

***Galnt5* Cas9-CKO Strategy**

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Project Overview

Project Name

Galnt5

Project type

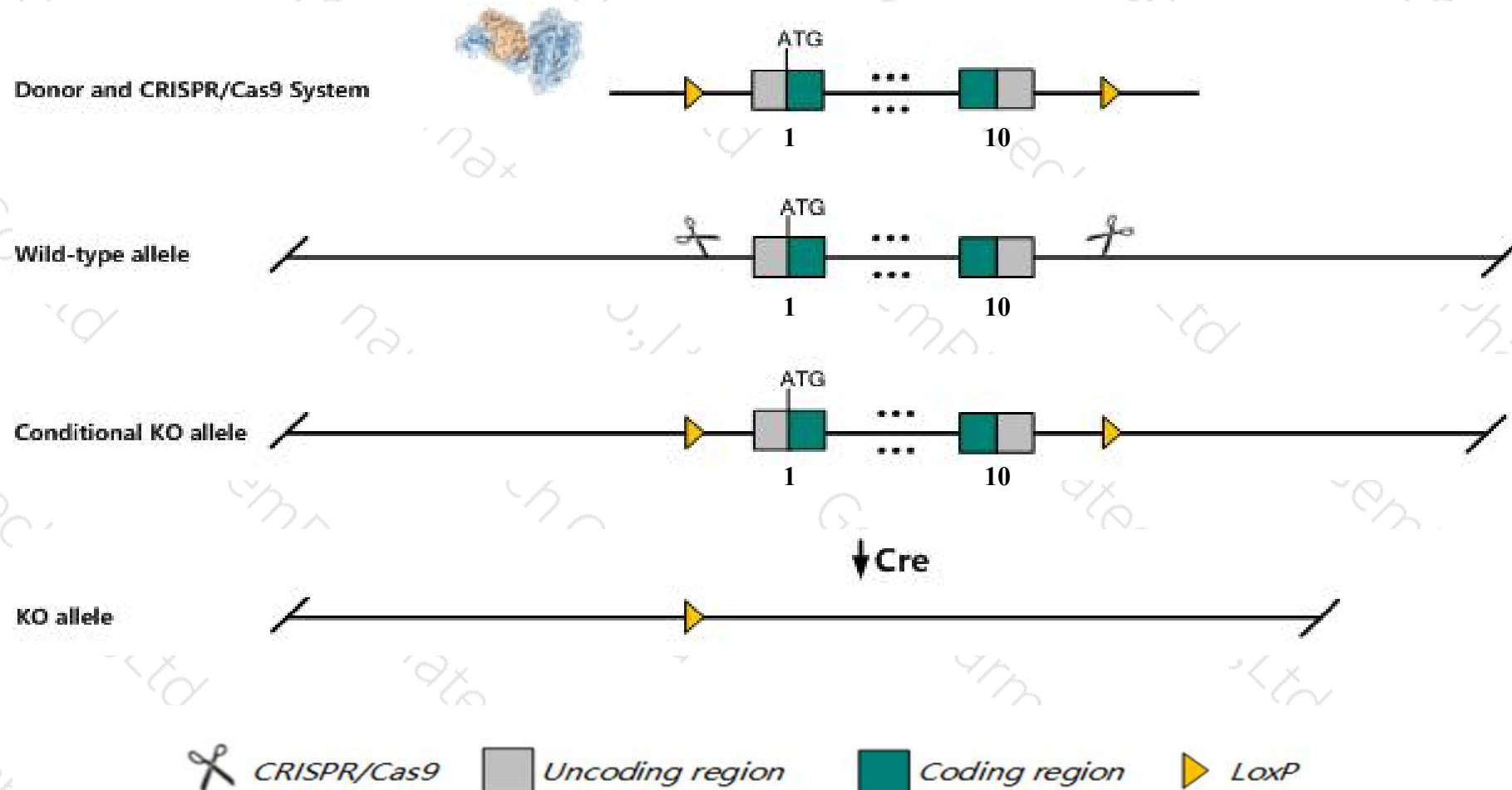
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Galnt5* gene. The schematic diagram is as follows:



Technical routes

- The *Galnt5* gene has 3 transcripts. According to the structure of *Galnt5* gene, exon1-exon10 of *Galnt5-201* (ENSMUST00000112616.7) transcript is recommended as the knockout region. The region contains all of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Galnt5* 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.

- According to the existing MGI data, An unpublished knockout mutation is reported to have no overt phenotypic consequences.
- The *Galnt5* gene is located on the Chr2. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)

Galnt5 polypeptide N-acetylgalactosaminyltransferase 5 [*Mus musculus* (house mouse)]

Gene ID: 241391, updated on 13-Mar-2020

Summary

Official Symbol	Galnt5 provided by MGI
Official Full Name	polypeptide N-acetylgalactosaminyltransferase 5 provided by MGI
Primary source	MGI:MGI:2179403
See related	Ensembl:ENSMUSG000000026828
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	4832424J23
Expression	Low expression observed in reference dataset See more
Orthologs	human all

Genomic context

Location: 2; 2 C1.1

Exon count: 11

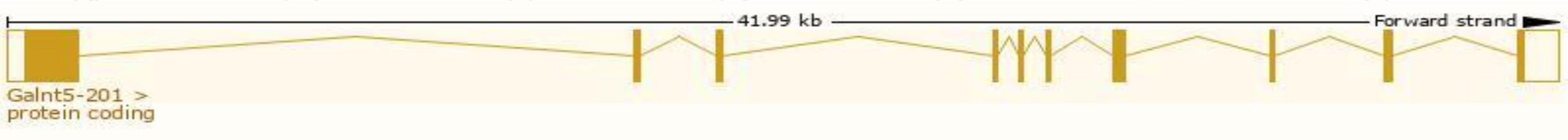
See Galnt5 in [Genome Data Viewer](#)

Transcript information (Ensembl)

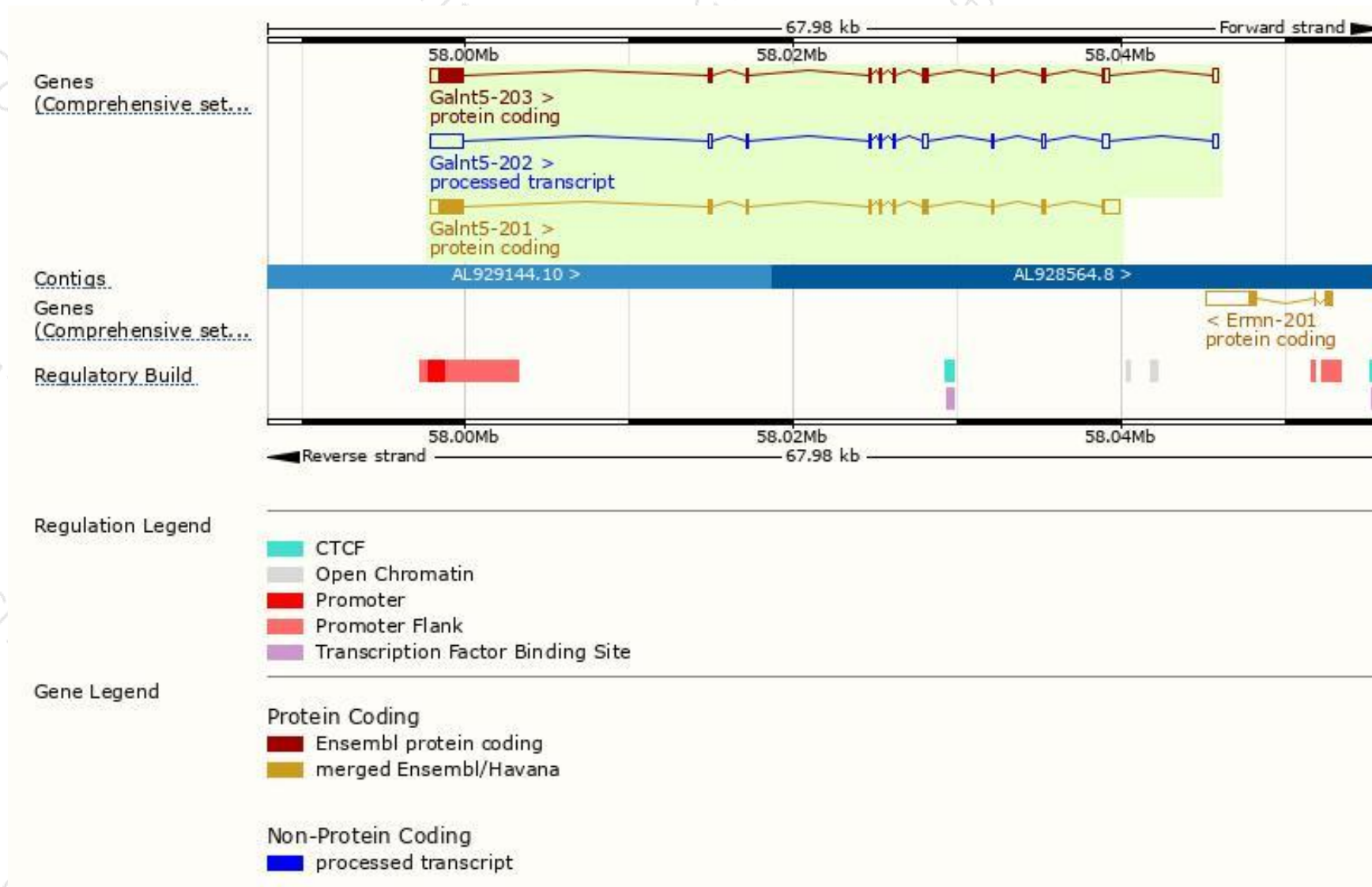
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Galnt5-201	ENSMUST00000112616.7	4225	930aa	Protein coding	CCDS38125	Q8C102	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 P1
Galnt5-203	ENSMUST00000166729.1	3812	930aa	Protein coding	CCDS38125	Q8C102	TSL:2 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 P1
Galnt5-202	ENSMUST00000144671.7	3812	No protein	Processed transcript	-	-	TSL:2

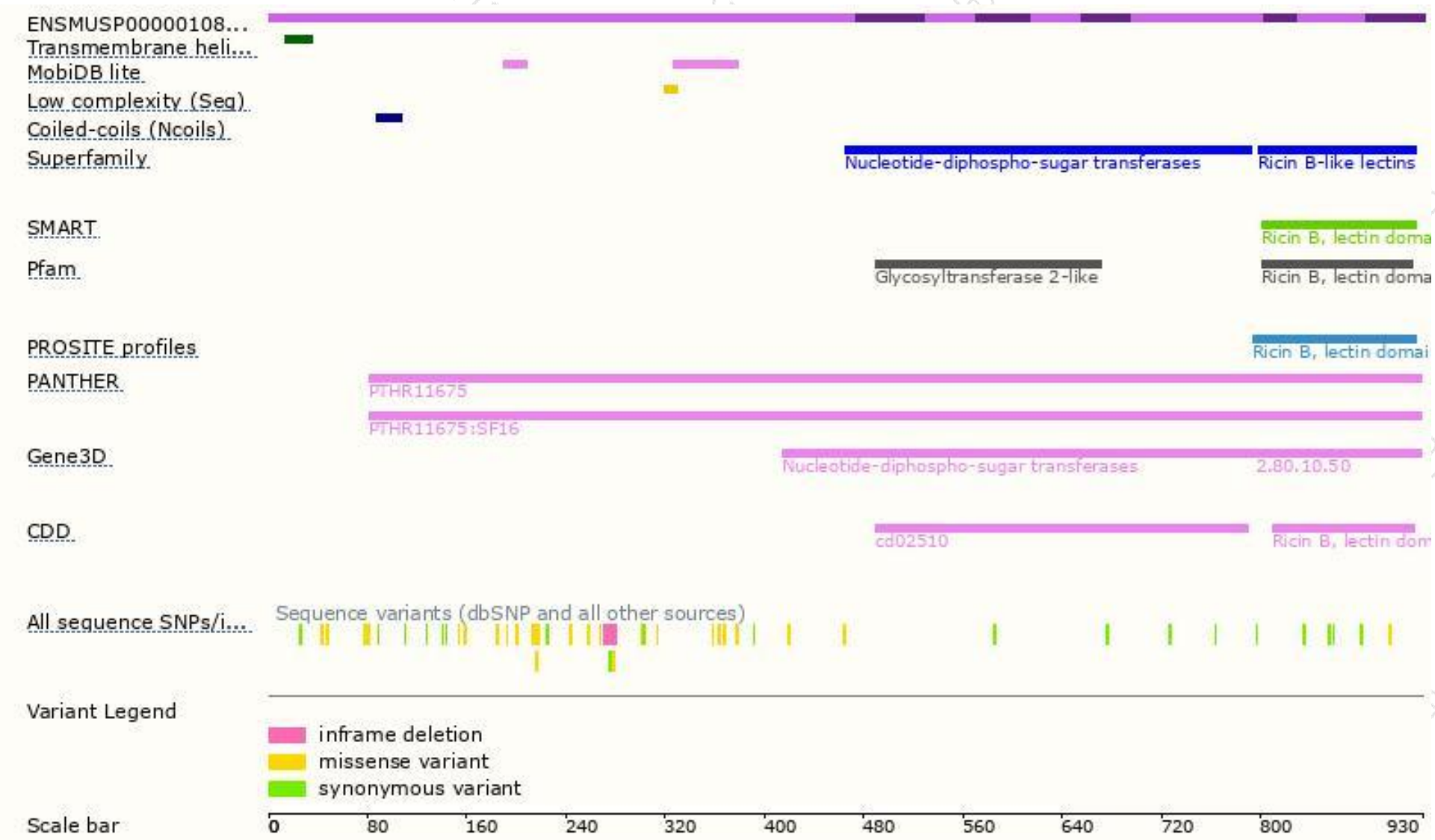
The strategy is based on the design of *Galnt5-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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