

Ganab Cas9-CKO Strategy

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Design Date:

2019-12-11

Project Overview

Project Name

Ganab

Project type

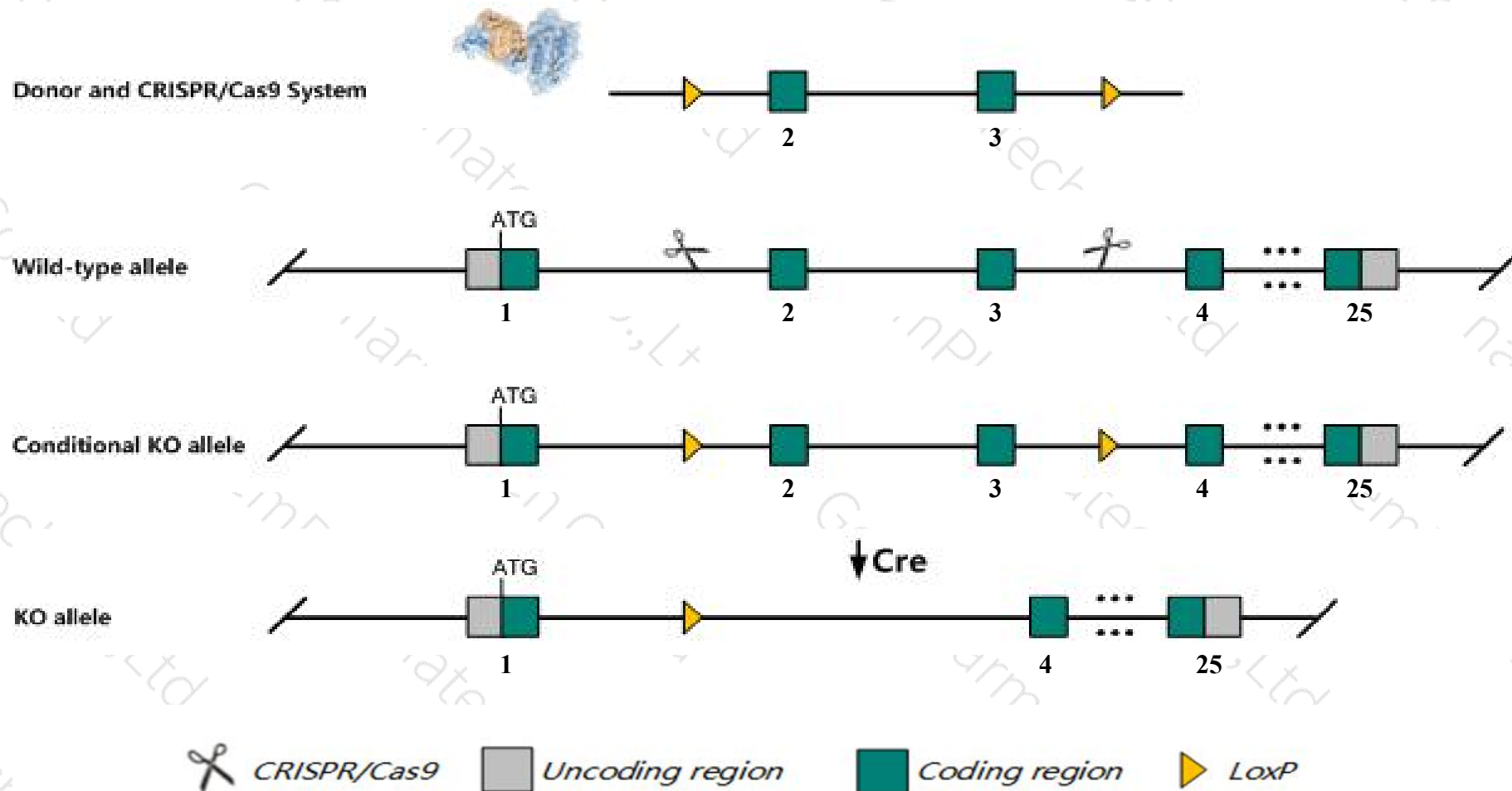
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

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



- The *Ganab* gene has 6 transcripts. According to the structure of *Ganab* gene, exon2-exon3 of *Ganab*-201 (ENSMUST00000096246.4) transcript is recommended as the knockout region. The region contains 214bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Ganab* 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.

Notice

- The *Ganab* gene is located on the Chr19. 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)

Ganab alpha glucosidase 2 alpha neutral subunit [Mus musculus (house mouse)]

Gene ID: 14376, updated on 3-Feb-2019

Summary



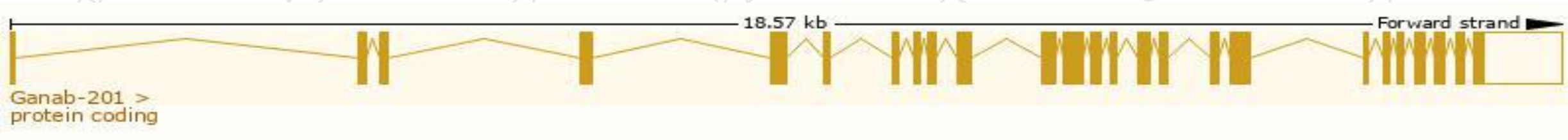
Official Symbol	Ganab provided by MGI
Official Full Name	alpha glucosidase 2 alpha neutral subunit provided by MGI
Primary source	MGI:MGI:1097667
See related	Ensembl:ENSMUSG00000071650
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	AU042638, G2an, Glull, mKIAA0088
Expression	Ubiquitous expression in limb E14.5 (RPKM 61.9), placenta adult (RPKM 49.8) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

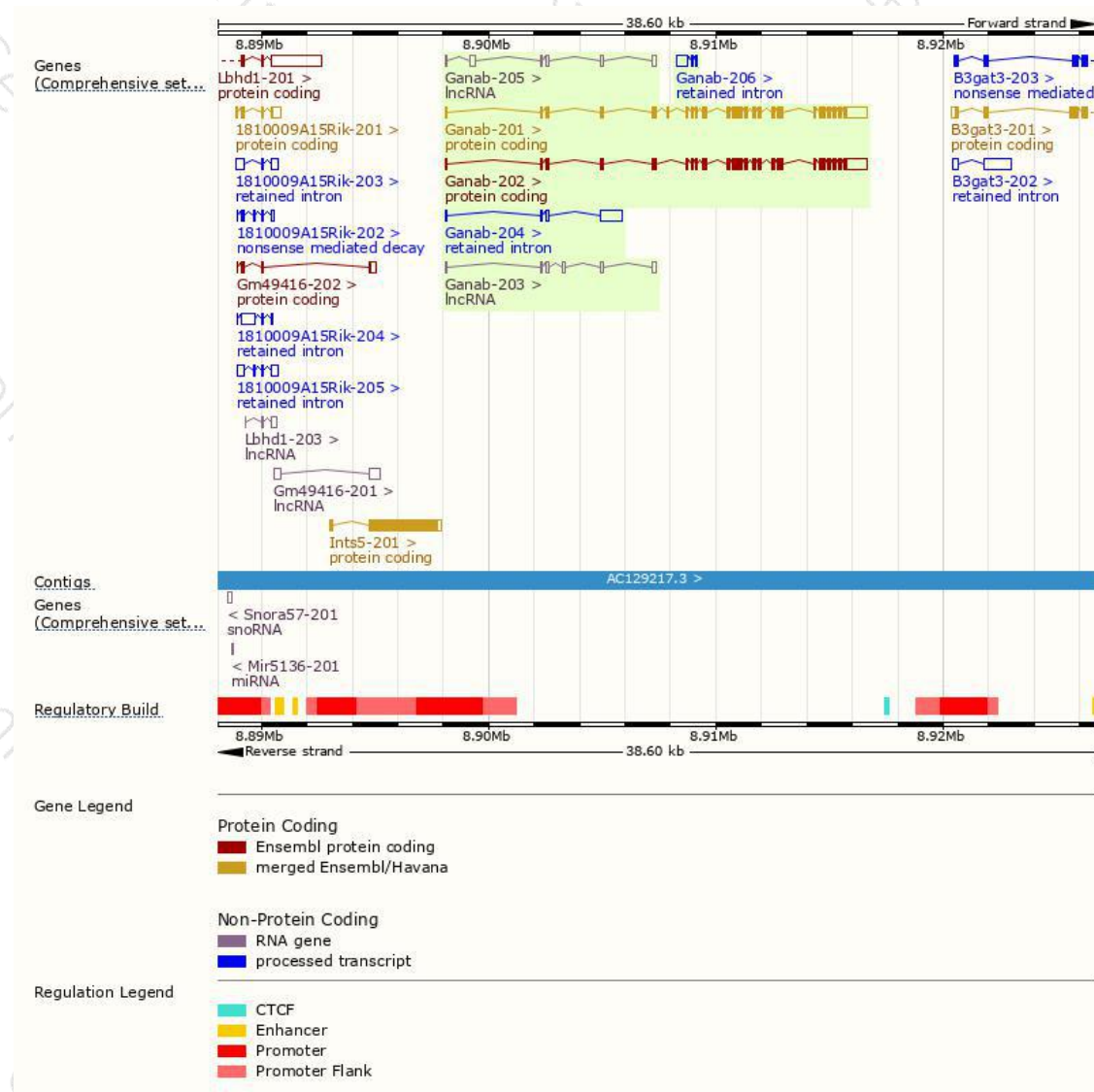
The gene has 6 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ganab-201	ENSMUST00000096246.4	3860	966aa	Protein coding	CCDS29557	A1A4T2 Q8BHN3	TSL:1 GENCODE basic APPRIS P2
Ganab-202	ENSMUST00000235274.1	3788	944aa	Protein coding	-	-	GENCODE basic APPRIS ALT 2
Ganab-204	ENSMUST00000236499.1	1226	No protein	Retained intron	-	-	
Ganab-206	ENSMUST00000238063.1	675	No protein	Retained intron	-	-	
Ganab-205	ENSMUST00000236563.1	808	No protein	lncRNA	-	-	
Ganab-203	ENSMUST00000235704.1	662	No protein	lncRNA	-	-	

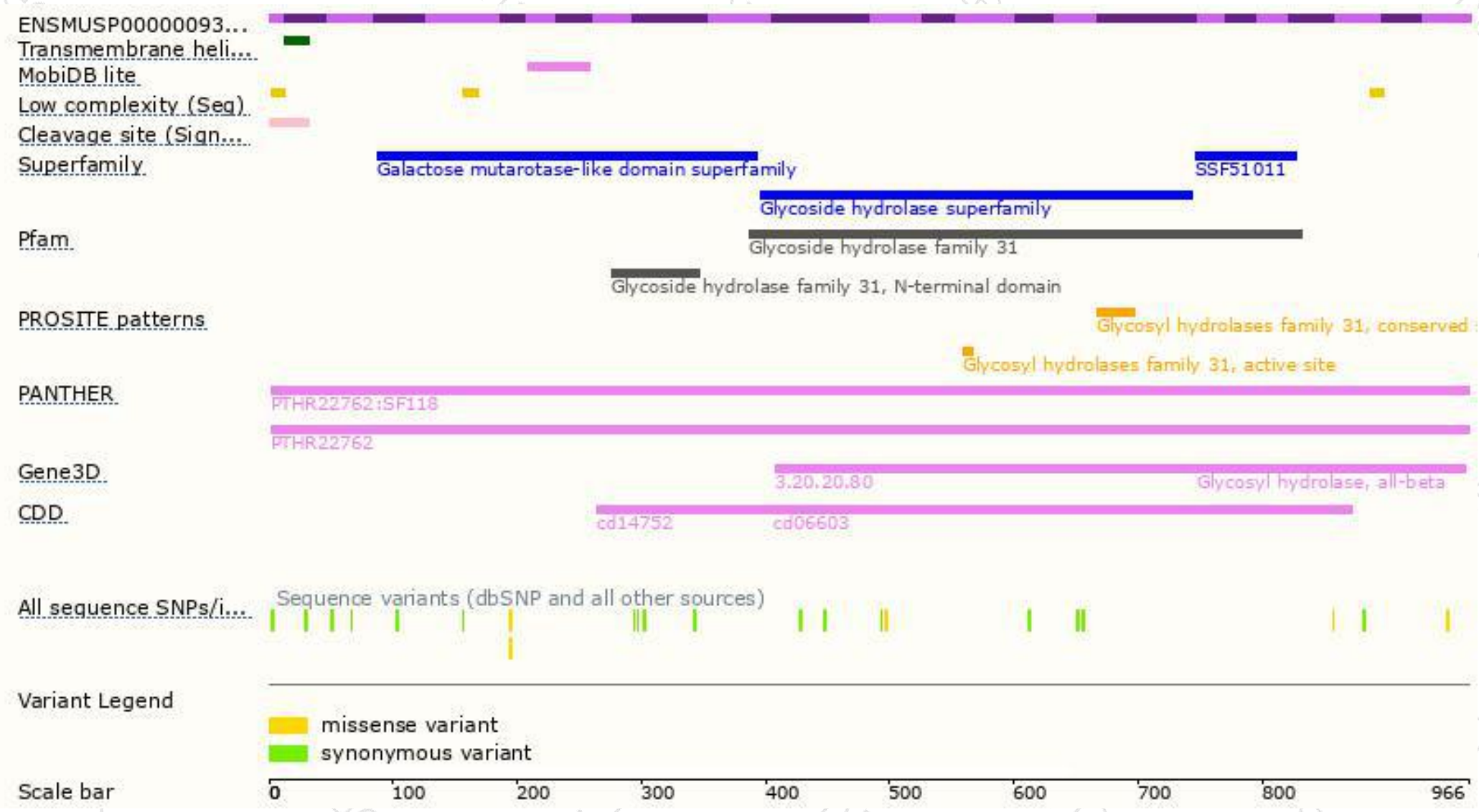
The strategy is based on the design of *Ganab-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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