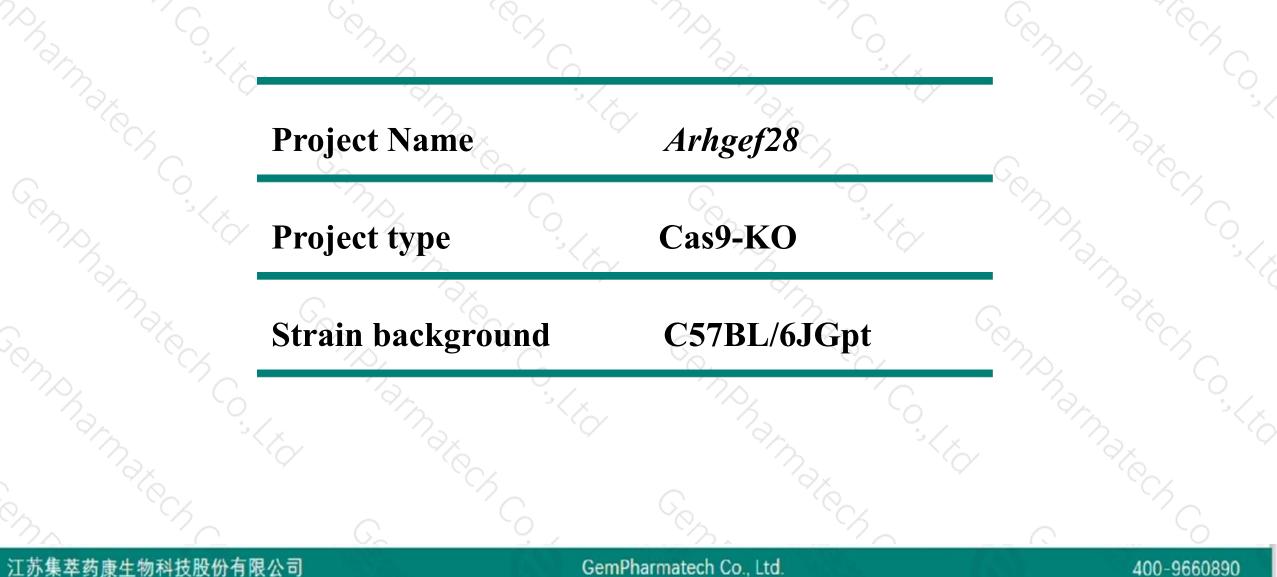


Arhgef28 Cas9-KO Strategy

Designer: Reviewer: Design Date: JiaYu Xiaojing Li 2020-3-17

Project Overview

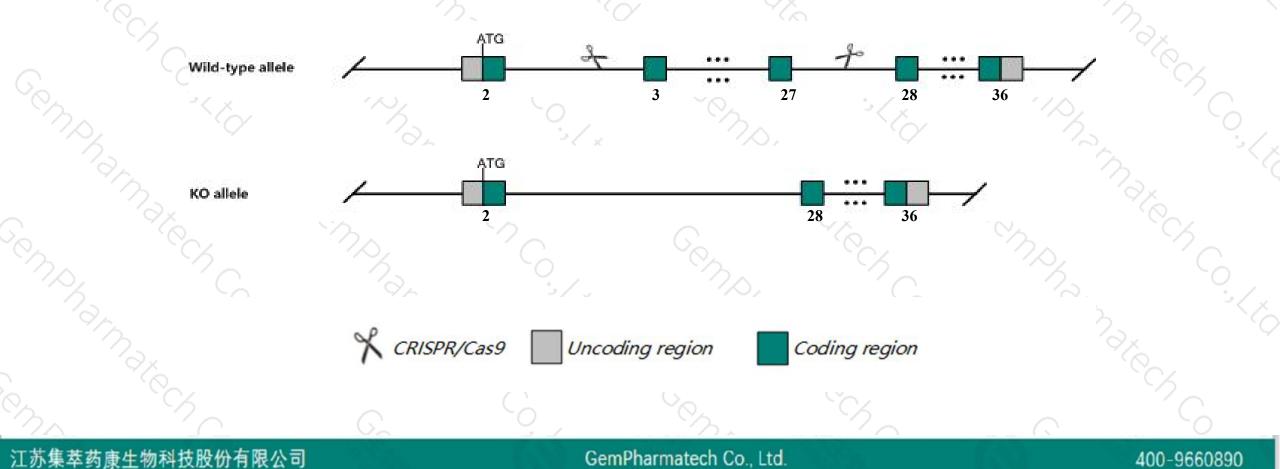




Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Arhgef28 gene. The schematic diagram is as follows:





- The Arhgef28 gene has 7 transcripts. According to the structure of Arhgef28 gene, exon3-exon27 of Arhgef28-201 (ENSMUST00000109426.2) transcript is recommended as the knockout region. The region contains 3524bp coding sequence. Knock out the region will result in disruption of protein function.
- > In this project we use CRISPR/Cas9 technology to modify Arhgef28 gene. The brief process is as follows: CRISPR/Cas9 system



- > According to the existing MGI data, Mice homozygous for a knock-out allele are born at lower than expected Mendelian ratios and exhibit a reduction in overall size that becomes negligible by 8 weeks of age. Mouse embryonic fibroblasts display defects in cell migration and focal adhesion formation.
- ➤ Transcript 202,203 CDS 3' incomplete the influences is unknown.
- The Arhgef28 gene is located on the Chr13.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

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Gene information (NCBI)



☆ ?

Arhgef28 Rho guanine nucleotide exchange factor (GEF) 28 [Mus musculus (house mouse)]

Gene ID: 110596, updated on 13-Mar-2020

Summary

Official SymbolArhgef28 provided by MGIOfficial Full NameRho guanine nucleotide exchange factor (GEF) 28 provided by MGIPrimary sourceMGI:MGI:1346016See relatedEnsembl:ENSMUSG0000021662Gene typeprotein codingRefSeq statusVALIDATEDOrganismMus musculusLineageEukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Myomorpha; Muroidea; Murinae; Mus; MusAlso known asRIP2; Rgnef; RhoGEF; Rhoip2; Al323540; p190RhoGEF; D13Bwg1089e; 9230110L08RikExpressionBroad expression in kidney adult (RPKM 5.2), ovary adult (RPKM 5.2) and 22 other tissues See more
human all

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Transcript information (Ensembl)



The gene has 7 transcripts, all transcripts are shown below:

Name 🖕	Transcript ID 👙	bp 🍦	Protein 🖕	Biotype 🖕	CCDS	UniProt 🍦	Flags
Arhgef28-201	ENSMUST00000109426.2	5387	<u>1700aa</u>	Protein coding	<u>CCDS36758</u> 교	<u>G5E8P2</u>	TSL:1 GENCODE basic APPRIS P2
Arhgef28-207	ENSMUST00000238746.1	4660	<u>1387aa</u>	Protein coding	52	7.1	APPRIS ALT2
Arhgef28-202	ENSMUST00000223849.1	2001	<u>643aa</u>	Protein coding		<u>A0A286YCY0</u> 团	CDS 3' incomplete
Arhgef28-203	ENSMUST00000224866.1	203	<u>11aa</u>	Protein coding		A0A286YD71团	CDS 3' incomplete
Arhgef28-206	ENSMUST00000225884.1	5307	<u>1324aa</u>	Nonsense mediated decay	52	A0A286YDI4 &	
Arhgef28-205	ENSMUST00000225663.1	4977	No protein	Processed transcript	52		
Arhgef28-204	ENSMUST00000225269.1	783	No protein	Retained intron	52		

The strategy is based on the design of Arhgef28-201 transcript, The transcription is shown below

< Arhgef28-201 protein coding

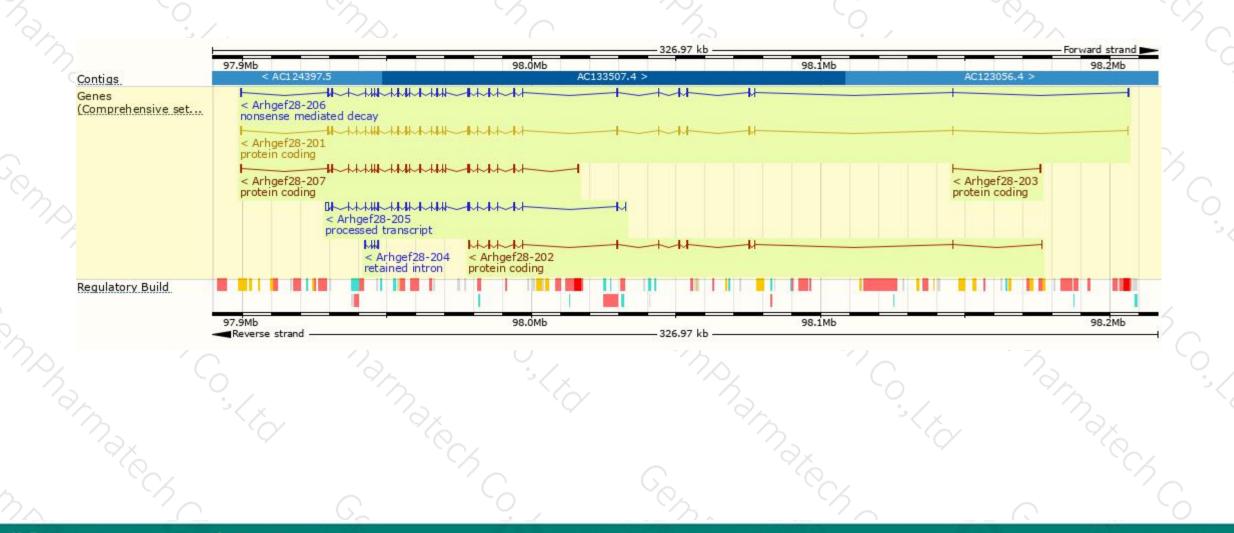
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Genomic location distribution



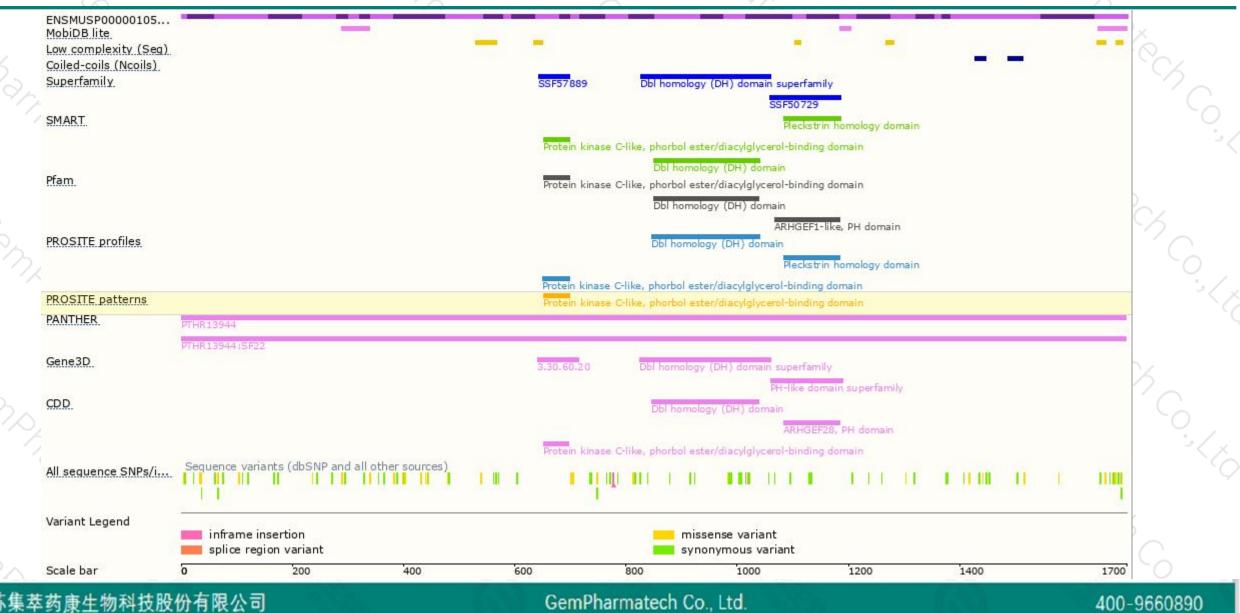
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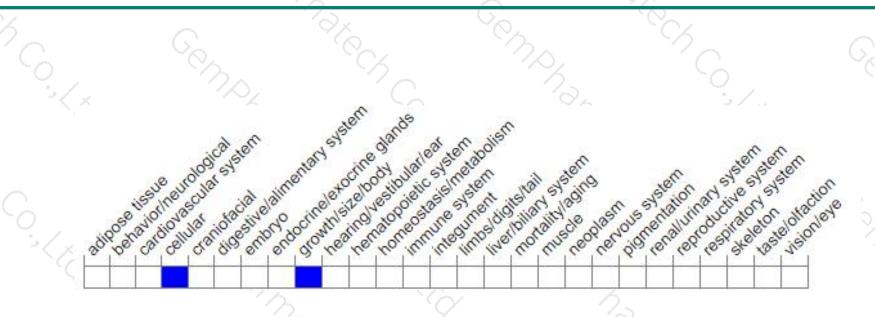
Protein domain





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 are born at lower than expected Mendelian ratios and exhibit a reduction in overall size that becomes negligible by 8 weeks of age. Mouse embryonic fibroblasts display defects in cell migration and focal adhesion formation.

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



