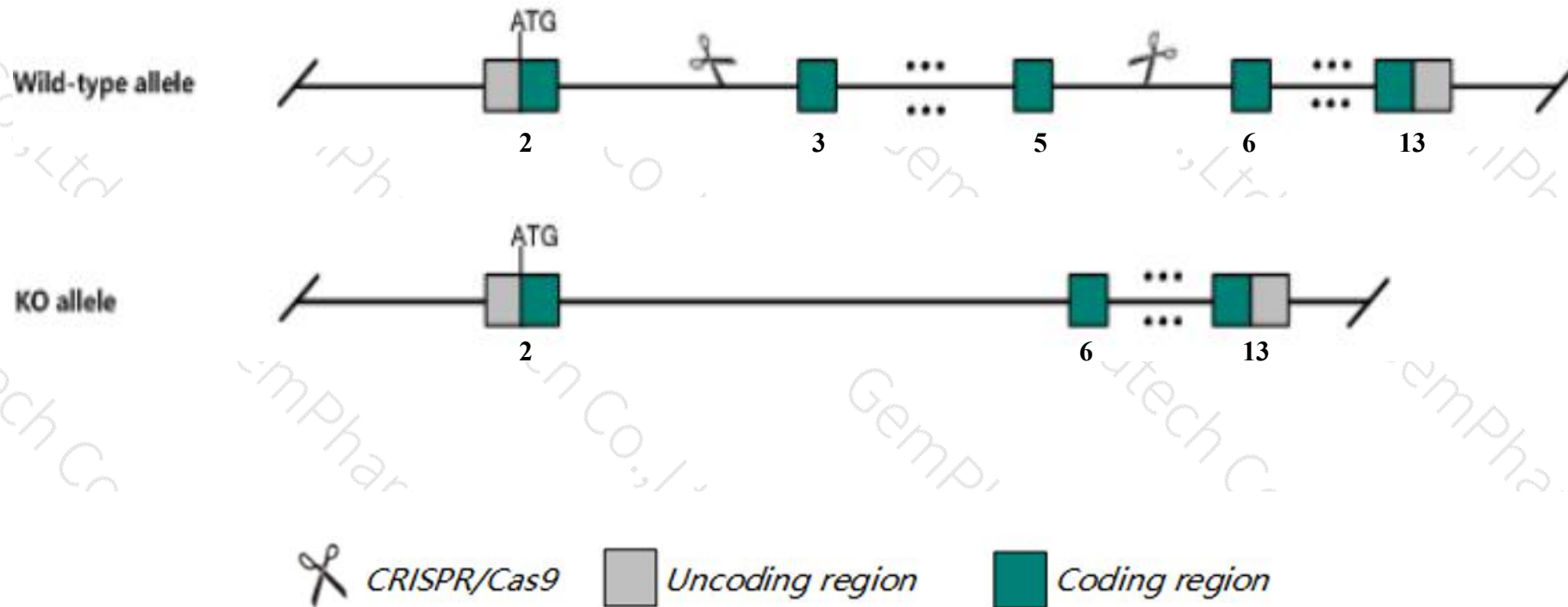
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Golga5* gene. The schematic diagram is as follows:



- The *Golga5* gene has 3 transcripts. According to the structure of *Golga5* gene, exon3-exon5 of *Golga5*-201(ENSMUST00000021609.9) transcript is recommended as the knockout region. The region contains 572bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Golga5* 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.

- According to the existing MGI data, homozygous knockout does not result in an obvious phenotype and does not affect Golgi apparatus morphology.
- Transcript 203 CDS 5' incomplete the influences is unknown.
- The *Golga5* gene is located on the Chr12. 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)

Golga5 golgi autoantigen, golgin subfamily a, 5 [Mus musculus (house mouse)]

Gene ID: 27277, updated on 13-Mar-2020

Summary



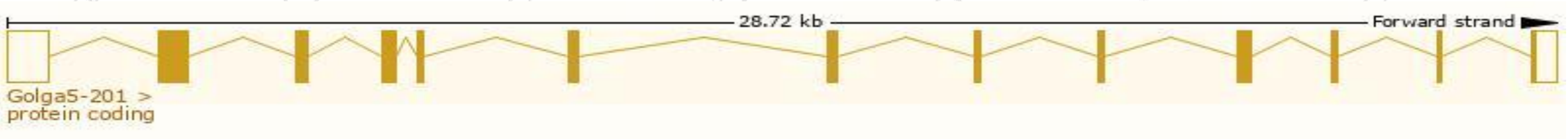
Official Symbol	Golga5 provided by MGI
Official Full Name	golgi autoantigen, golgin subfamily a, 5 provided by MGI
Primary source	MGI:MGI:1351475
See related	Ensembl:ENSMUSG00000021192
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Ret-II
Expression	Ubiquitous expression in bladder adult (RPKM 11.9), testis adult (RPKM 11.5) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

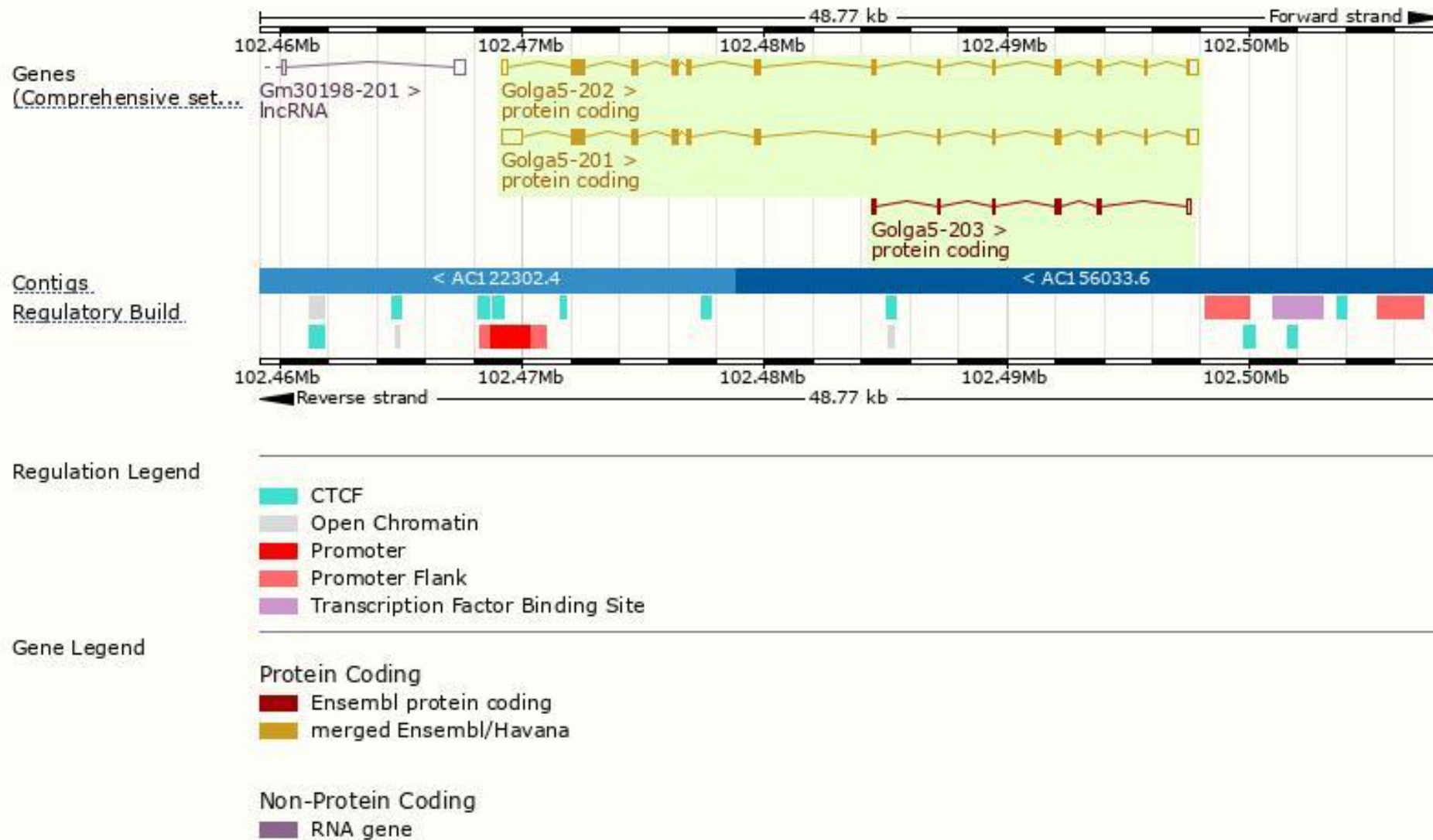
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Golga5-201	ENSMUST00000021609.9	3390	729aa	Protein coding	CCDS36526	Q9QYE6	TSL:1 GENCODE basic APPRIS P1
Golga5-202	ENSMUST00000179218.8	2827	729aa	Protein coding	CCDS36526	Q9QYE6	TSL:1 GENCODE basic APPRIS P1
Golga5-203	ENSMUST00000222744.1	814	234aa	Protein coding	-	A0A1Y7VMD3	CDS 5' incomplete TSL:3

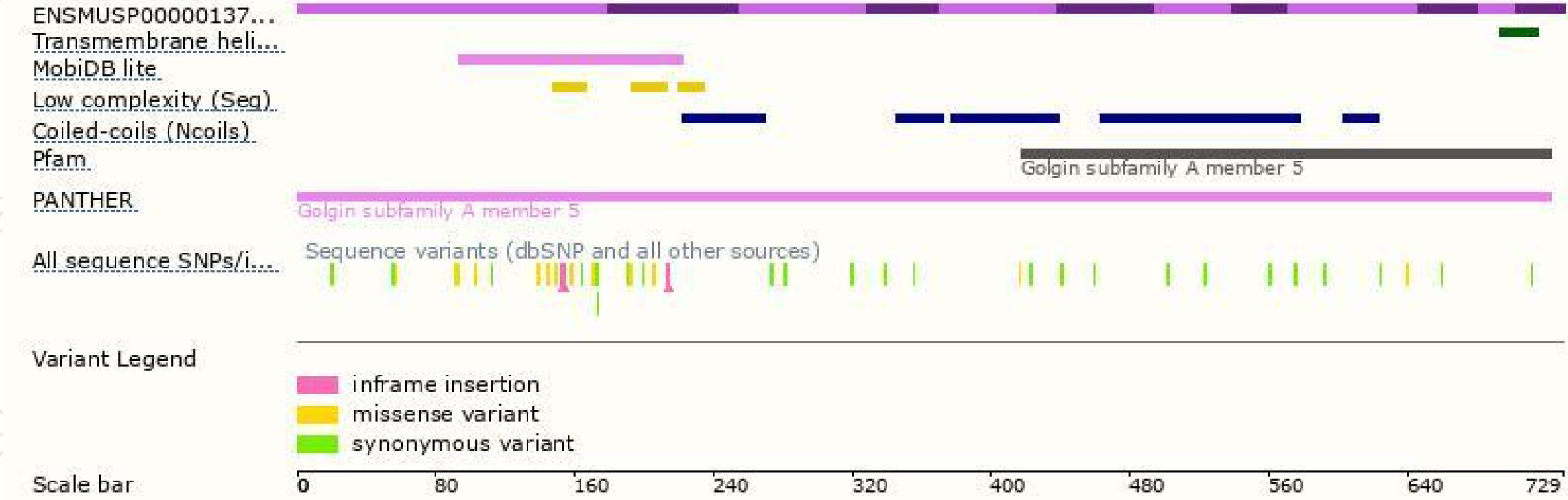
The strategy is based on the design of *Golga5-201* transcript,the transcription is shown below:



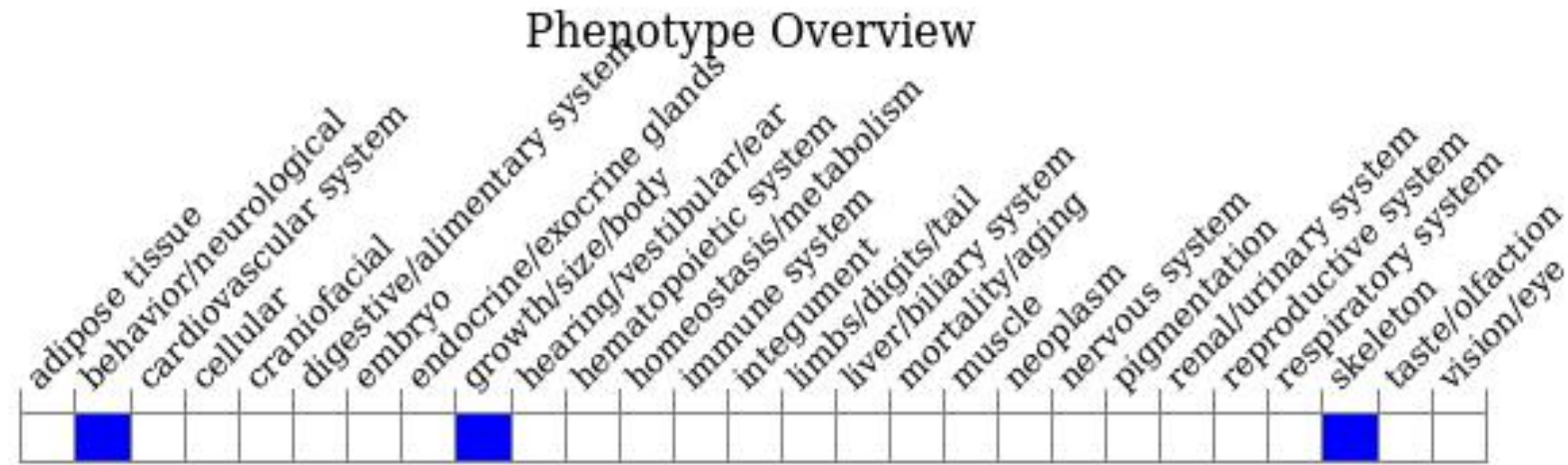
Genomic location distribution



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 knockout does not result in an obvious phenotype and does not affect Golgi apparatus morphology.

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

Tel: 400-9660890

