

# ***Cnr1 Cas9-CKO Strategy***

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**Reviewer:**

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**Design Date:**

**2019-10-30**

# Project Overview

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**Project Name**

***Cnr1***

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**Project type**

**Cas9-CKO**

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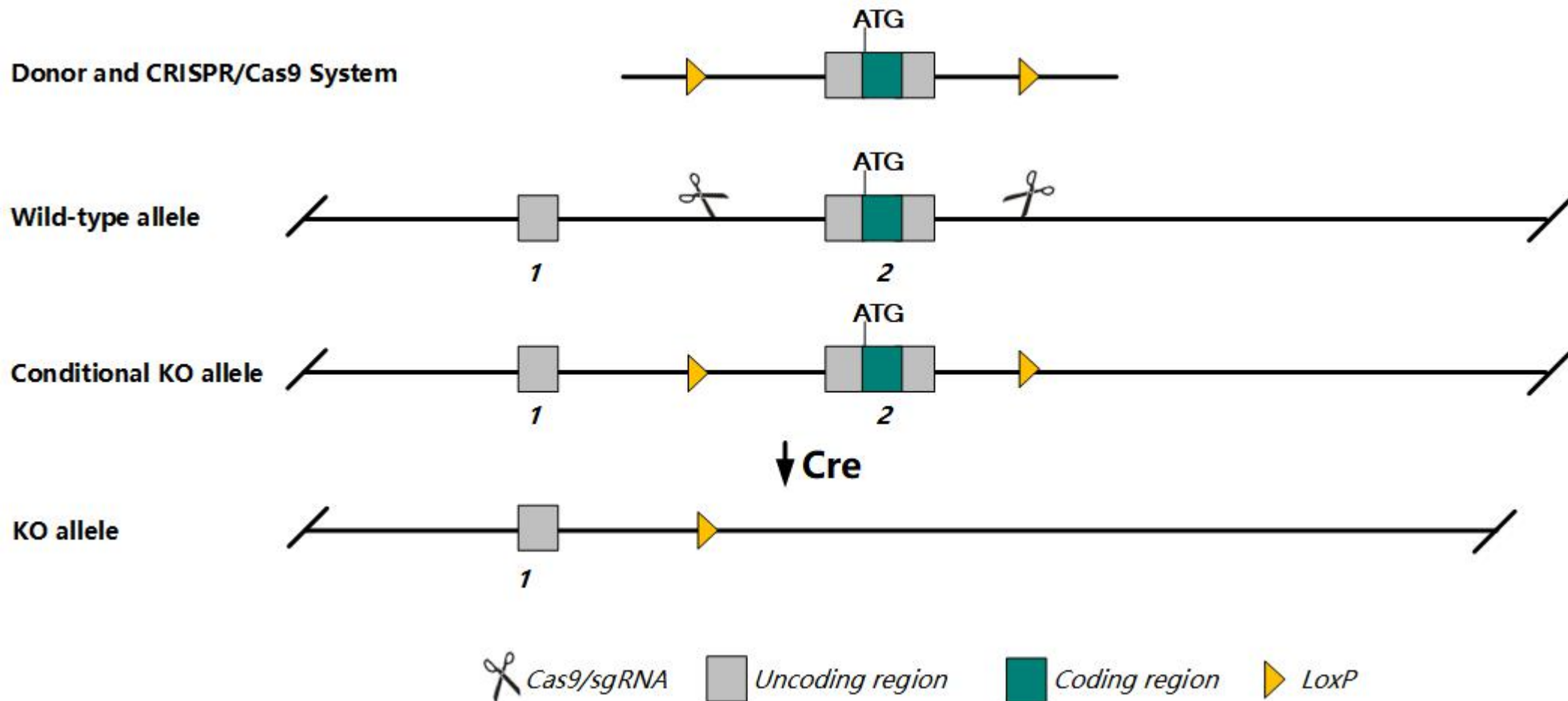
**Strain background**

**C57BL/6JGpt**

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# Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Cnr1* gene. The schematic diagram is as follows:



- The *Cnr1* gene has 3 transcripts. According to the structure of *Cnr1* gene, exon2 of *Cnr1*-201 (ENSMUST00000057188.6 ) transcript is recommended as the knockout region. The region contains all coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Cnr1* gene. The brief process is as follows: gRNA was transcribed in vitro, donor was constructed. Cas9, gRNA 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 or cell types.

- According to the existing MGI data , Mice homozygous for a null allele exhibit abnormal behaviors, altered long term depression and susceptibility to induced seizure.
- The *Cnr1* gene is located on the Chr4. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.



# Gene information ( NCBI )

## Cnr1 cannabinoid receptor 1 (brain) [ *Mus musculus* (house mouse) ]

Gene ID: 12801, updated on 29-Oct-2019

### Summary



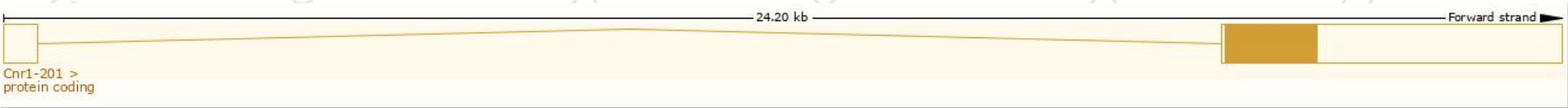
Official Symbol	Cnr1 provided by <a href="#">MGI</a>
Official Full Name	cannabinoid receptor 1 (brain) provided by <a href="#">MGI</a>
Primary source	<a href="#">MGI:MGI:104615</a>
See related	<a href="#">Ensembl:ENSMUSG00000044288</a>
Gene type	protein coding
RefSeq status	VALIDATED
Organism	<a href="#">Mus musculus</a>
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	CB1; CB-R; CB1A; CB1B; CB1R
Expression	Biased expression in cerebellum adult (RPKM 24.0), CNS E18 (RPKM 17.4) and 7 other tissues <a href="#">See more</a>
Orthologs	<a href="#">human</a> <a href="#">all</a>

# Transcript information ( Ensembl )

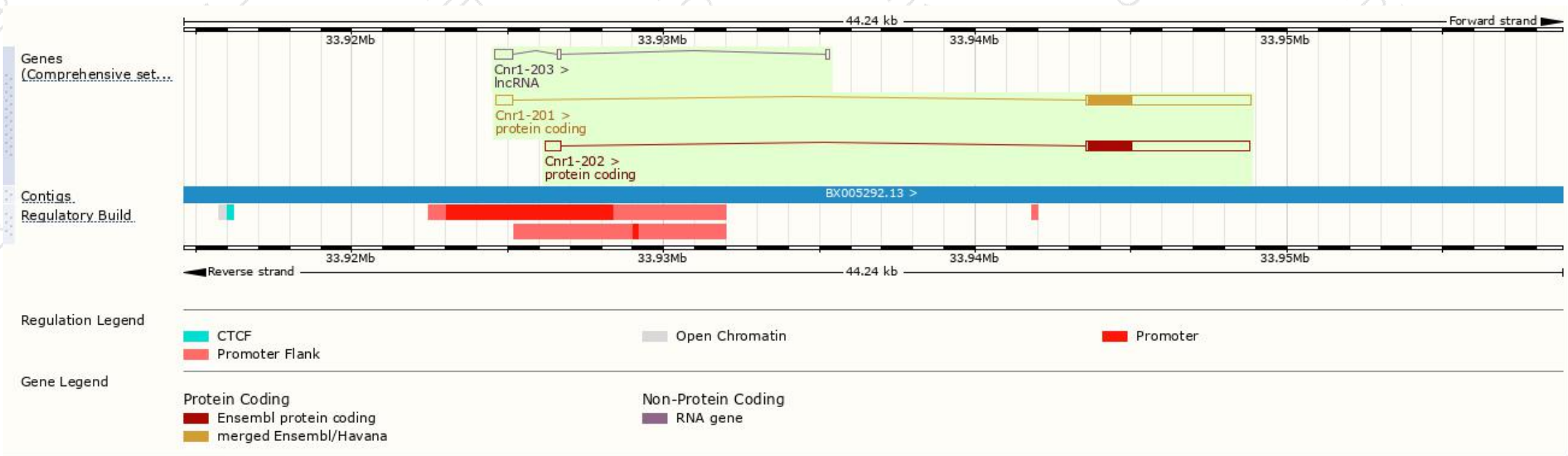
The gene has 3 transcripts, and all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Cnr1-203	<a href="#">ENSMUST00000133462.1</a>	757	No protein	lncRNA	-	-	TSL:3
Cnr1-202	<a href="#">ENSMUST00000084736.2</a>	5761	<a href="#">473aa</a>	Protein coding	<a href="#">CCDS18025</a>	<a href="#">A0A385KNU8</a> <a href="#">P47746</a>	TSL:2 GENCODE basic APPRIS P1
Cnr1-201	<a href="#">ENSMUST00000057188.6</a>	5807	<a href="#">473aa</a>	Protein coding	<a href="#">CCDS18025</a>	<a href="#">A0A385KNU8</a> <a href="#">P47746</a>	TSL:1 GENCODE basic APPRIS P1

The strategy is based on the design of *Cnr1*-201 transcript, The transcription is shown below



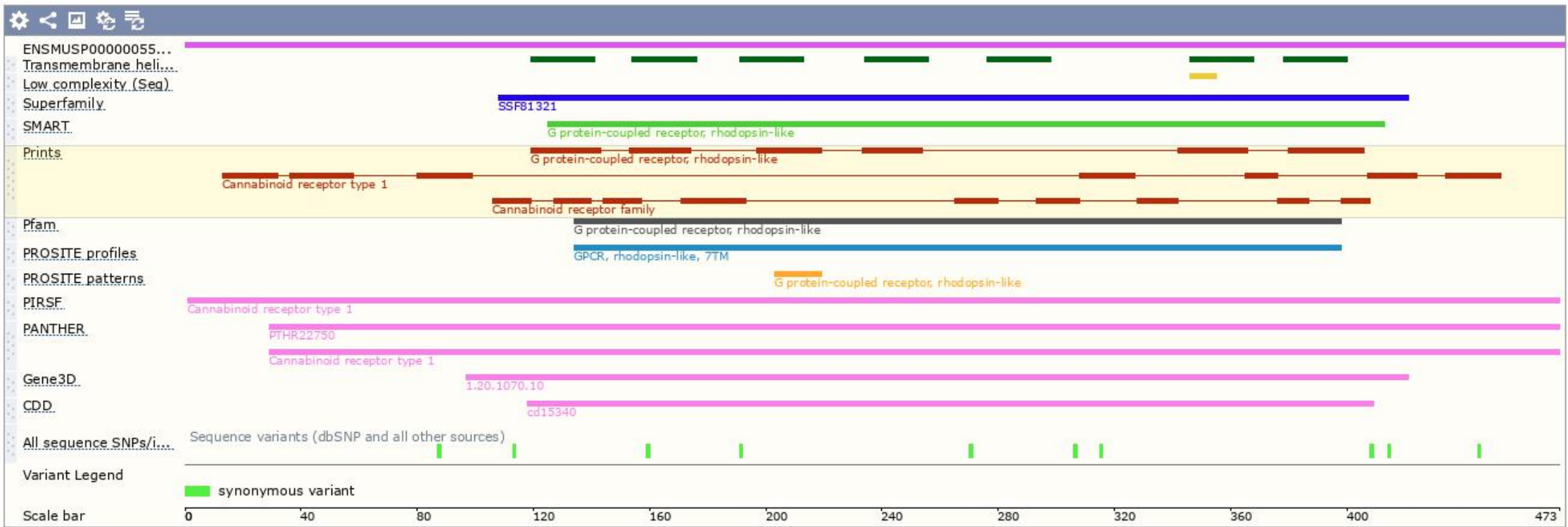
# Genomic location distribution



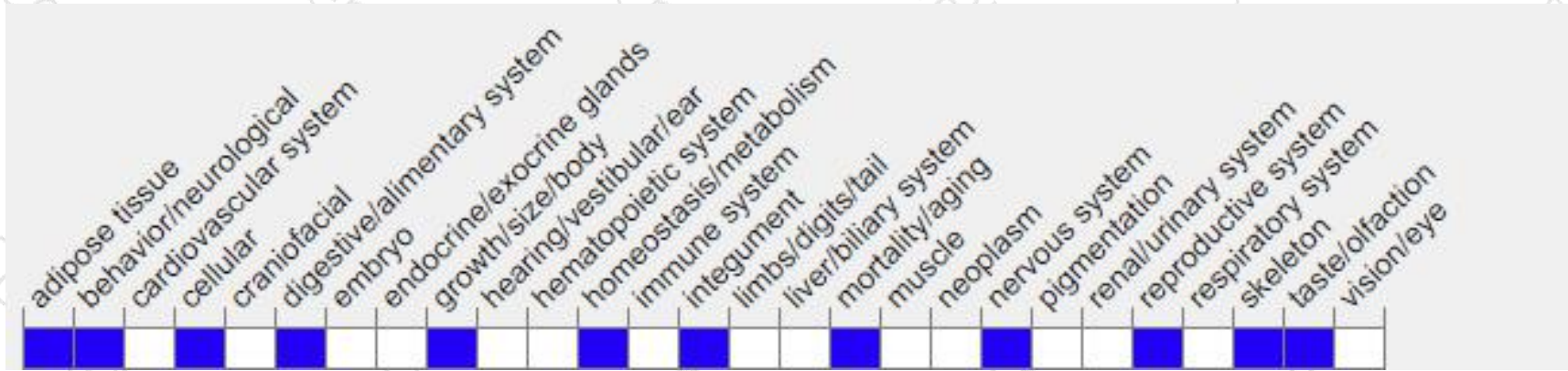


# Protein domain

Protein domains for ENSMUSP00000055797.6



# Mouse phenotype description(MGI)



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

Mice homozygous for a null allele exhibit abnormal behaviors, altered long term depression and susceptibility to induced seizure.

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