

***Galnt7* Cas9-KO Strategy**

Designer: Daohua Xu

Reviewer: Huimin Su

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Project Overview

Project Name

Galnt7

Project type

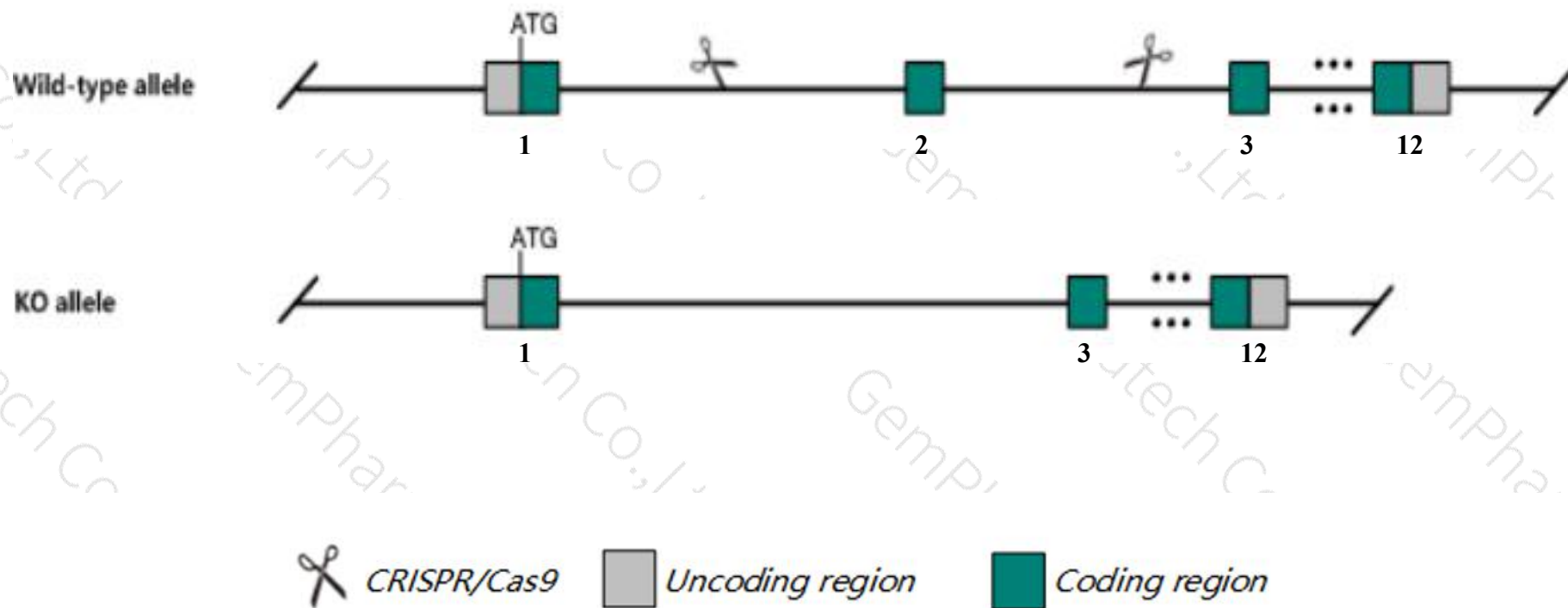
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Galnt7* gene. The schematic diagram is as follows:



- The *Galnt7* gene has 4 transcripts. According to the structure of *Galnt7* gene, exon2 of *Galnt7-201* (ENSMUST00000034021.11) transcript is recommended as the knockout region. The region contains 461bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Galnt7* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Galnt7* gene is located on the Chr8. 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)

Galnt7 polypeptide N-acetylgalactosaminyltransferase 7 [Mus musculus (house mouse)]

Gene ID: 108150, updated on 13-Mar-2020

Summary



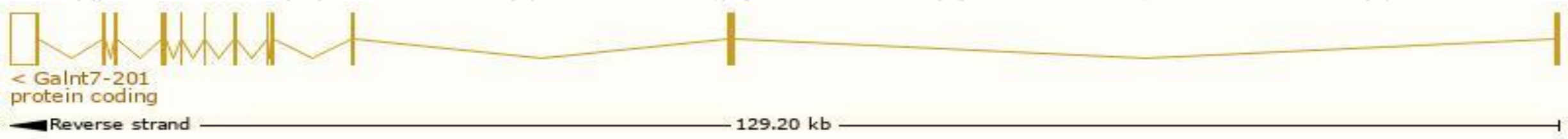
Official Symbol	Galnt7 provided by MGI
Official Full Name	polypeptide N-acetylgalactosaminyltransferase 7 provided by MGI
Primary source	MGI:MGI:1349449
See related	Ensembl:ENSMUSG000000031608
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	AI225872
Expression	Ubiquitous expression in colon adult (RPKM 6.3), placenta adult (RPKM 5.3) and 27 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

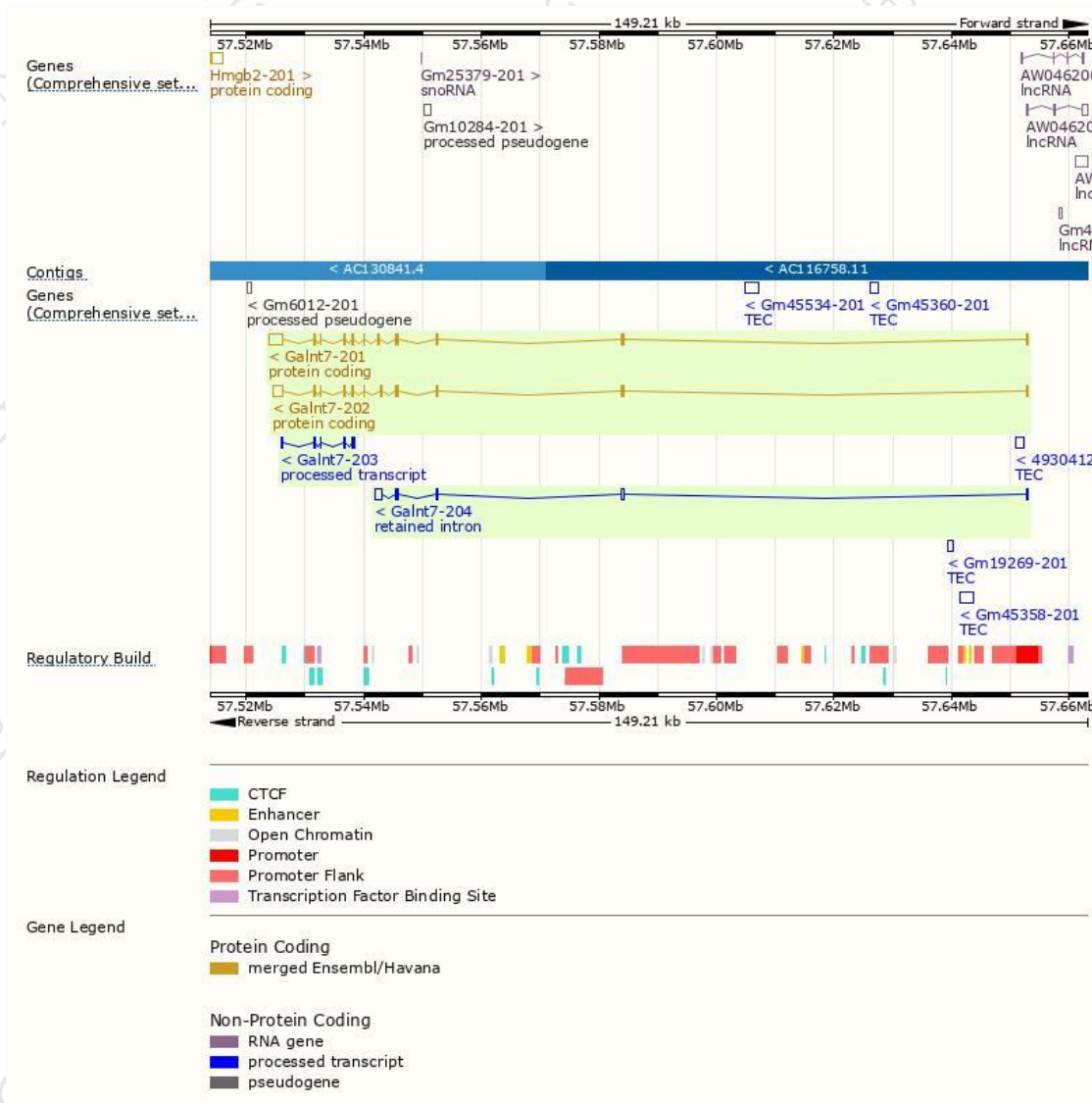
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Galnt7-201	ENSMUST00000034021.11	4313	657aa	Protein coding	CCDS22317	Q80VA0	TSL:1 GENCODE basic APPRIS is a system to annotate alternatively spliced transcripts based on a range of computational methods to identify the most functionally important transcript(s) of a gene. APPRIS P3
Galnt7-202	ENSMUST00000110316.2	3413	657aa	Protein coding	CCDS52552	Q80VA0	TSL:1 GENCODE basic APPRIS is a system to annotate alternatively spliced transcripts based on a range of computational methods to identify the most functionally important transcript(s) of a gene. APPRIS ALT1
Galnt7-203	ENSMUST00000139417.1	897	No protein	Processed transcript	-	-	TSL:5
Galnt7-204	ENSMUST00000156907.1	2042	No protein	Retained intron	-	-	TSL:1

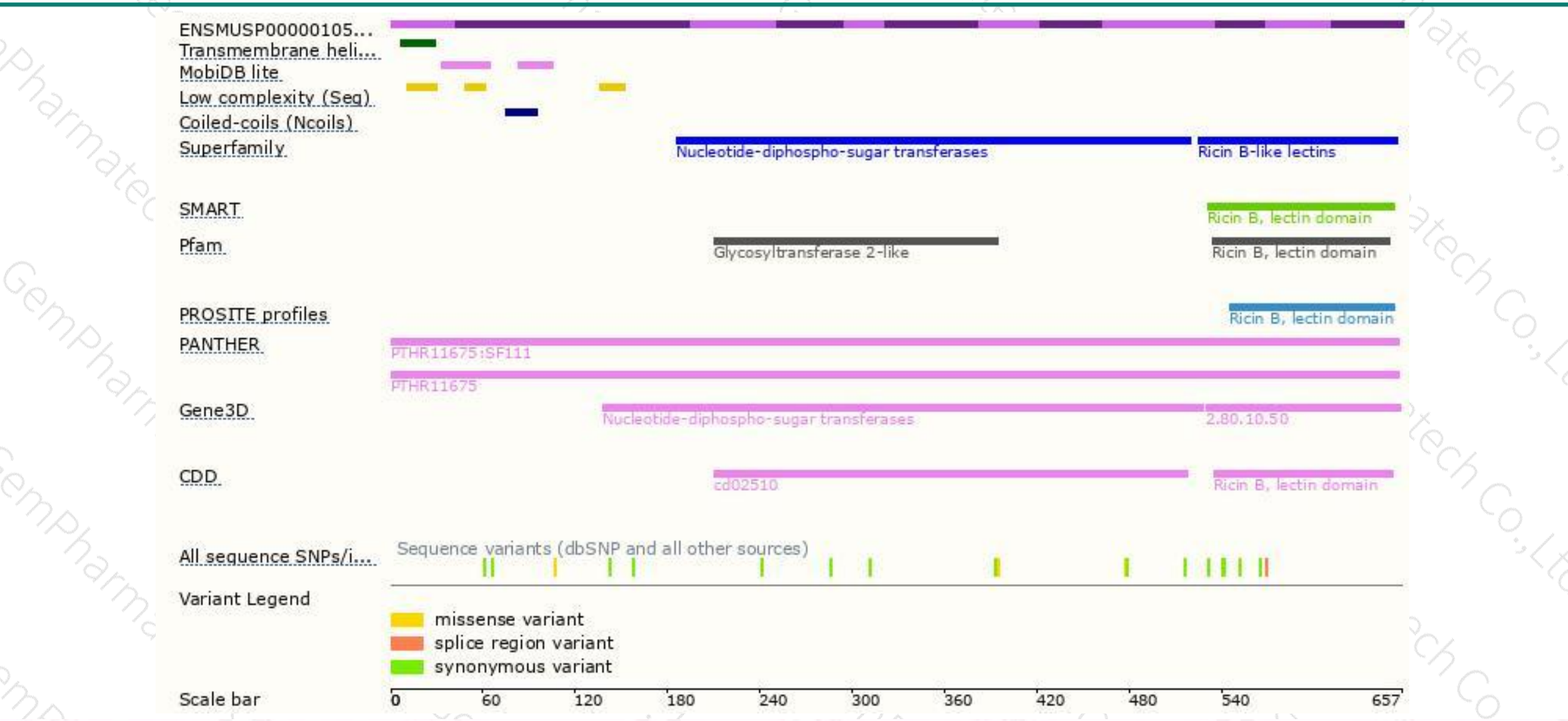
The strategy is based on the design of *Galnt7-201* transcript,the transcription is shown below:



Genomic location distribution



Protein domain



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

Tel: 400-9660890

