

Golga4 Cas9-CKO Strategy

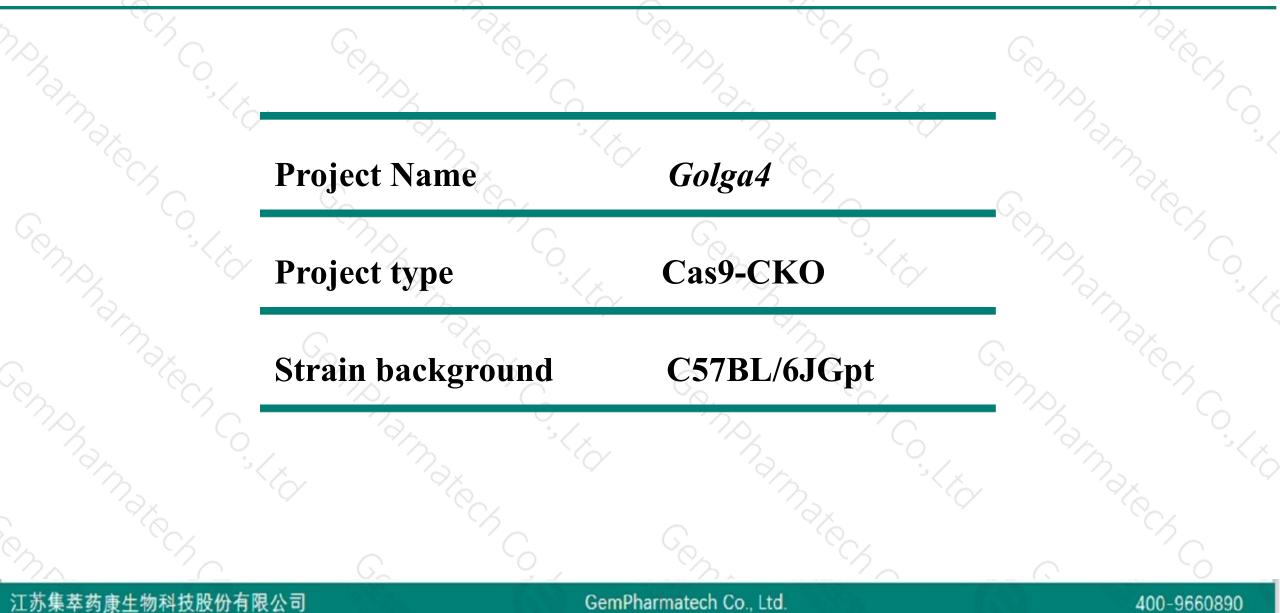
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Reviewer: Xiaojing Li

Design Date: 2020-8-25

Project Overview



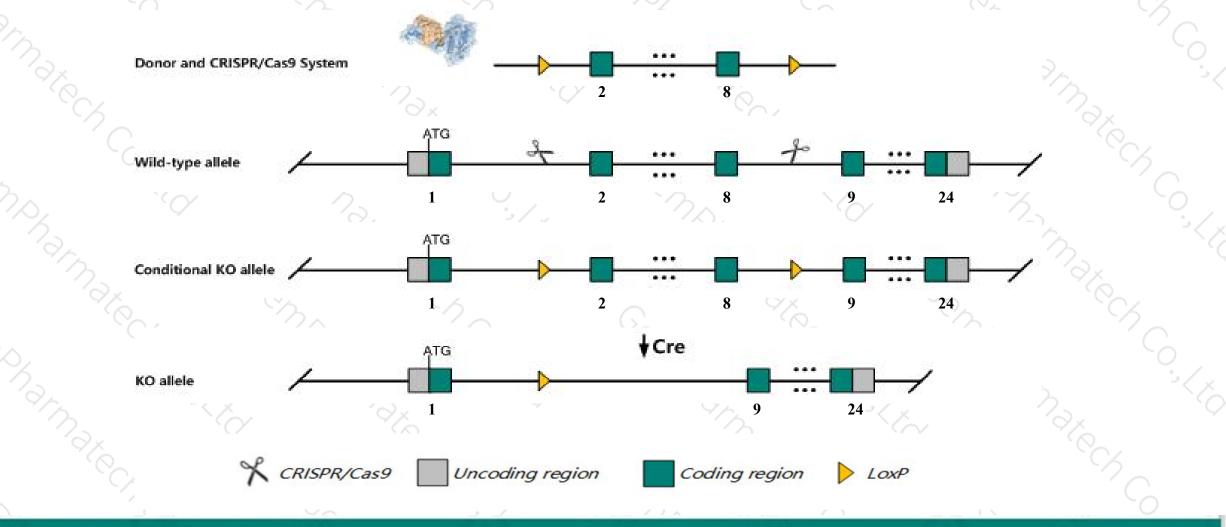


Conditional Knockout strategy



400-9660890

This model will use CRISPR/Cas9 technology to edit the Golga4 gene. The schematic diagram is as follows:



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The Golga4 gene has 13 transcripts. According to the structure of Golga4 gene, exon2-exon8 of Golga4-201(ENSMUST0000084820.5) transcript is recommended as the knockout region. The region contains 805bp coding sequence. Knock out the region will result in disruption of protein function.

➤ In this project we use CRISPR/Cas9 technology to modify *Golga4* 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.



- > The Golga4 gene is located on the Chr9. 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.
- ≻Transcript 204 CDS 5' incomplete the influences is unknown.
- ≻Transcript 206 CDS 3' incomplete the influences is unknown.
- ≻Transcript 210 CDS 5' and 3' incomplete the influences is unknown.
- > 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)



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Golga4 golgi autoantigen, golgin subfamily a, 4 [Mus musculus (house mouse)]

Gene ID: 54214, updated on 13-Mar-2020

Summary

Official Symbol	Golga4 provided by MGI
Official Full Name	golgi autoantigen, golgin subfamily a, 4 provided by MGI
Primary source	MGI:MGI:1859646
See related	Ensembl:ENSMUSG0000038708
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	AI225887, AU019508, Olp-1
Expression	Ubiquitous expression in CNS E11.5 (RPKM 6.1), placenta adult (RPKM 5.1) and 28 other tissues See more
Orthologs	human all

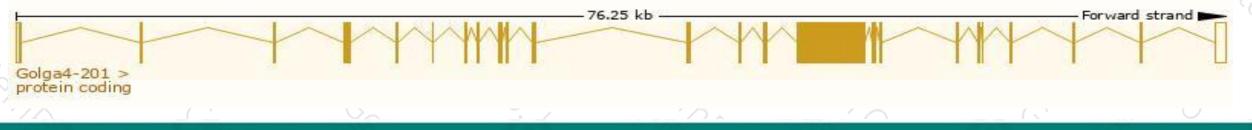
Transcript information (Ensembl)



The gene has 13 transcripts, all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Golga4-201	ENSMUST0000084820.5	7583	<u>2238aa</u>	Protein coding	CCDS40801	<u>Q91VW5</u>	TSL:1 GENCODE basic APPRIS P1
Golga4-204	ENSMUST00000211840.1	4461	<u>1251aa</u>	Protein coding		A0A1D5RLV4	CDS 5' incomplete TSL:1
Golga4-206	ENSMUST00000212097.1	2858	<u>941aa</u>	Protein coding	828	A0A1D5RME8	CDS 3' incomplete TSL:1
Golga4-210	ENSMUST00000212461.1	1363	<u>454aa</u>	Protein coding		A0A1D5RLC6	CDS 5' and 3' incomplete TSL:5
Golga4-208	ENSMUST00000212183.1	1237	<u>111aa</u>	Nonsense mediated decay	-	A0A1D5RM19	CDS 5' incomplete TSL:5
Golga4-205	ENSMUST00000211846.1	796	<u>118aa</u>	Nonsense mediated decay	1.70	A0A1D5RL95	CDS 5' incomplete TSL:3
Golga4-209	ENSMUST00000212274.1	615	<u>131aa</u>	Nonsense mediated decay	0.00	A0A1D5RLH4	CDS 5' incomplete TSL:5
Golga4-212	ENSMUST00000212620.1	2963	No protein	Retained intron	1	(22)	TSL:NA
Golga4-211	ENSMUST00000212593.1	2309	No protein	Retained intron	1070	070	TSL:1
Golga4-202	ENSMUST00000211826.1	687	No protein	Retained intron	-		TSL:3
Golga4-207	ENSMUST00000212151.1	611	No protein	Retained intron	828	828	TSL:2
Golga4-213	ENSMUST00000212913.1	582	No protein	Retained intron			TSL:3
Golga4-203	ENSMUST00000211838.1	362	No protein	Retained intron	-	120	TSL:2

The strategy is based on the design of Golga4-201 transcript, the transcription is shown below:

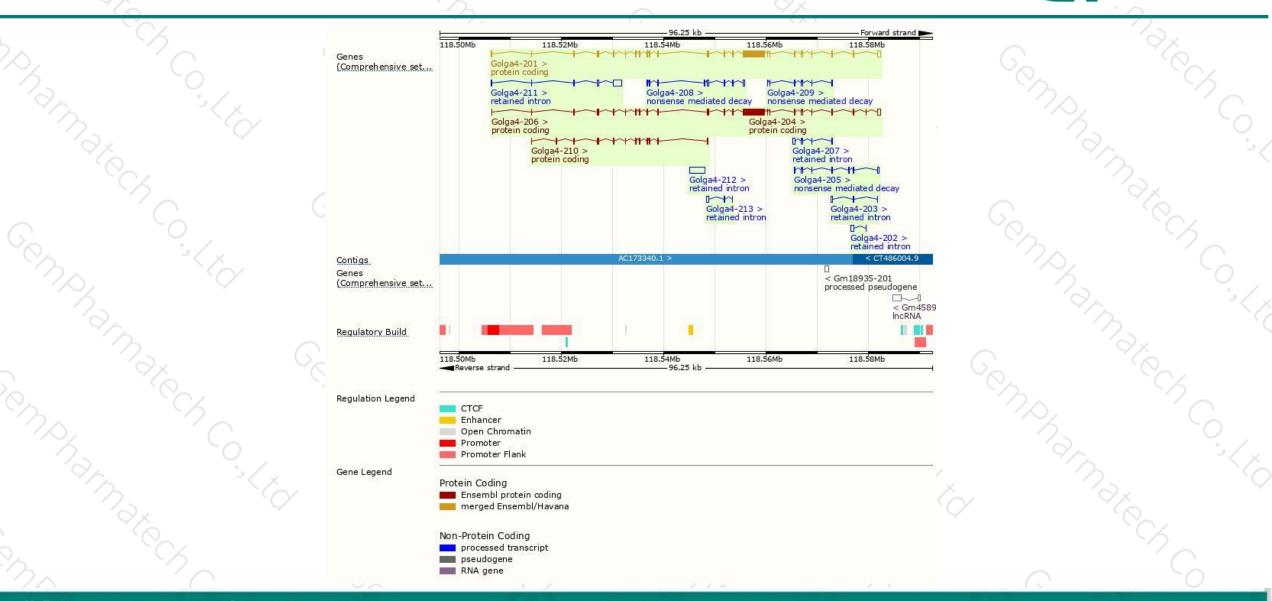


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Genomic location distribution



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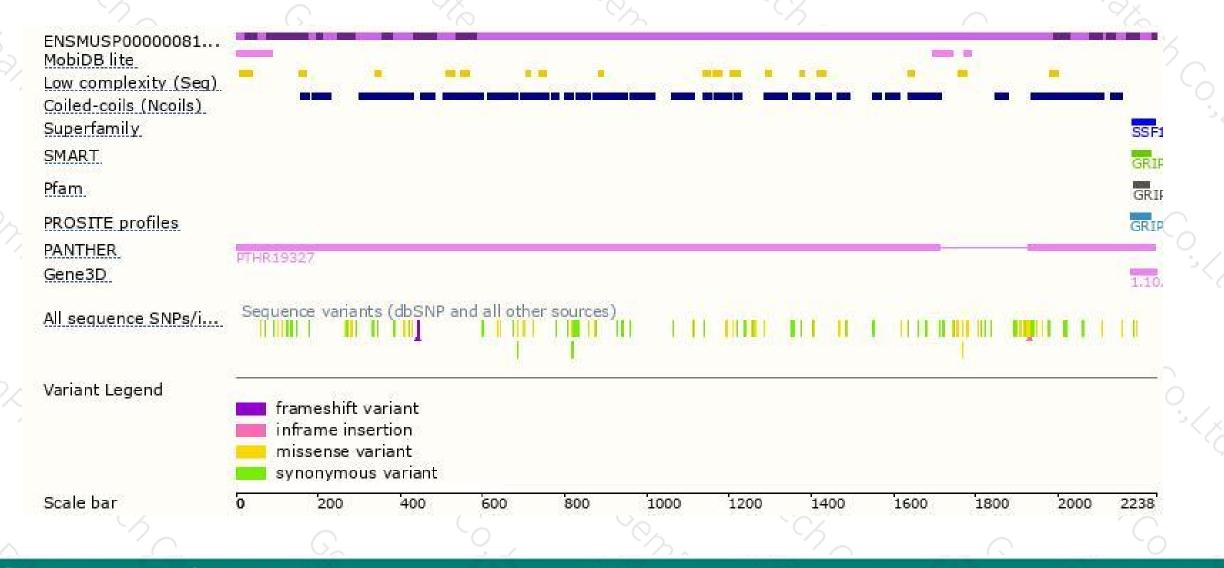
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Protein domain





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



