

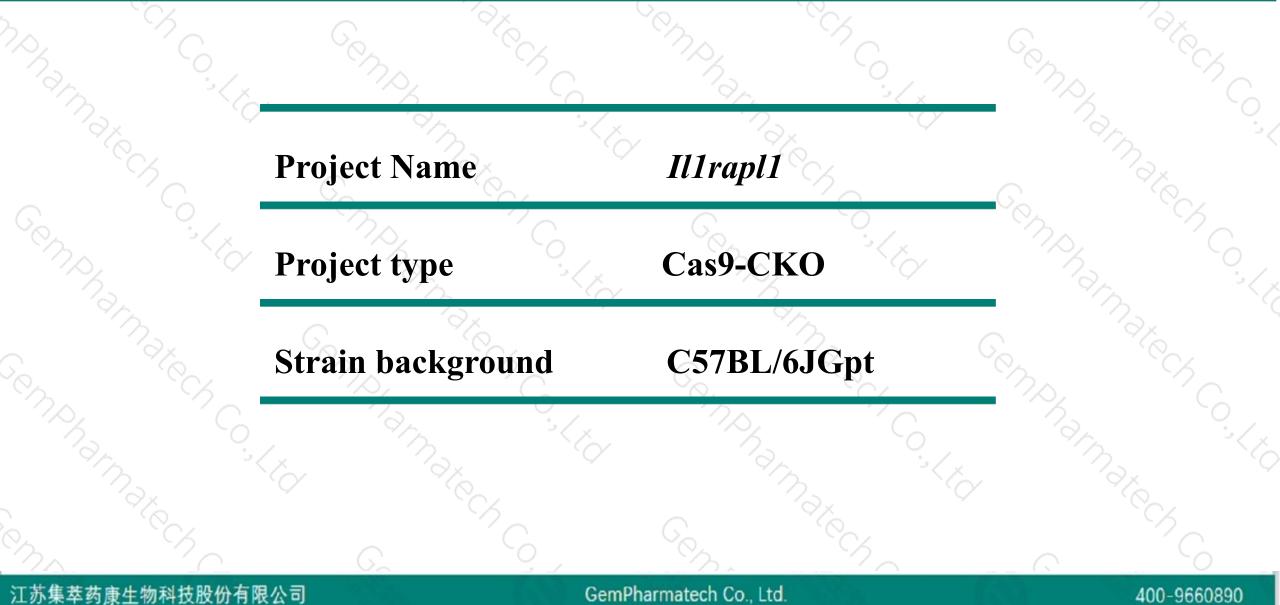
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Project Overview



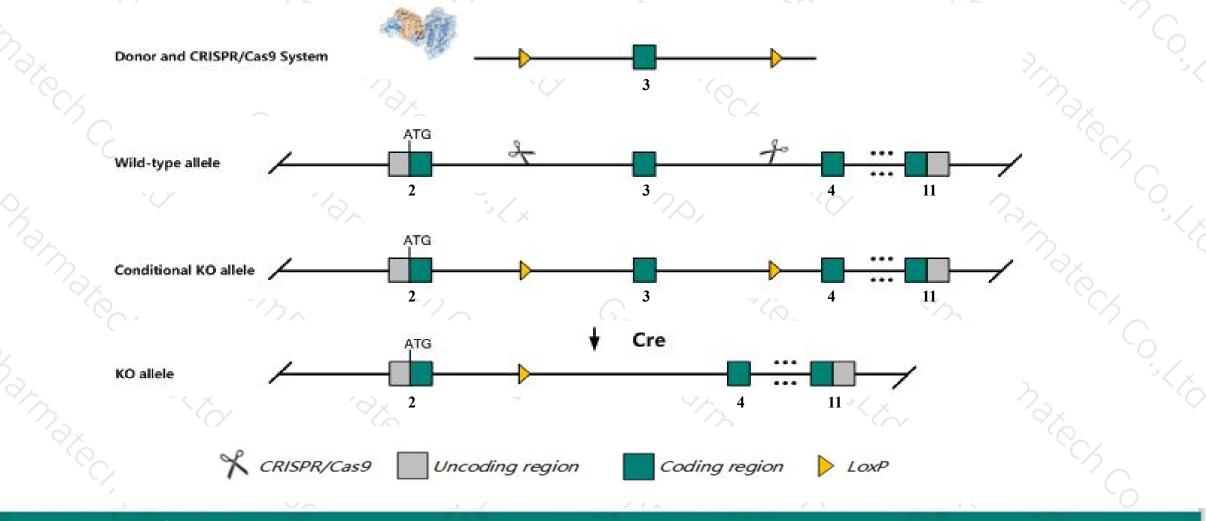


Conditional Knockout strategy



400-9660890

This model will use CRISPR/Cas9 technology to edit the *Illrapl1* gene. The schematic diagram is as follows:



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The *Illrapl1* gene has 3 transcripts. According to the structure of *Illrapl1* gene, exon3 of *Illrapl1-203* (ENSMUST00000113966.7) transcript is recommended as the knockout region. The region contains 280bp coding sequence. Knock out the region will result in disruption of protein function.

In this project we use CRISPR/Cas9 technology to modify *Il1rapl1* 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, Mice homozygous for a knock-out allele exhibit premature giant inhibitory postsynaptic currents and parallel fiber-mediated recruitment of molecular layer interneurons.
- The *Illrapl1* gene is located on the ChrX. 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)



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Il1rapl1 interleukin 1 receptor accessory protein-like 1 [Mus musculus (house mouse)]

Gene ID: 331461, updated on 13-Mar-2019

Summary

Official Symbol	II1rapi1 provided by MGI
Official Full Name	interleukin 1 receptor accessory protein-like 1 provided by MGI
Primary source	MGI:MGI:2687319
See related	Ensembl:ENSMUSG00000052372
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	6330532G10Rik, C030039N24, IL1R8, IL1RAPL, IL1RAPL-1, MRX34, OPHN4, TIGIRR-2
Expression	Biased expression in frontal lobe adult (RPKM 1.3), cortex adult (RPKM 1.1) and 5 other tissues See more
Orthologs	human all

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Transcript information (Ensembl)



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

Name	Transcript ID	bp 💧	Protein	Biotype	CCDS	UniProt	Flags
ll1rapl1-203	ENSMUST00000113966.7	9133	<u>696aa</u>	Protein coding	CCDS53128@	B1ASU0@	TSL:5 GENCODE basic APPRIS P1
ll1rapl1-201	ENSMUST0000078875.7	2091	<u>696aa</u>	Protein coding	CCDS53128@	B1ASU0@	TSL:5 GENCODE basic APPRIS P1
ll1rapl1-202	ENSMUST00000113964.1	1936	<u>383aa</u>	Protein coding	1992	Q0VDP7	TSL:1 GENCODE basic

The strategy is based on the design of *Illrapl1-203* transcript, The transcription is shown below

< Il1rapl1-203 protein coding Reverse strand

-1.37 Mb

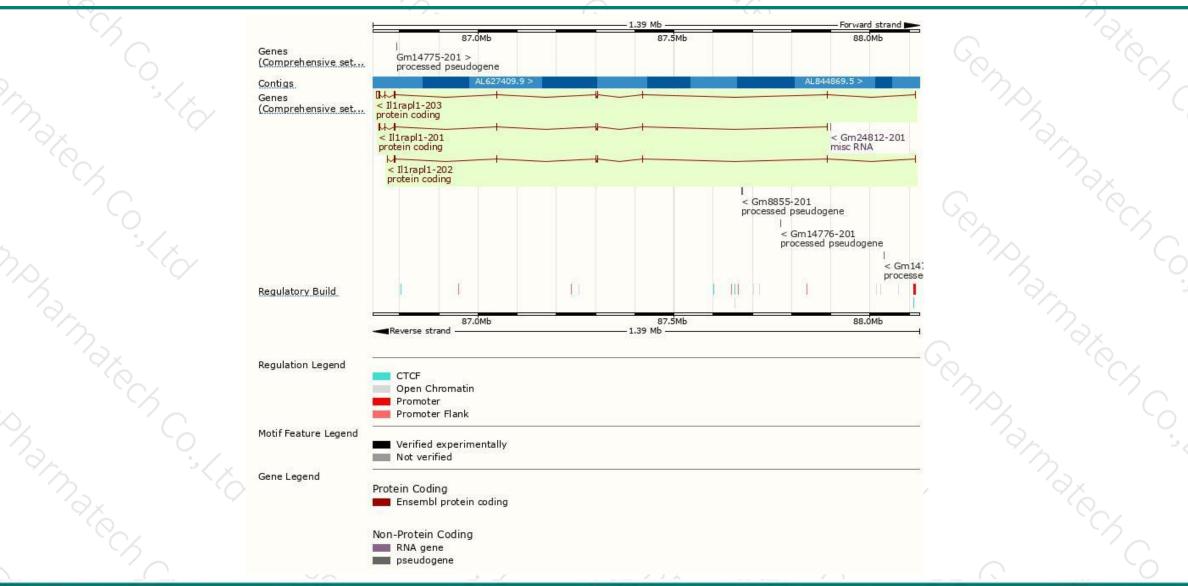
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Genomic location distribution



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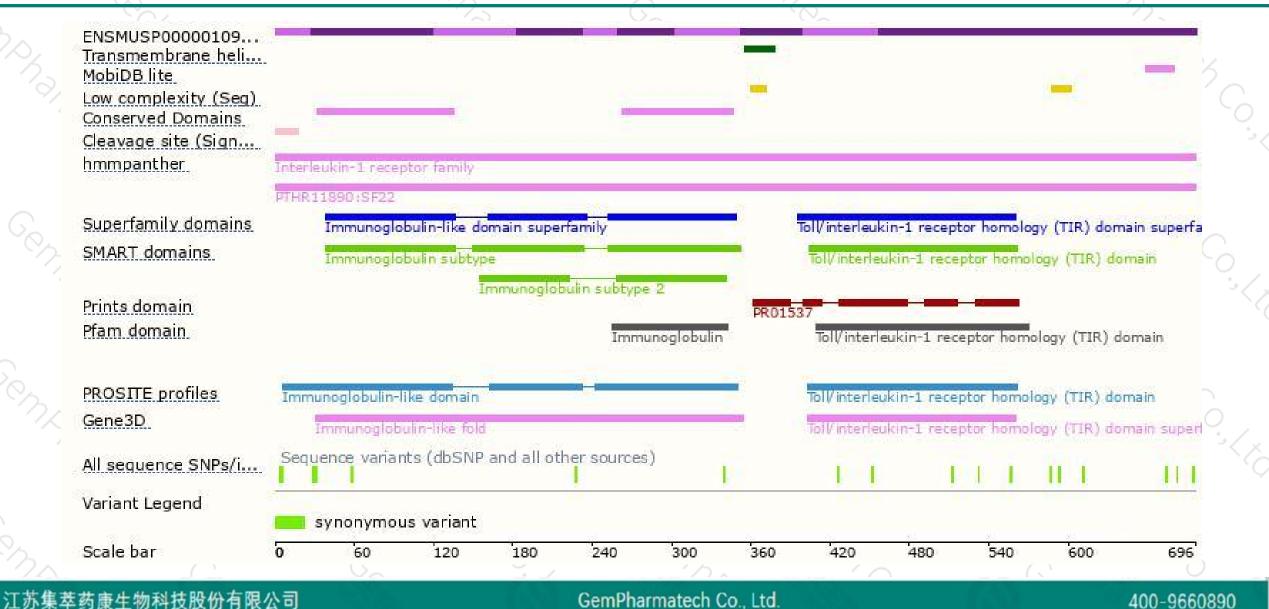


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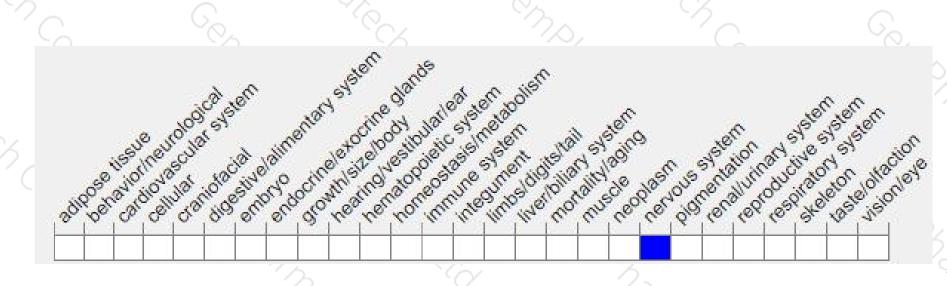
Protein domain





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 a knock-out allele exhibit premature giant inhibitory postsynaptic currents and parallel fiber-mediated recruitment of molecular layer interneurons.



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



