

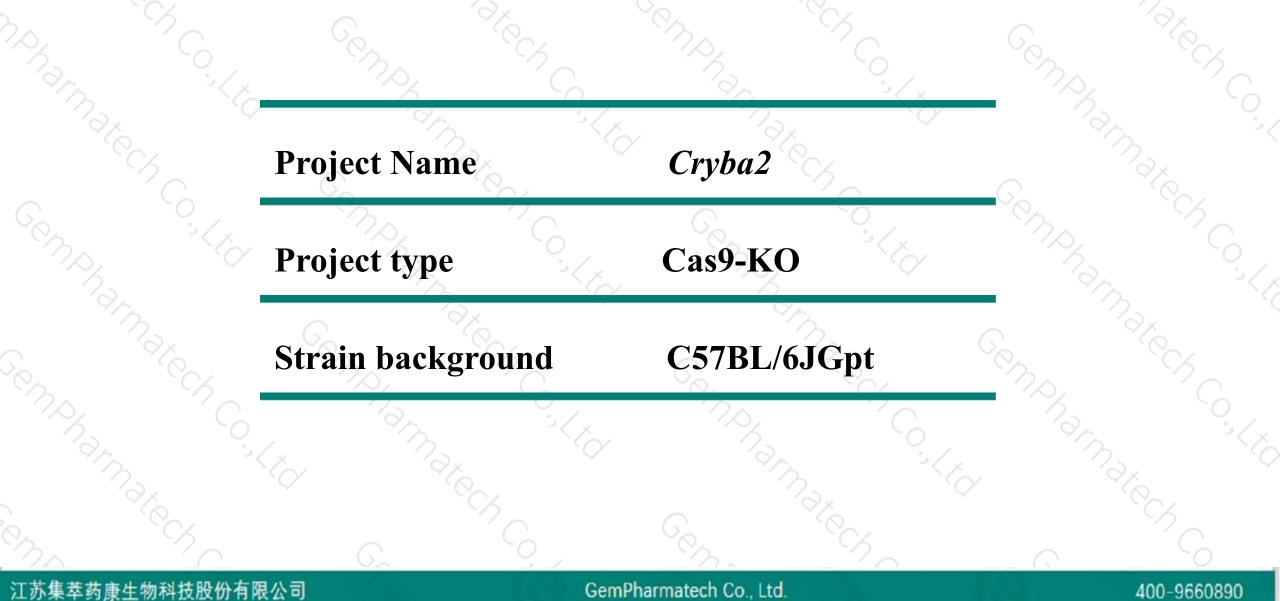
Cryba2 Cas9-KO Strategy

Designer: Reviewer: Design Date:

JiaYu Xiaojing Li 2020-2-19

Project Overview

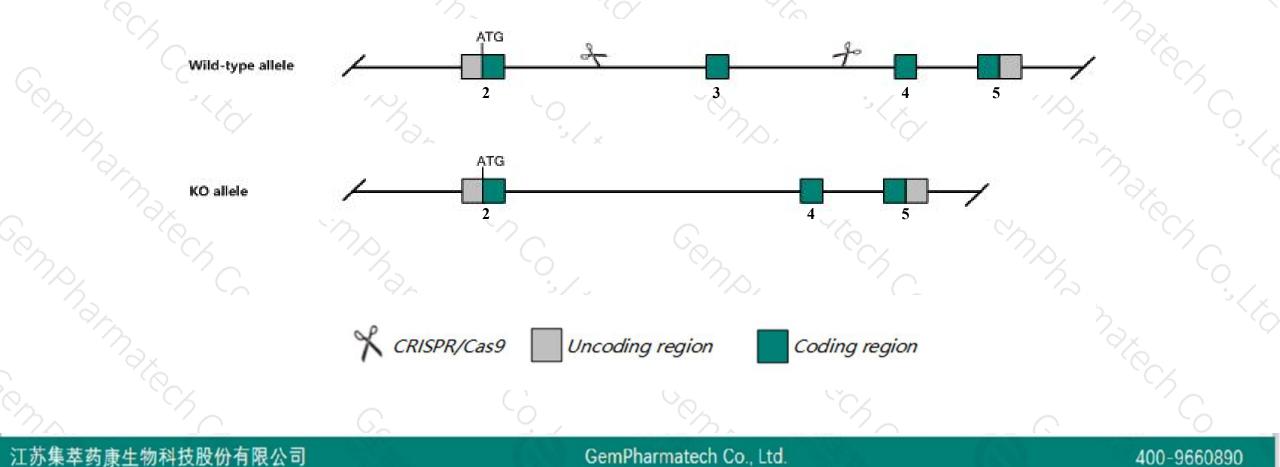




Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Cryba2 gene. The schematic diagram is as follows:





- The Cryba2 gene has 2 transcripts. According to the structure of Cryba2 gene, exon3 of Cryba2-201 (ENSMUST0000006721.2) transcript is recommended as the knockout region. The region contains 142bp coding sequence. Knock out the region will result in disruption of protein function.
- > In this project we use CRISPR/Cas9 technology to modify Cryba2 gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice heterozygous or homozygous for an ENU-induced allele exhibit small lens and cataracts.
- The Cryba2 gene is located on the Chr1. 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 gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Notice

Gene information (NCBI)



☆ ?

Cryba2 crystallin, beta A2 [Mus musculus (house mouse)]

Gene ID: 12958, updated on 31-Jan-2019

Summary

Official SymbolCryba2 provided by MGIOfficial Full Namecrystallin, beta A2 provided byMGIPrimary sourceMGI:MGI:104336Primary sourceEnsembl:ENSMUSG0000006546Gene typeprotein codingRefSeq statusVALIDATEDOrganismMus musculusLineageEukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Golires; Rodentia; Myomorpha;
Muroidea; Murinae; Mus; MusAlso knownasE130107M19RikExpressionLow expression observed in reference datasetSee more
human all

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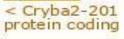
Transcript information (Ensembl)



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

Name Transcript ID		bp	Protein	Biotype	CCDS	UniProt	Flags	
Cryba2-201	ENSMUST0000006721.2	720	<u>197aa</u>	Protein coding	CCDS15059	Q9JJV1	TSL:1 GENCODE basic APPRIS P1	
Cryba2-202	ENSMUST00000133833.2	923	<u>54aa</u>	Nonsense mediated decay		A0A087WQQ2	TSL:3	

The strategy is based on the design of Cryba2-201 transcript, The transcription is shown below



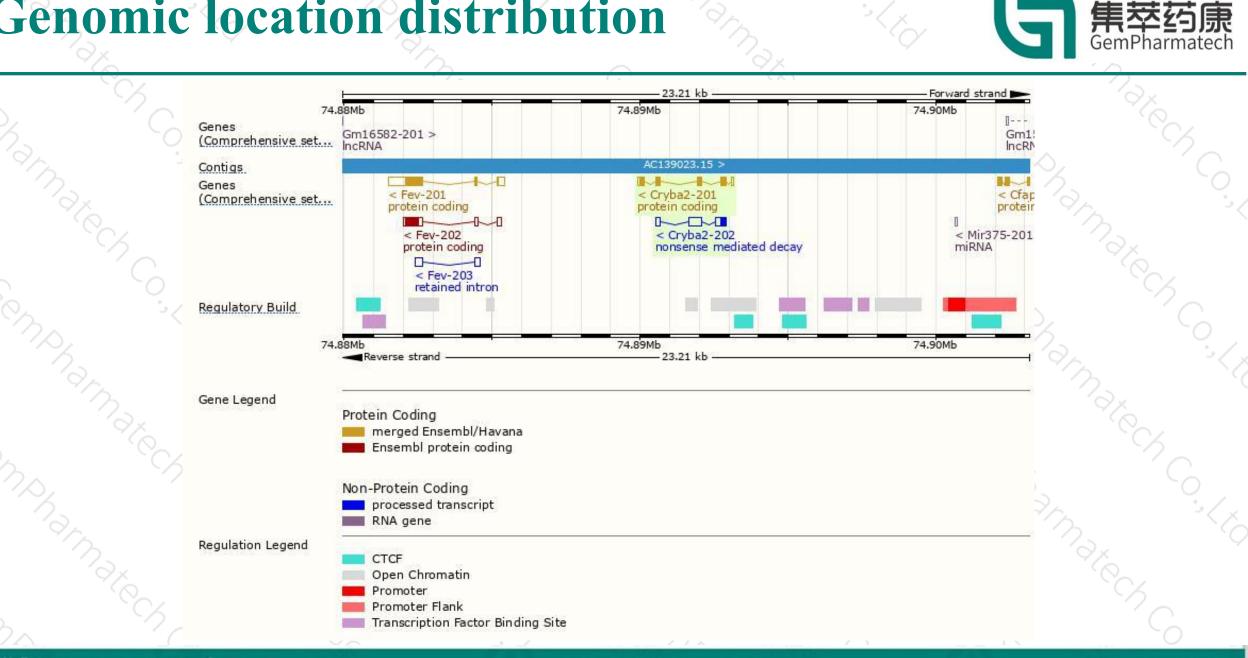
Reverse strand

- 3.21 kb -

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Genomic location distribution



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Protein domain



ENSMUSP0000006... Low complexity (Seg) Superfamily SMART Prints Pfam PROSITE profiles PANTHER

Gene3D

All sequence SNPs/i...

Variant Legend

Scale bar

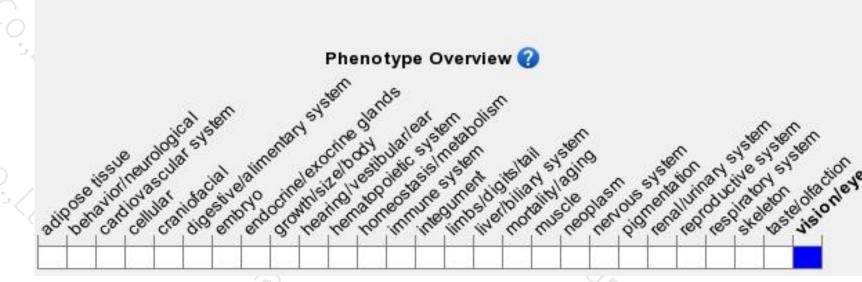
Gamma	-crystallin-like							
Beta/ga	amma crystallin	Beta/gamma	crystallin				-	_
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			80	100	120	140	160	19

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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, Mice heterozygous or homozygous for an ENU-induced allele exhibit small lens and

cataracts.



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



