

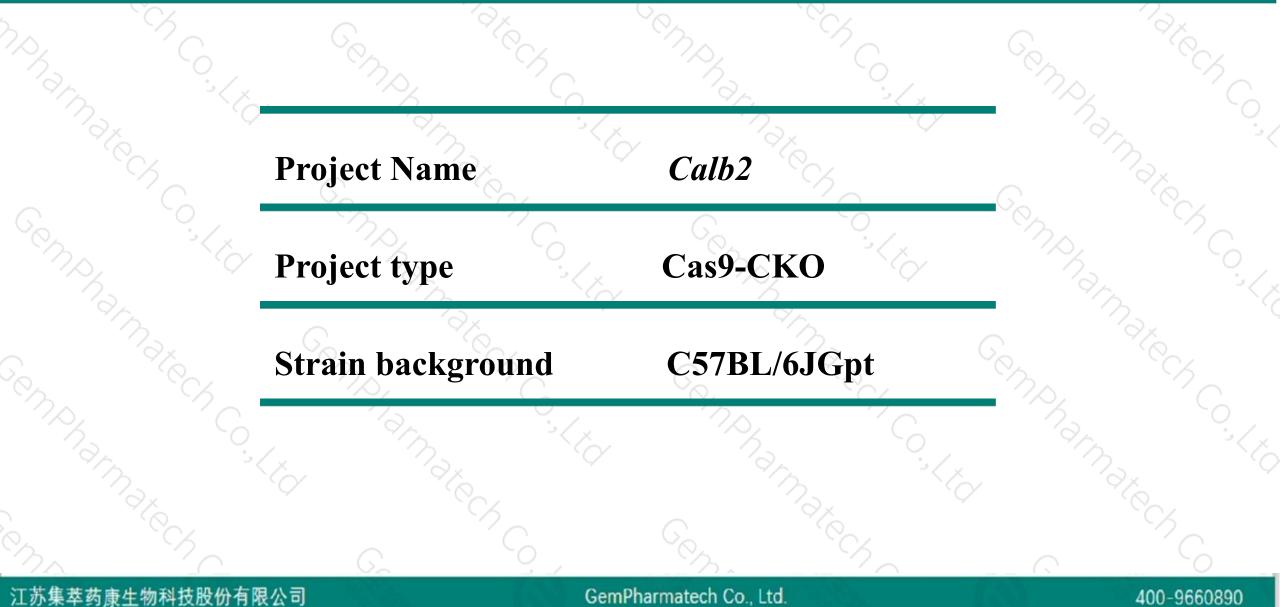
# Cemphamatery of Calb2 Cas9-CKO Strategy Romphamater Contraction

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# **Project Overview**

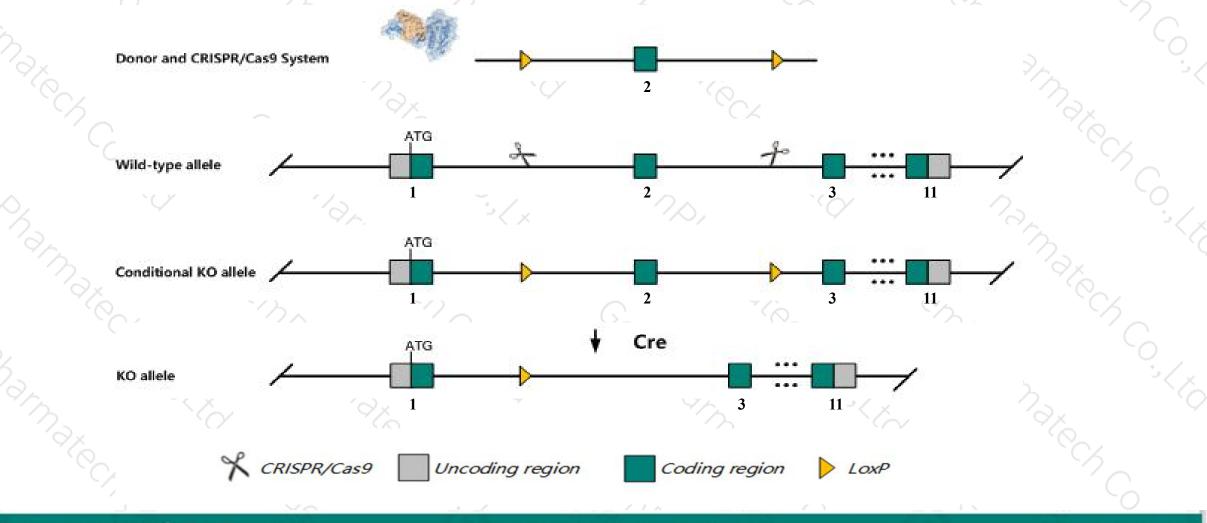




# **Conditional Knockout strategy**



This model will use CRISPR/Cas9 technology to edit the Calb2 gene. The schematic diagram is as follows:



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The Calb2 gene has 2 transcripts. According to the structure of Calb2 gene, exon2 of Calb2-201 (ENSMUST00000003754.7) transcript is recommended as the knockout region. The region contains 77bp coding sequence. Knock out the region will result in disruption of protein function.

In this project we use CRISPR/Cas9 technology to modify *Calb2* 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 targeted mutants showed normal growth, normal brain histology, and generally normal behavior. Impaired motor coordination was observed in wheel running in young and old mutant mice, and progressive impairment was seen on the runway and horizontal stationary rod tests in older mice. Abnormalities are observed in Purkinje cell firing, altering both simple and complex spikes.
- The Calb2 gene is located on the Chr8. 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)



☆ ?

# Calb2 calbindin 2 [Mus musculus (house mouse)]

Gene ID: 12308, updated on 26-Mar-2019

#### Summary

Official SymbolCalb2 provided by MGIOfficial Full Namecalbindin 2 provided by MGIPrimary soureMGI:MGI:101914See relatedEnsembl:ENSMUSG000003657Gene typeprotein codingVal DATEDVal DATEDOrganismMus musculusLineageEukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Golenta; Myomorpha;<br/>Muroidea; Murinae; Mus; MusAlso knownasCRExpressionBiased expression in cerebellum adult (RPKM 131.6), frontal lobe adult (RPKM 90.9) and 3 other tissuesSee more<br/>human all

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# **Transcript information (Ensembl)**



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

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Calb2-201	ENSMUST0000003754.7	1431	<u>271aa</u>	Protein coding	CCDS22661	<u>Q08331</u>	TSL:1 GENCODE basic APPRIS P1
Calb2-202	ENSMUST00000212297.1	1080	<u>242aa</u>	Protein coding	-5	Q8CCS7	TSL:1 GENCODE basic

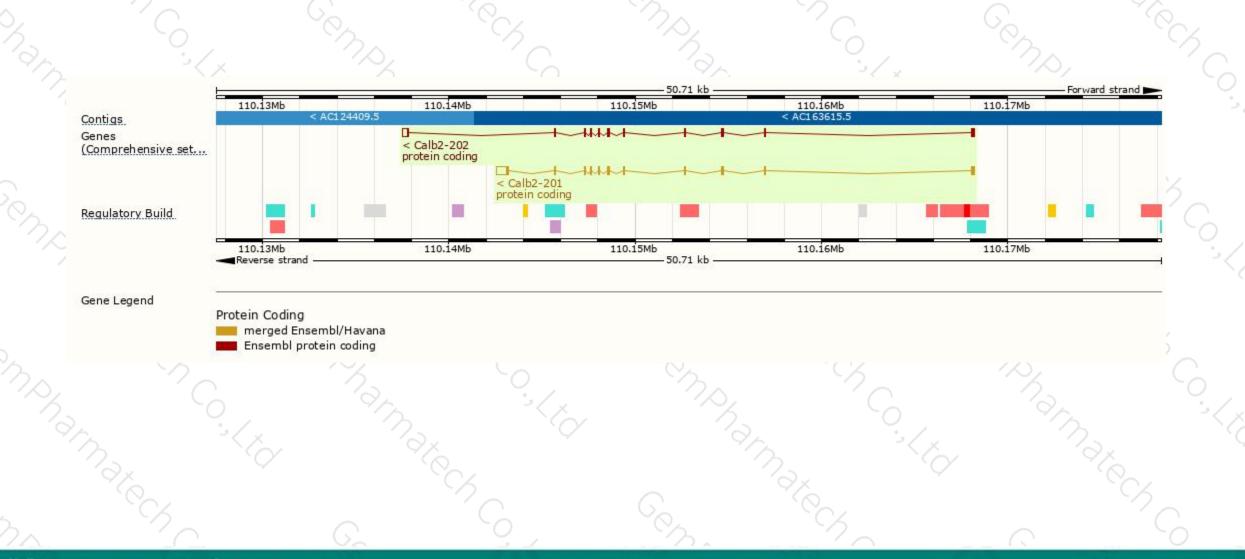
The strategy is based on the design of *Calb2-201* transcript, The transcription is shown below

< Calb2-201 protein coding

Reverse strand 25.67 kb 400-9660890 江苏集萃药康生物科技股份有限公司

# **Genomic location distribution**





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# **Protein domain**



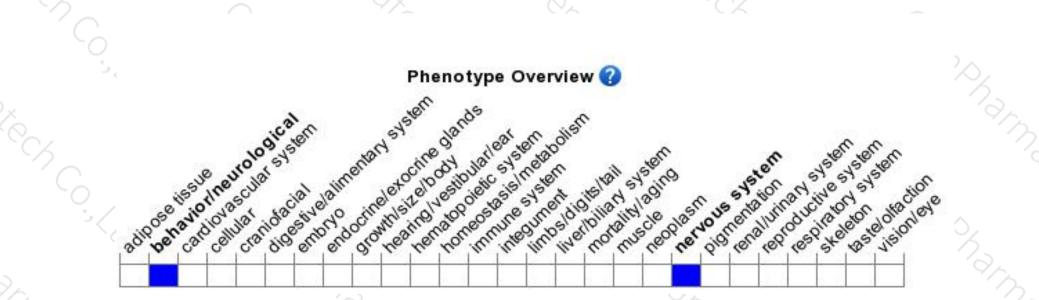
ENSMUSP0000003... Conserved Domains hmmpanther Calretinin PTHR19972 Superfamily domains EF-hand domain pair SMART domains EF-hand domain Pfam domain EF-hand domain EF-hand domain PROSITE profiles EF-hand domain PROSITE patterns EF-Hand 1, calcium-binding site Gene3D 1.10.238.10 Sequence variants (dbSNP and all other sources) All sequence SNPs/i... Variant Legend synonymous variant Scale bar 40 120 160 271 0 200 80

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# 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 targeted mutants showed normal growth, normal brain histology, and generally normal behavior. Impaired motor coordination was observed in wheel running in young and old mutant mice, and progressive impairment was seen on the runway and horizontal stationary rod tests in older mice. Abnormalities are observed in Purkinje cell firing, altering both simple and complex spikes.

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



