

Itgb8 Cas9-KO Strategy

Designer:

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Design Date:

2019-10-18

Project Overview

Project Name

Itgb8

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Itgb8* gene. The schematic diagram is as follows:



- The *Itgb8* gene has 3 transcripts. According to the structure of *Itgb8* gene, exon2 of *Itgb8-201* (ENSMUST00000026360.8) transcript is recommended as the knockout region. The region contains 86bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Itgb8* gene. The brief process is as follows: CRISPR/Cas9 system v

- According to the existing MGI data, Mice homozygous for disruption of this gene either die before E11.5 as a result of circulatory abnormalities in the placenta or die within the first few days after birth and display intracerebral hemorrhaging.
- The *Itgb8* 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)

Itgb8 integrin beta 8 [Mus musculus (house mouse)]

Gene ID: 320910, updated on 26-Feb-2019

Summary



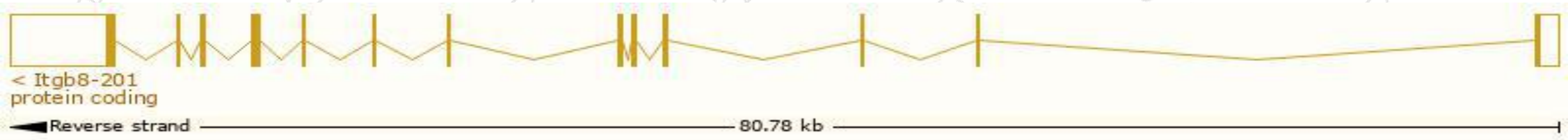
Official Symbol	Itgb8 provided by MGI
Official Full Name	integrin beta 8 provided by MGI
Primary source	MGI:MGI:1338035
See related	Ensembl:ENSMUSG00000025321
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	4832412O06Rik
Expression	Biased expression in kidney adult (RPKM 9.7), CNS E11.5 (RPKM 3.6) and 9 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

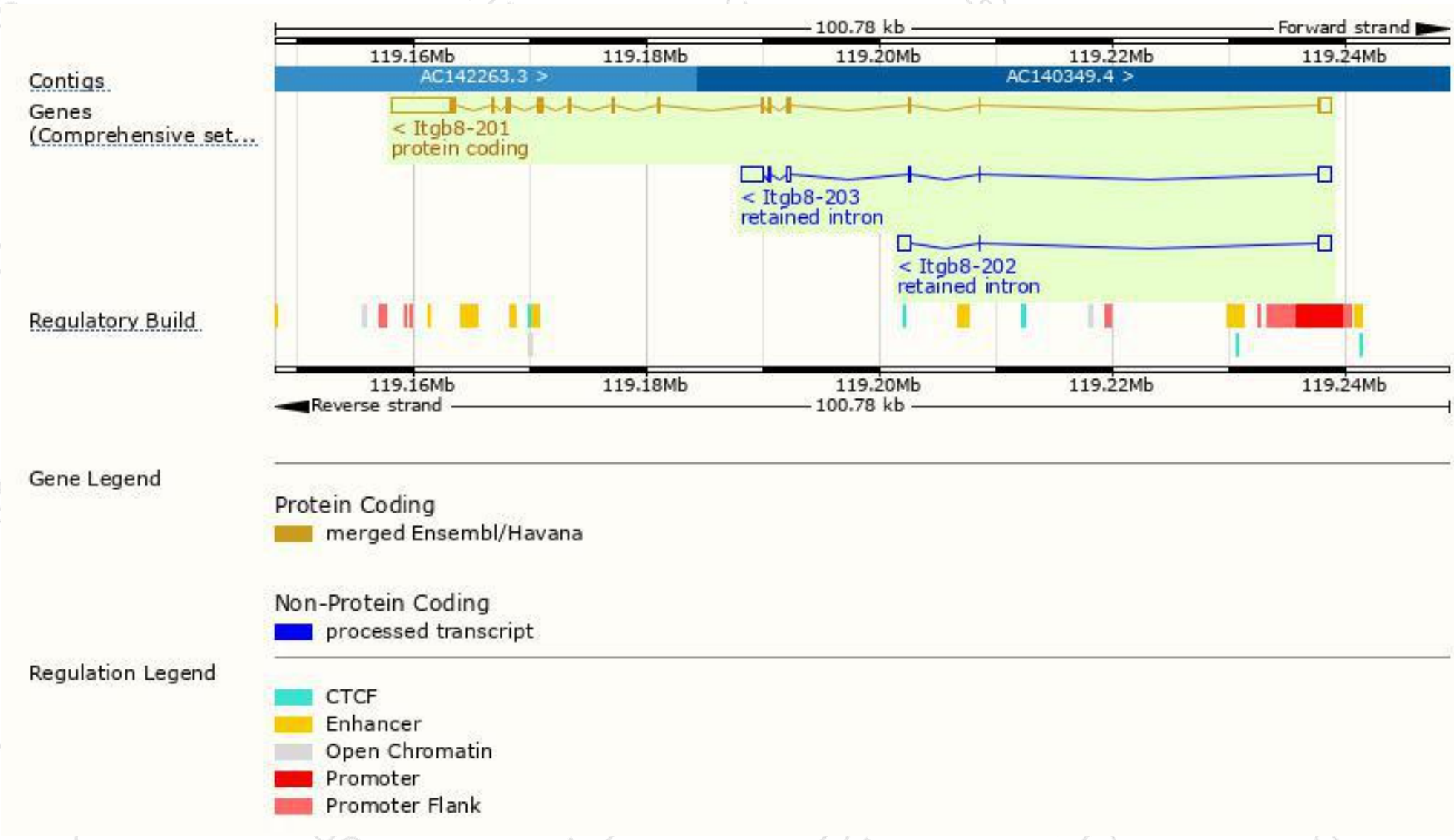
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Itgb8-201	ENSMUST00000026360.8	8403	767aa	Protein coding	CCDS36580	Q0VBD0	TSL:1 GENCODE basic APPRIS P1
Itgb8-203	ENSMUST00000151023.7	3676	No protein	Retained intron	-	-	TSL:1
Itgb8-202	ENSMUST00000137804.1	2304	No protein	Retained intron	-	-	TSL:1

The strategy is based on the design of *Itgb8-201* transcript,The transcription is shown below



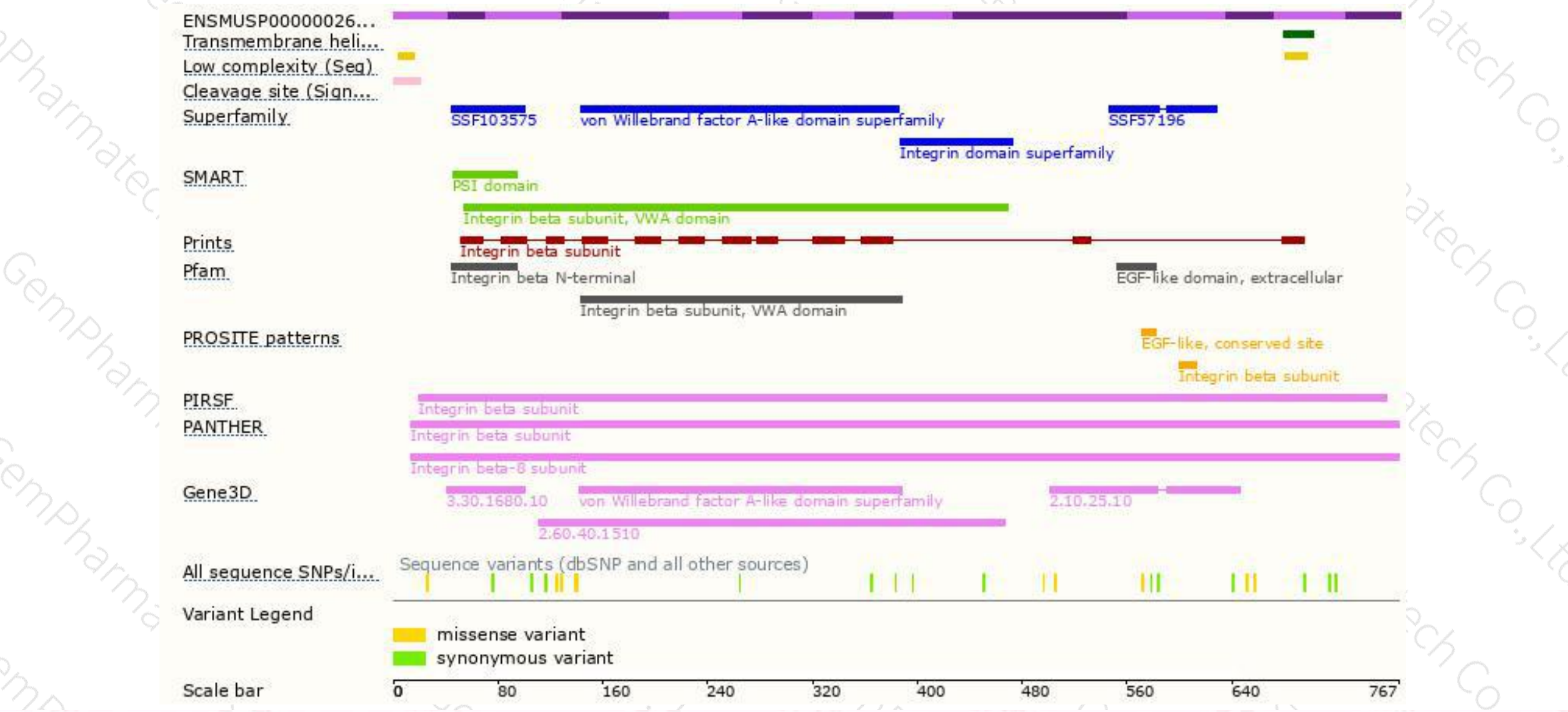
Genomic location distribution



Protein domain



集萃药康
GemPharmatech



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, Mice homozygous for disruption of this gene either die before E11.5 as a result of circulatory abnormalities in the placenta or die within the first few days after birth and display intracerebral hemorrhaging.

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

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