

# Efhc2 Cas9-KO Strategy

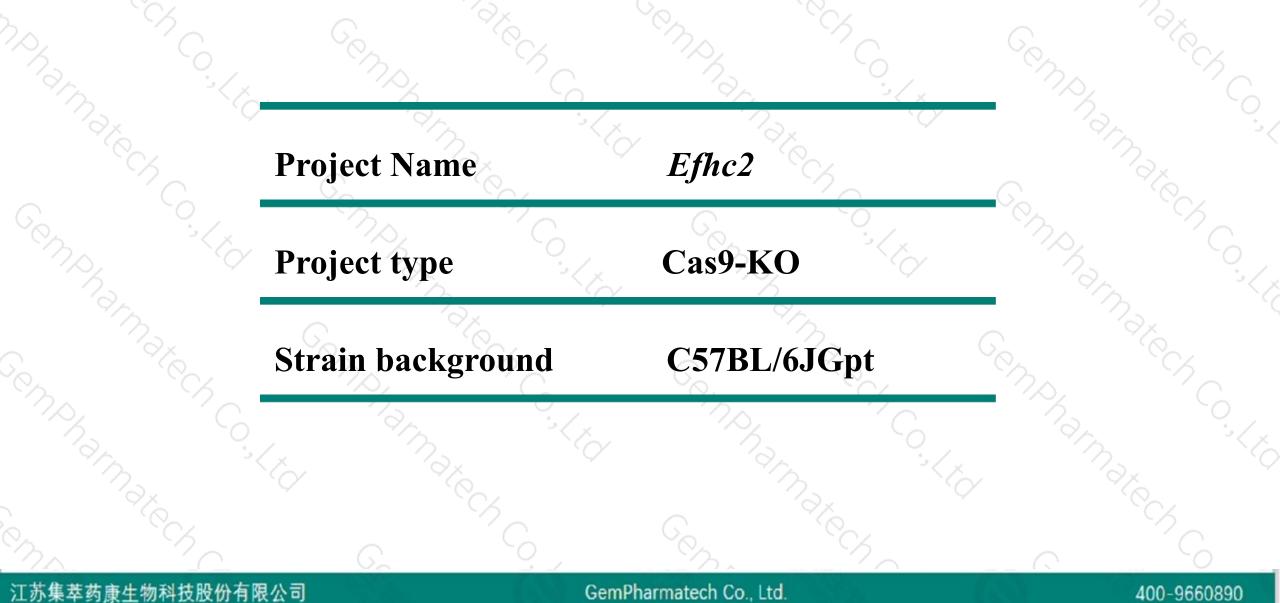
Designer: Reviewer:

**Design Date:** 

Daohua Xu Huimin Su 2020-1-1

# **Project Overview**

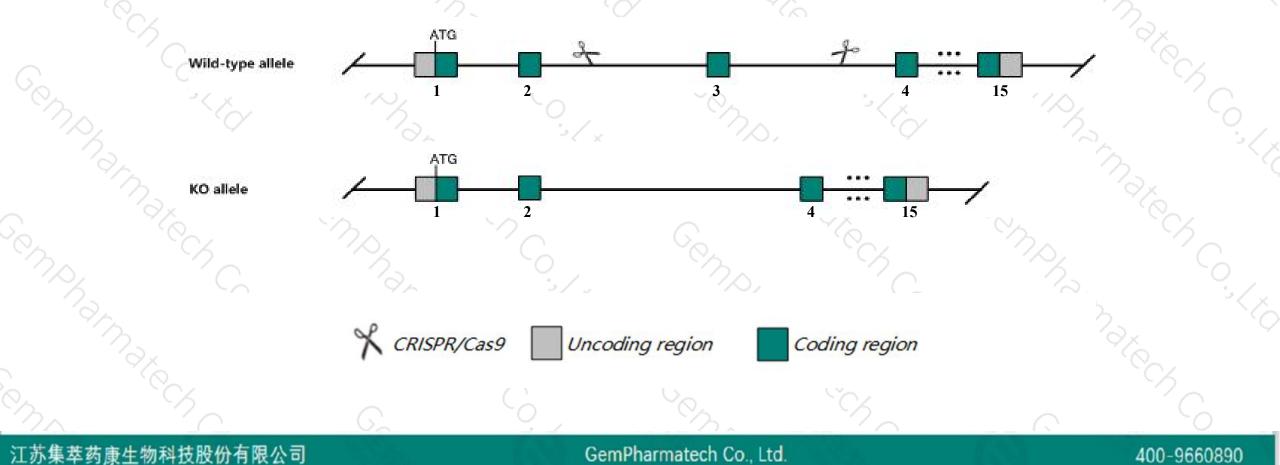




# **Knockout strategy**



This model will use CRISPR/Cas9 technology to edit the *Efhc2* gene. The schematic diagram is as follows:





- The *Efhc2* gene has 1 transcript. According to the structure of *Efhc2* gene, exon3 of *Efhc2-201* (ENSMUST0000026014.7) transcript is recommended as the knockout region. The region contains 151bp coding sequence. Knock out the region will result in disruption of protein function.
- > In this project we use CRISPR/Cas9 technology to modify *Efhc2* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Efhc2* 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Notice

# **Gene information (NCBI)**



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#### Efhc2 EF-hand domain (C-terminal) containing 2 [Mus musculus (house mouse)]

Gene ID: 74405, updated on 31-Jan-2019

#### Summary

Official Symbol	Efhc2 provided by MGI
<b>Official Full Name</b>	EF-hand domain (C-terminal) containing 2 provided by MGI
<b>Primary source</b>	MGI:MGI:1921655
See related	Ensembl:ENSMUSG0000025038
Gene type	protein coding
<b>RefSeq status</b>	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	4933407D04Rik, mRib72-2
Expression	Biased expression in testis adult (RPKM 2.7), placenta adult (RPKM 0.8) and 6 other tissuesSee more
Orthologs	human all

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# **Transcript information (Ensembl)**



The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Efhc2-201	ENSMUST0000026014.7	2554	<u>750aa</u>	Protein coding	CCDS40883	Q059K2 Q9D485	TSL:1 GENCODE basic APPRIS P1
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The strateg	y is based on the design	of Efh	<i>c2-201</i> tr	anscript, The t	ranscription	is shown below	$\gamma_{\rm e} = 0$
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Efhc2-201			MM				
rotein codin	9						

Reverse strand

- 187.32 kb -

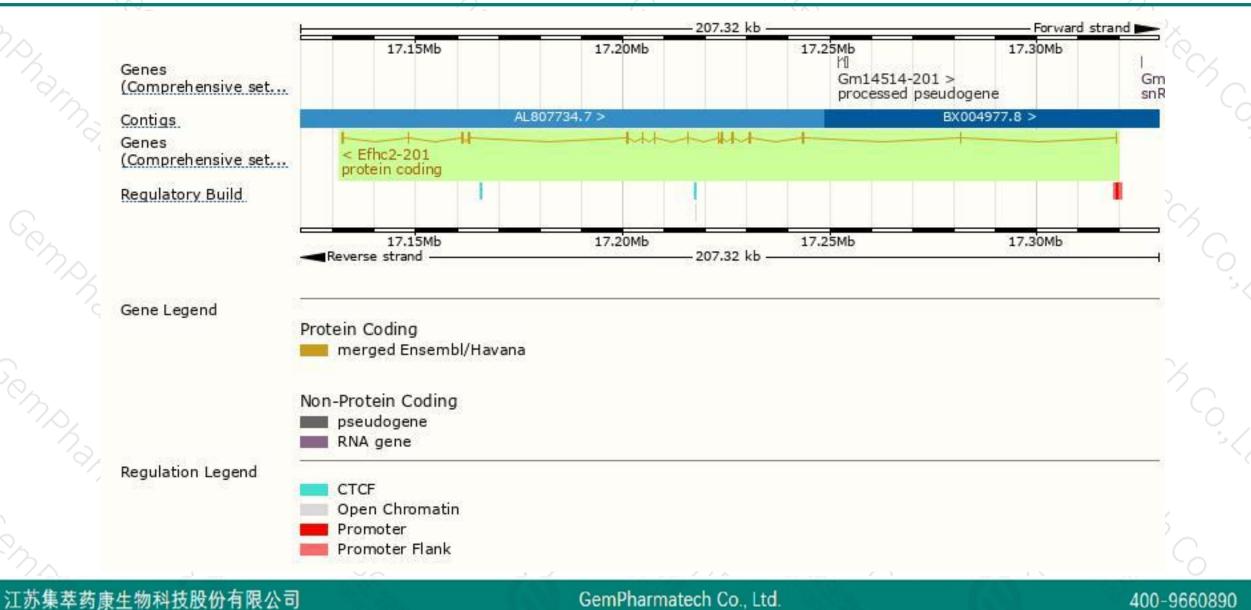
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### **Genomic location distribution**





### **Protein domain**



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



