

Adcyl Cas9-CKO Strategy

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Project Overview



Project Name

Adcy1

Project type

Cas9-CKO

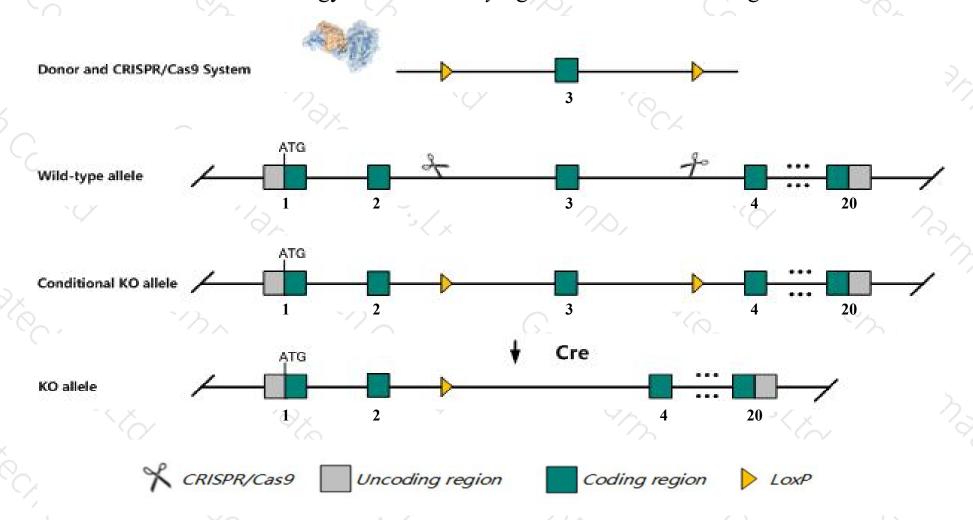
Strain background

C57BL/6JGpt

Conditional Knockout strategy



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



Technical routes



- The *Adcy1* gene has 2 transcripts. According to the structure of *Adcy1* gene, exon3 of *Adcy1-201*(ENSMUST00000020706.4) transcript is recommended as the knockout region. The region contains 119bp coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Adcy1* 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.

Notice



- > According to the existing MGI data, Mice homozygous for an insertional or null mutation fail to develop normal patterned distribution of neurons in the brain and display behavioral and learning abnormalities.
- The *Adcy1* gene is located on the Chr11. 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)



Adcy1 adenylate cyclase 1 [Mus musculus (house mouse)]

Gene ID: 432530, updated on 12-Aug-2019

Summary

△ ?

Official Symbol Adcy1 provided by MGI

Official Full Name adenylate cyclase 1 provided by MGI

Primary source MGI:MGI:99677

See related Ensembl: ENSMUSG00000020431

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 AC1; brl; I-AC; D11Bwg1392e

Expression Biased expression in cerebellum adult (RPKM 78.9), cortex adult (RPKM 50.8) and 4 other tissues See more

Orthologs human all

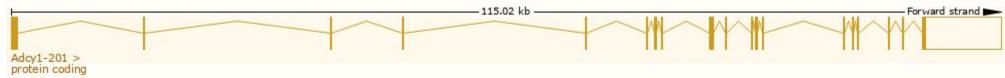
Transcript information (Ensembl)



The gene has 2 transcripts, all transcripts are shown below:

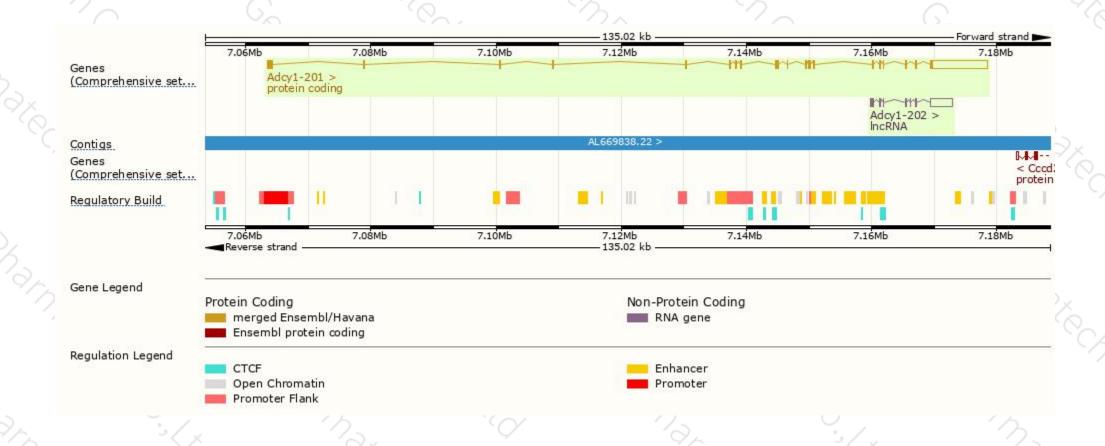
Name 🌲	Transcript ID 👙	bp 🌲	Protein 4	Biotype 🍦	CCDS	UniProt 👙	Flags		
Adcy1-201	ENSMUST00000020706.4	12259	<u>1118aa</u>	Protein coding	CCDS24426 ₽	<u>088444</u> ₽	TSL:2	GENCODE basic	APPRIS P1
Adcy1-202	ENSMUST00000135398.1	4333	No protein	I IncRNA	-	-		TSL:1	

The strategy is based on the design of Adcy1-201 transcript, The transcription is shown below



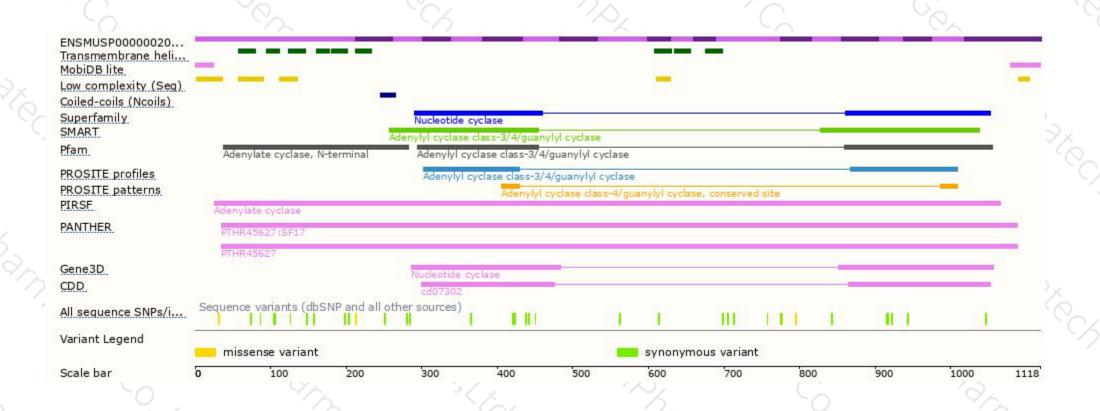
Genomic location distribution





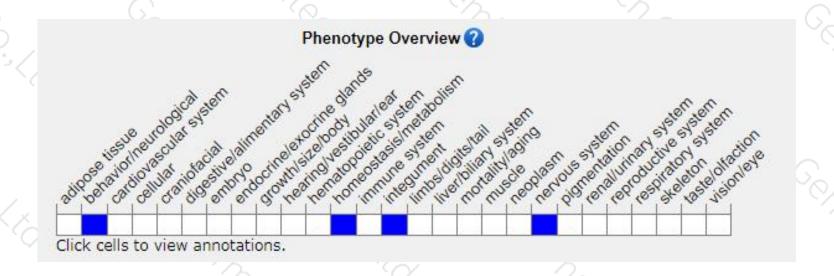
Protein domain





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, Mice homozygous for an insertional or null mutation fail to develop normal patterned distribution of neurons in the brain and display behavioral and learning abnormalities.



If you have any questions, you are welcome to inquire. Tel: 400-9660890





