Prmt8 Cas9-KO Strategy

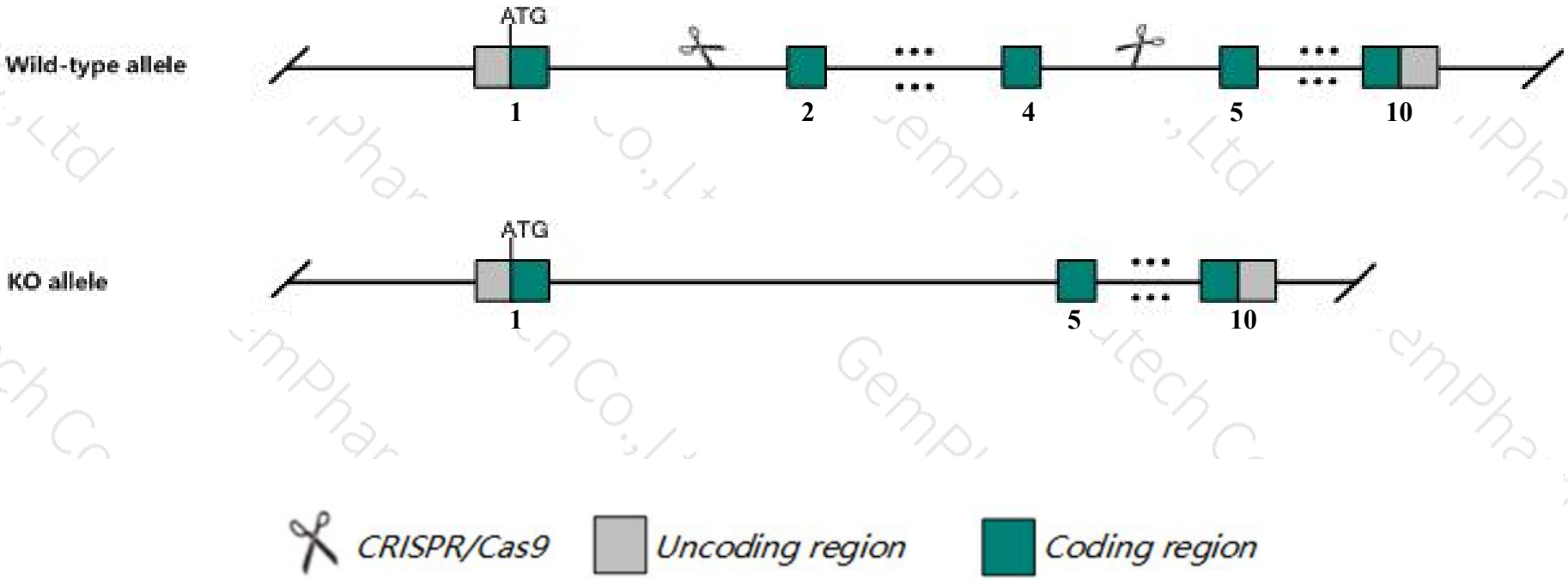
Designer:Xiaojing Li
Reviewer:JiaYu
Design Date:2020-2-12

Project Overview

Project Name	***Prmt8***
Project type	**Cas9-KO**
Strain background	**C57BL/6JGpt**

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Prmt8* gene. The schematic diagram is as follows:



Technical routes

➢ The *Prmt8* gene has 1 transcript. According to the structure of *Prmt8* gene, exon2-exon4 of *Prmt8-201* (ENSMUST00000032500.8) transcript is recommended as the knockout region.The region contains 406bp coding sequence. Knock out the region will result in disruption of protein function.

➢ In this project we use CRISPR/Cas9 technology to modify *Prmt8* gene. The brief process is as follows: CRISPR/Cas9 system

Notice

- According to the existing MGI data,Mice homozygous for a knockout allele exhibit abnormal Purkinje cell dendrite morphology, hyperactivity, limb grasping and gait abnormalities, and show reduced levels of acetylcholine and choline along with increased phosphatidylcholine levels in the cerebellum.

- The *Prmt8* gene is located on the Chr6. 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）

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Prmt8 protein arginine N-methyltransferase 8 [*Mus musculus* (house mouse)]

Gene ID: 381813, updated on 19-Oct-2019

Summary

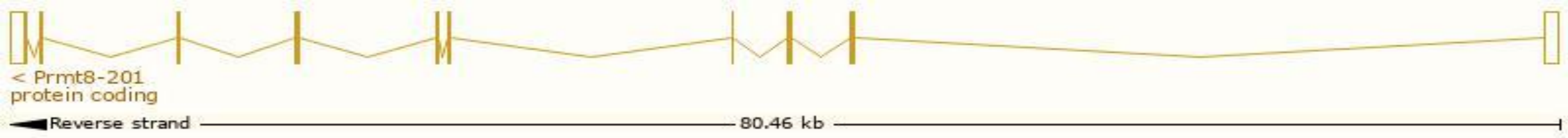
Official Symbol	Prmt8 provided by MGI
Official Full Name	protein arginine N-methyltransferase 8 provided by MGI
Primary source	MGI:MGI:3043083
See related	Ensembl:ENSMUSG00000030350
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Hrmt1l3; Hrmt1l4
Expression	Biased expression in cortex adult (RPKM 13.0), frontal lobe adult (RPKM 11.1) and 6 other tissues See more
Orthologs	human all

Transcript information（Ensembl）

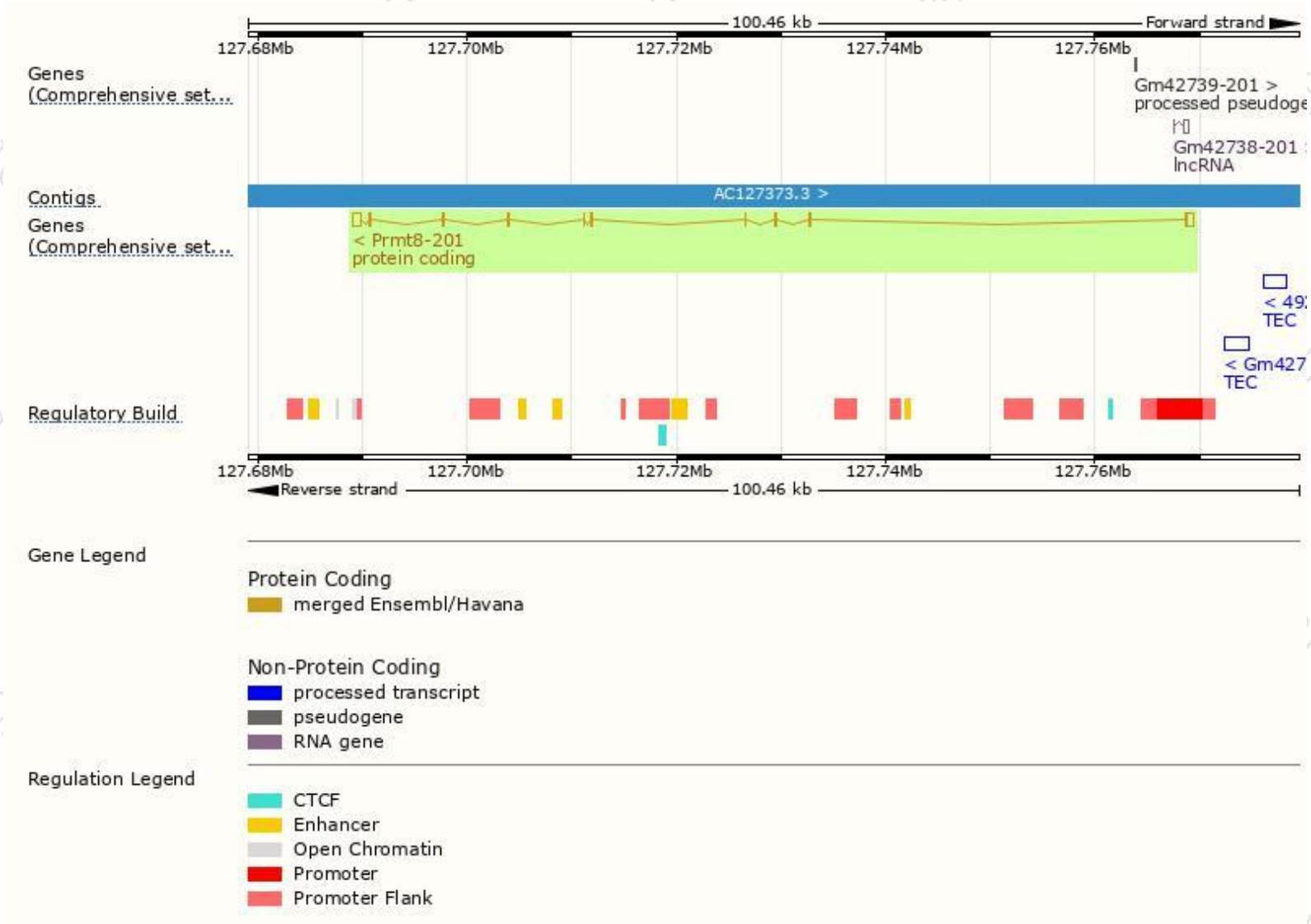
The gene has 1 transcript,and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Prmt8-201	ENSMUST00000032500.8	2724	394aa	Protein coding	CCDS57449	Q6PAK3	TSL:1 GENCODE basic APPRIS P1

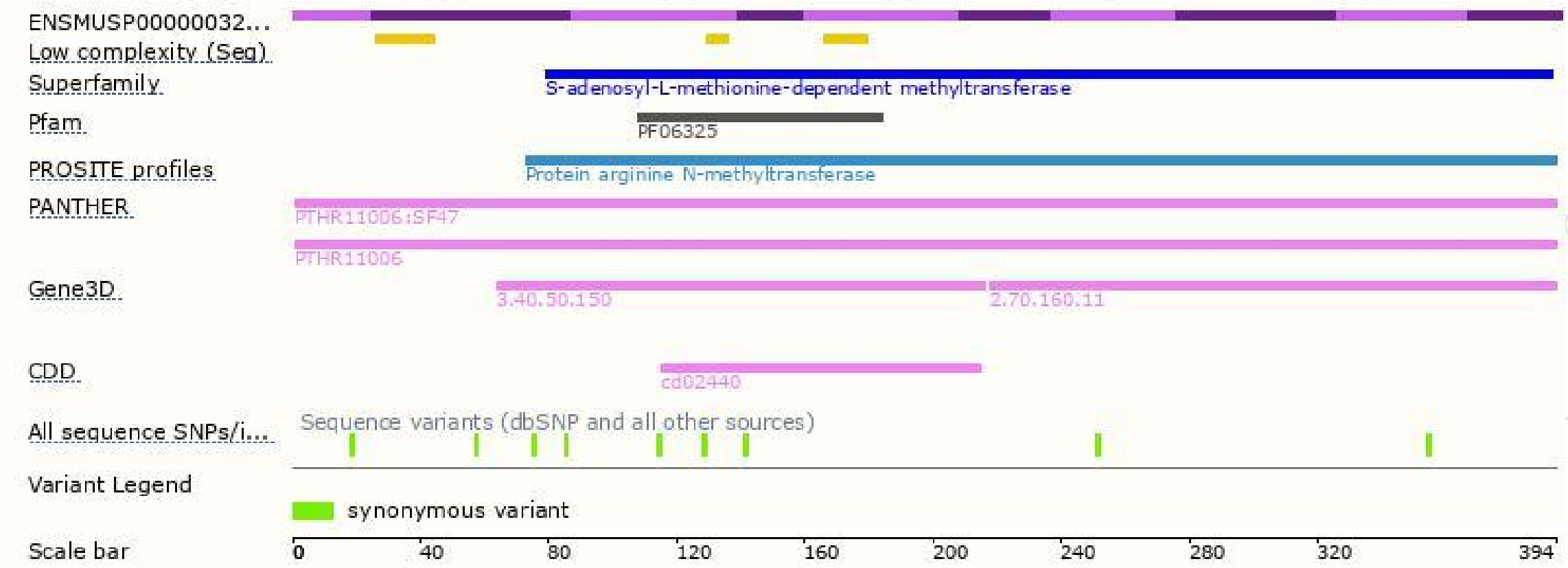
The strategy is based on the design of *Prmt8-201* transcript,The transcription is shown below



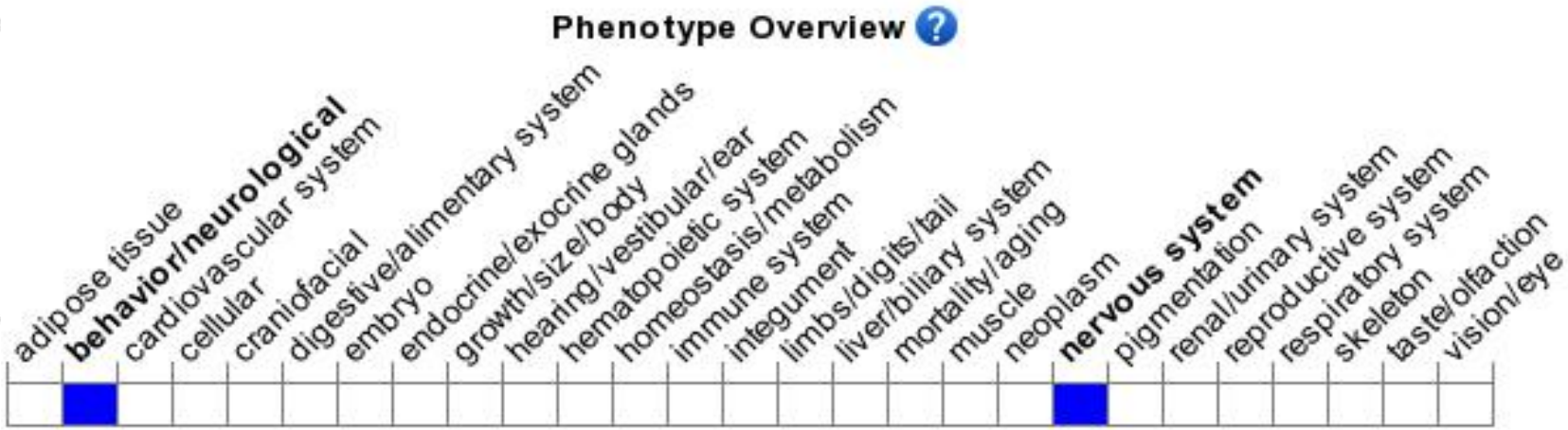
Genomic location distribution



Protein domain



Mouse phenotype description(MGI)



Phenotypes affected by the gene are marked in blue.Data quoted from MGI database(http://www.informatics.jax.org/).

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集萃药康
GemPharmatech

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

Tel: 400-9660890

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GemPharmatech Co.,Ltd