

Crb2 Cas9-CKO Strategy

Designer:

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Design Date:

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Project Overview

Project Name

Crb2

Project type

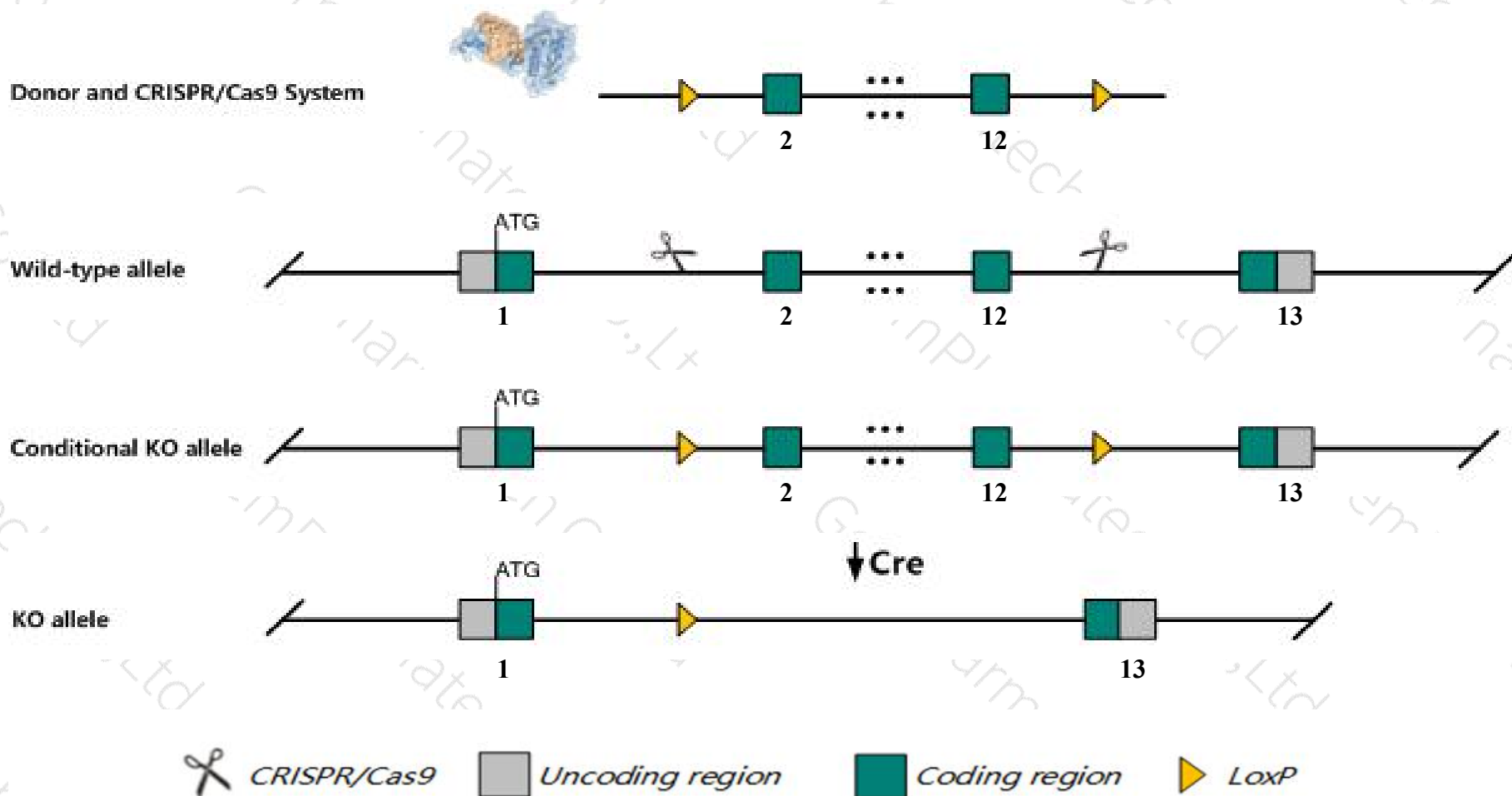
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

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



- 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 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, 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)

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

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

Summary



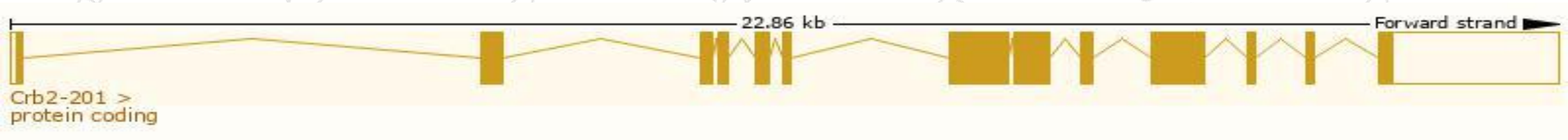
Official Symbol	Crb2 provided by MGI
Official Full Name	crumbs family member 2 provided by MGI
Primary source	MGI:MGI:2679260
See related	Ensembl:ENSMUSG00000035403
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; 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)

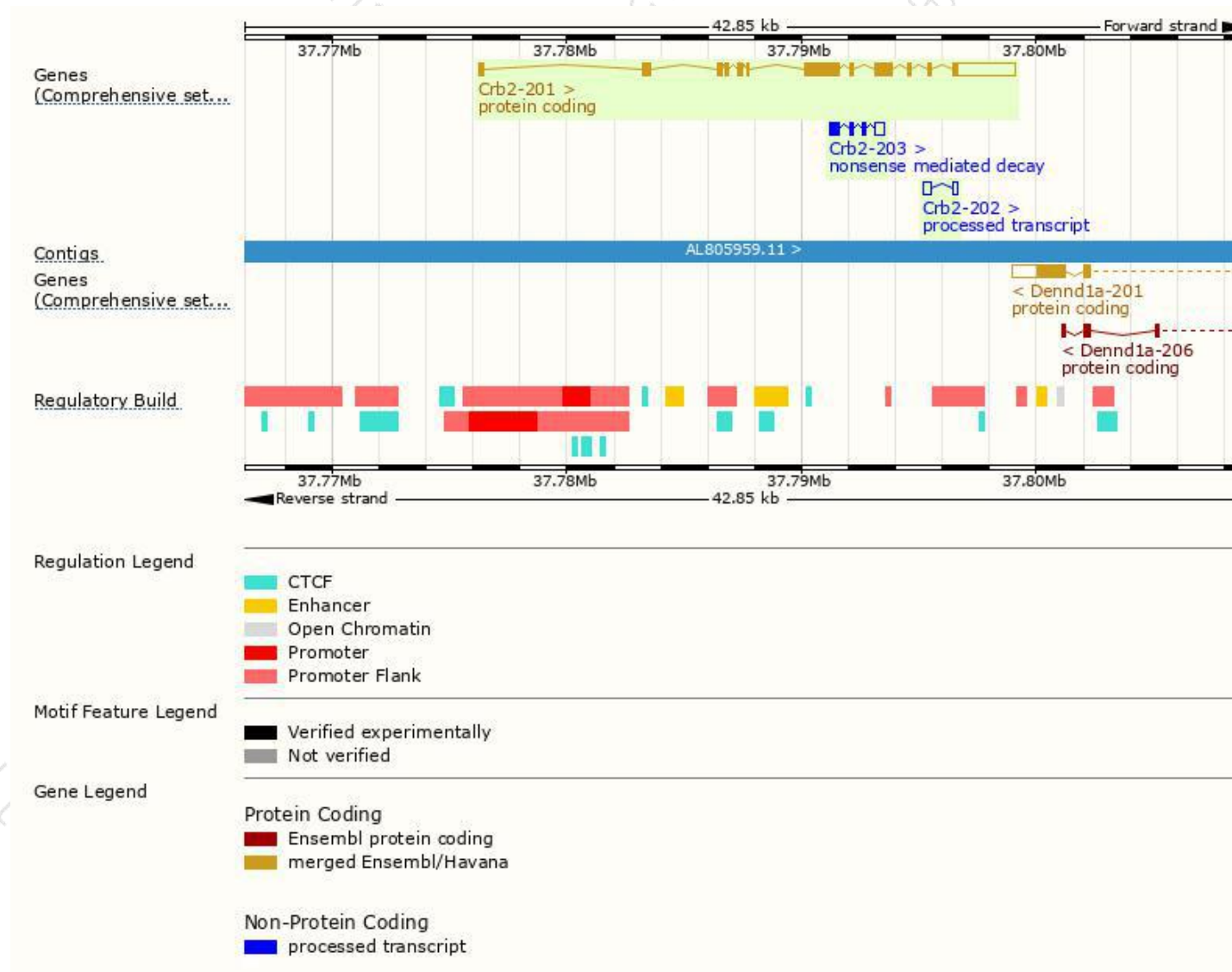
The gene has 3 transcripts,all transcripts are shown below:

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

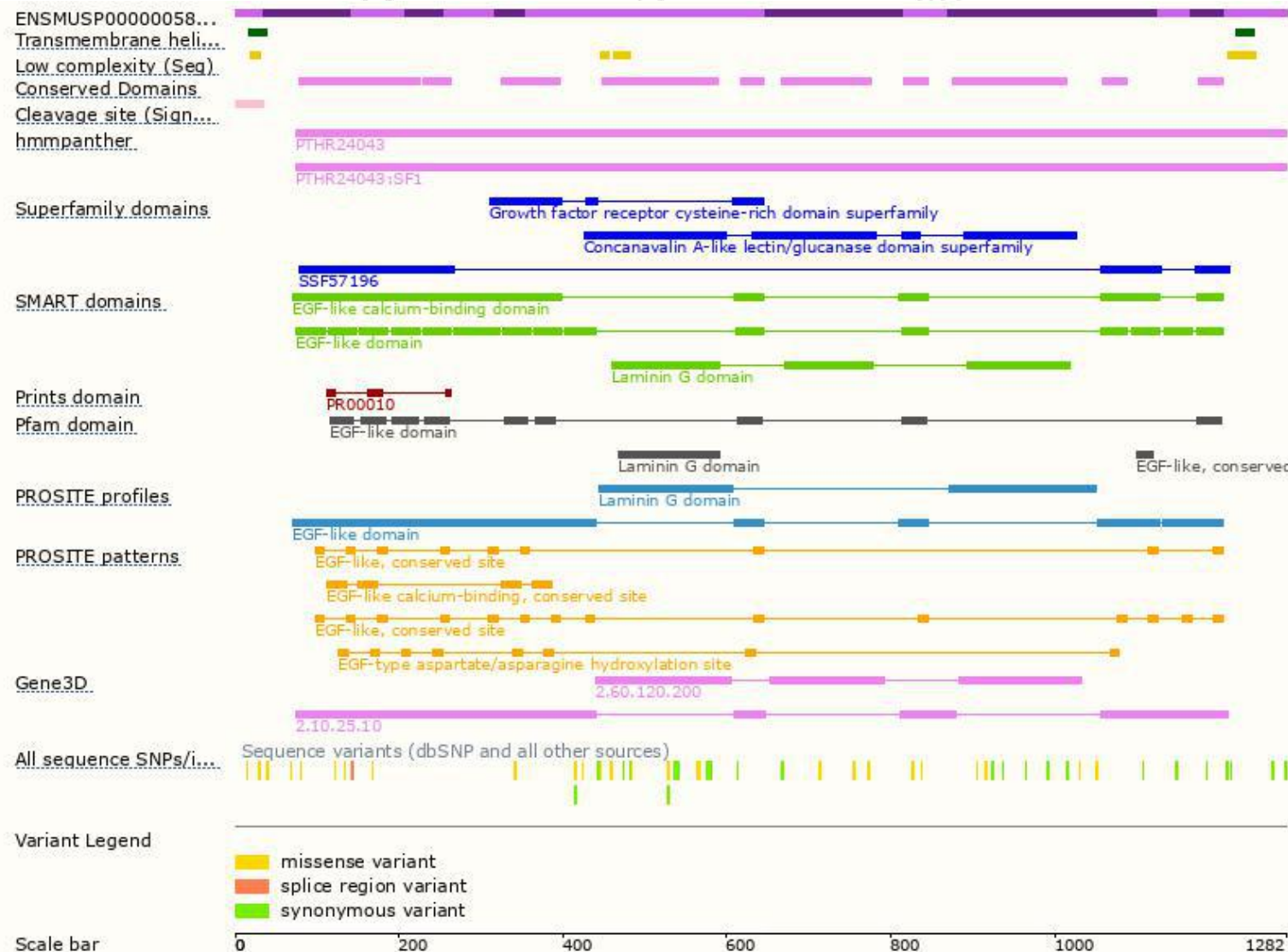
The strategy is based on the design of *Crb2-201* transcript,The transcription is shown below



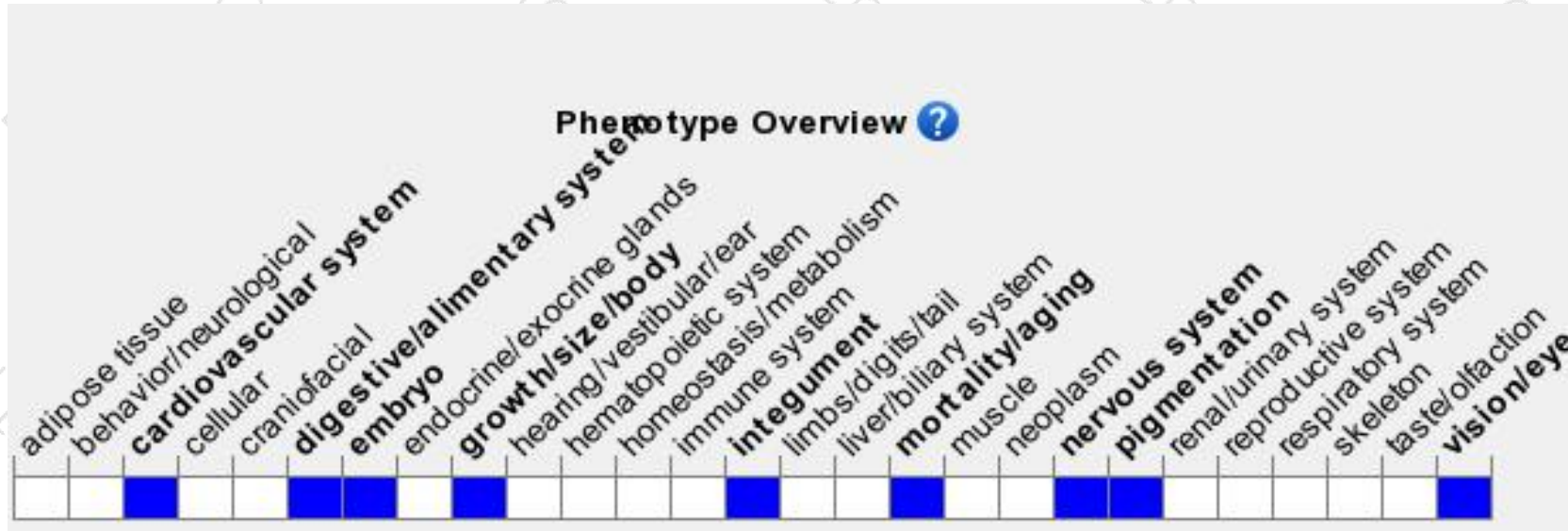
Genomic location distribution



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/>).

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.

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

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