Crb2 Cas9-KO Strategy

Designer: Huan Fan

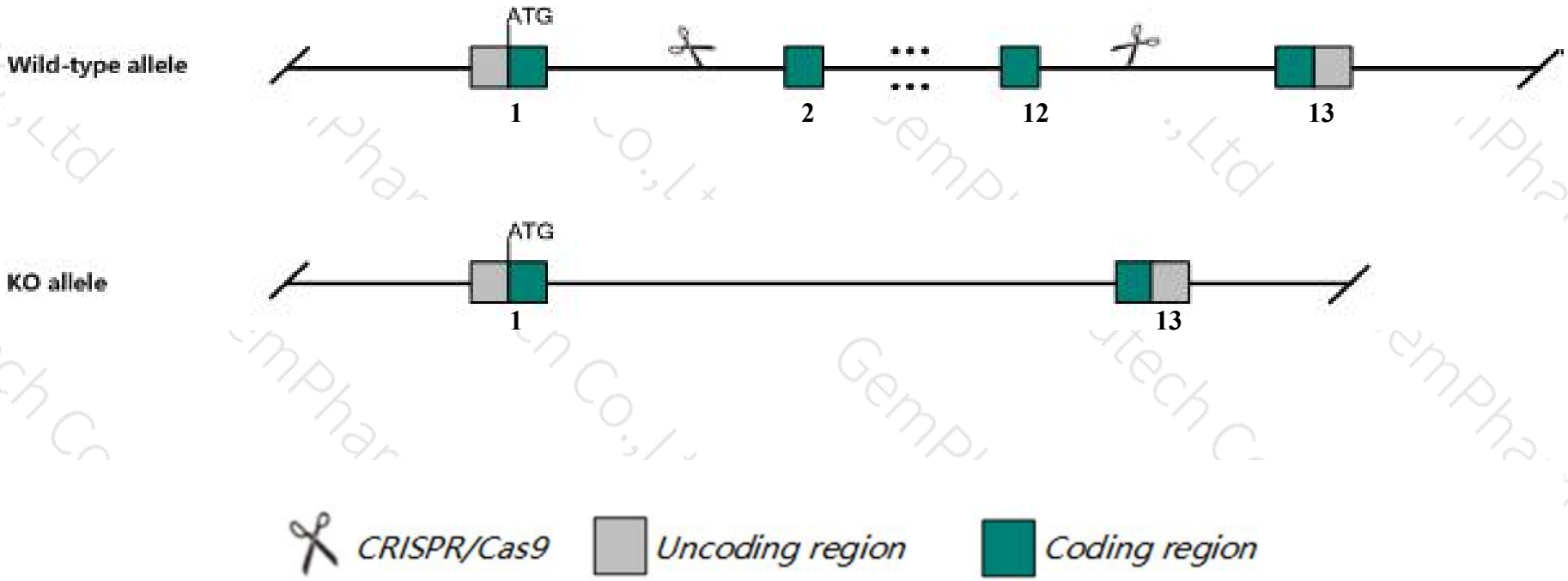
Design Date: 2019-7-25

Project Overview

Project Name	***Crb2***
Project type	**Cas9-KO**
Strain background	**C57BL/6JGpt**

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Crb2* gene. The schematic diagram is as follows:



Technical routes

➢ The *Crb2* gene has 3 transcripts. According to the structure of *Crb2* gene, exon2-exon12 of *Crb2-201* (ENSMUST00000050372.9) transcript is recommended as the knockout region.The region contains 3518bp coding sequence. Knock out the region will result in disruption of protein function.

➢ In this project we use CRISPR/Cas9 technology to modify *Crb2* gene. The brief process is as follows: CRISPR/Cas9 system v

Notice

- According to the existing MGI data,Homozygous inactivation of this gene causes severe gastrulation defects, impaired somitogenesis and organogenesis. and complete embryonic death by E12.5. Several organ primordia, including neuroepithelium, gut, and heart, fail to form properly.

- The *Crb2* 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information（NCBI）

集萃药康 GemPharmatech

Crb2 crumbs family member 2 [Mus musculus (house mouse)]

Gene ID: 241324, updated on 16-Mar-2019

Summary

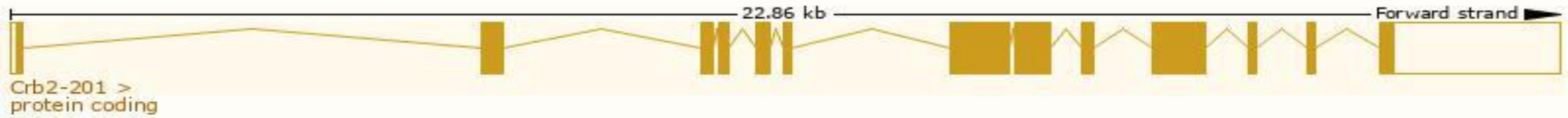
Official Symbol	Crb2 provided by MGI
Official Full Name	crumbs family member 2 provided by MGI
Primary source	MGI:MGI:2679260
See related	Ensembl:ENSMUSG00000035403
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	5930402A21, BC043114
Expression	Biased expression in CNS E11.5 (RPKM 6.1), whole brain E14.5 (RPKM 3.0) and 13 other tissues See more
Orthologs	human all

Transcript information（Ensembl）

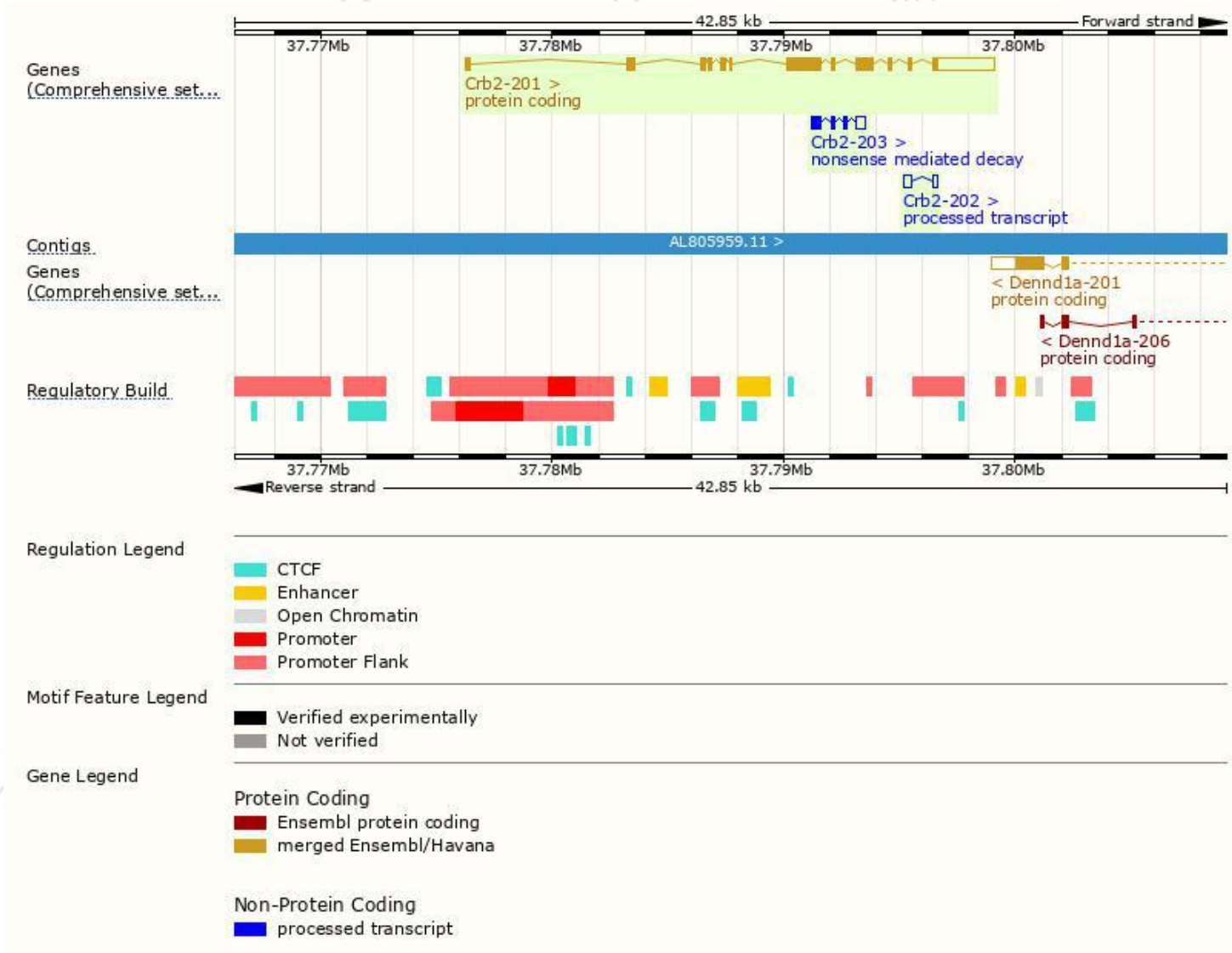
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Crb2-201	ENSMUST00000050372.9	6372	1282aa	Protein coding	CCDS50580	Q80YA8	TSL:5 GENCODE basic APPRIS P1
Crb2-203	ENSMUST00000147600.2	1132	197aa	Nonsense mediated decay	-	A0A0N4SUJ6	CDS 5' incomplete TSL:3
Crb2-202	ENSMUST00000137693.1	502	No protein	Processed transcript	-	-	TSL:5

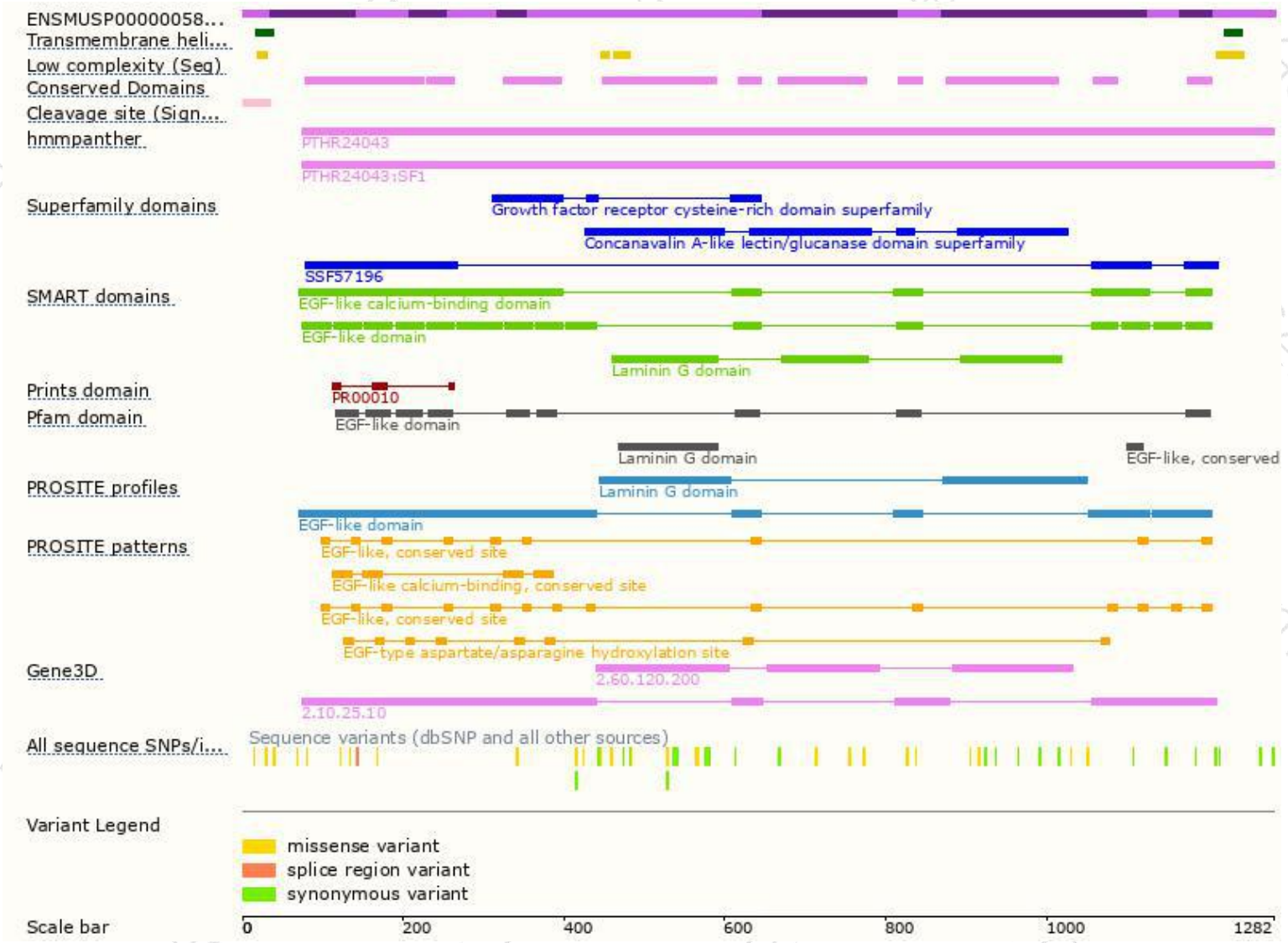
The strategy is based on the design of *Crb2-201* transcript,The transcription is shown below



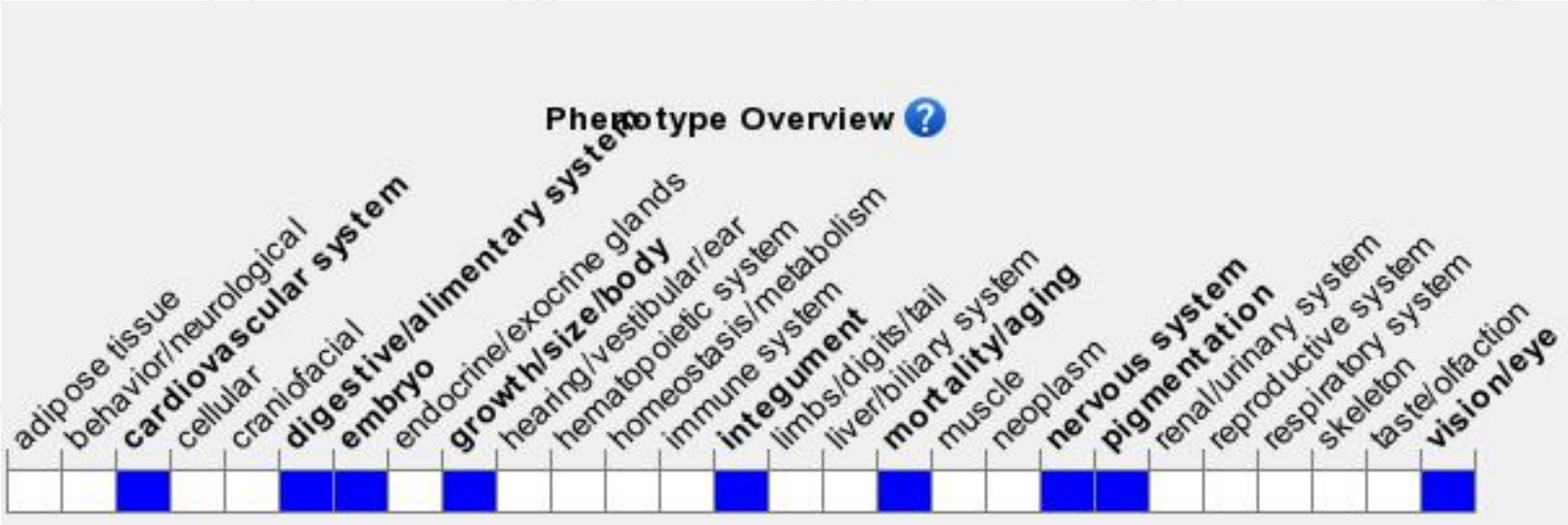
Genomic location distribution



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

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