

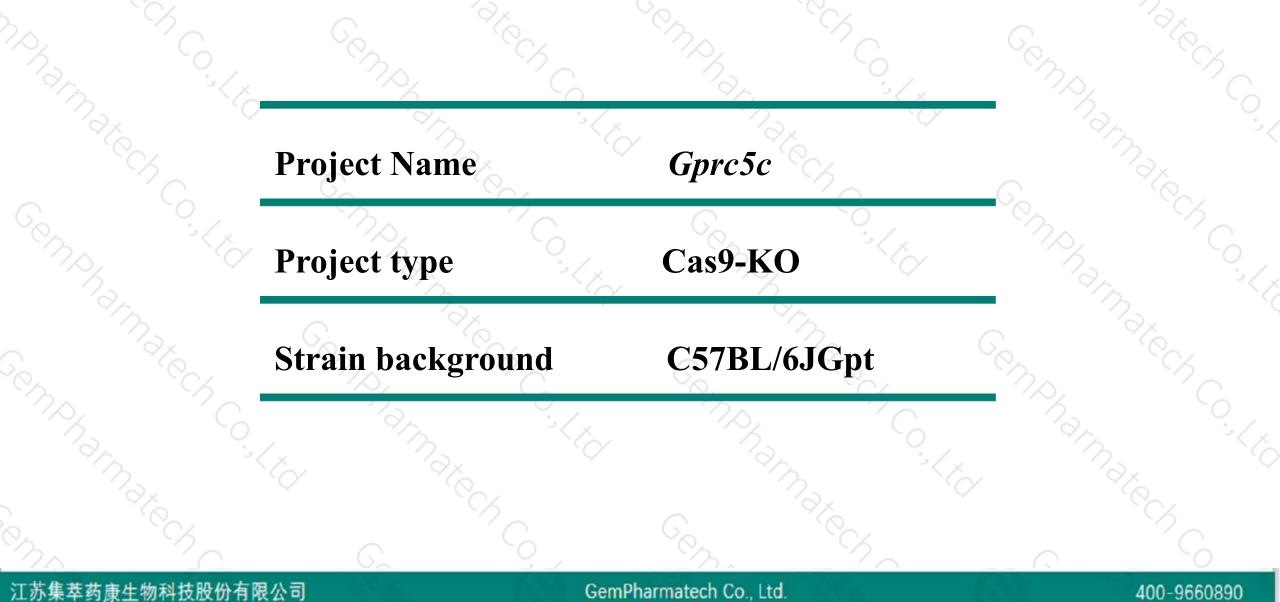
# **Gprc5c** Cas9-KO Strategy

Designer: Xueting Zhang Design Date: 2019-8-5

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### **Project Overview**

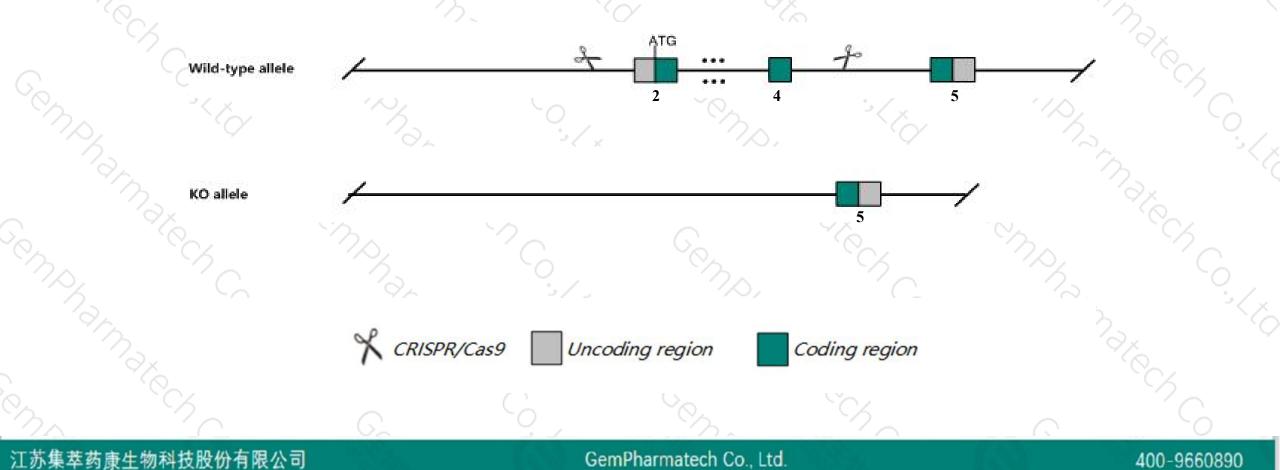




# **Knockout** strategy



This model will use CRISPR/Cas9 technology to edit the Gprc5c gene. The schematic diagram is as follows:





The Gprc5c gene has 8 transcripts. According to the structure of Gprc5c gene, exon2-exon4 of Gprc5c-202 (ENSMUST00000053361.11) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.

> In this project we use CRISPR/Cas9 technology to modify Gprc5c gene. The brief process is as follows: CRISPR/Cas9 system



- According to the existing MGI data, Mice homozygous for a reporter allele are behaviorally normal but exhibit reticulocytosis, increased mean corpuscular volume, increased percentage of basophils, decreased mean corpuscular hemogloblin concentration, and increased alkaline phophatase and lactic dehydrogenase levels.
- The Gprc5c gene is located on the Chr11. 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.

# **Gene information (NCBI)**



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### Gprc5c G protein-coupled receptor, family C, group 5, member C [Mus musculus (house mouse)]

Gene ID: 70355, updated on 31-Jan-2019

#### Summary

Official Symbol	Gprc5c provided by MGI
Official Full Name	G protein-coupled receptor, family C, group 5, member C provided by MGI
<b>Primary source</b>	MGI:MGI:1917605
See related	Ensembl:ENSMUSG00000051043
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	1110028I06Rik, 3200002M13Rik, Raig3
Expression	Biased expression in kidney adult (RPKM 60.6), genital fat pad adult (RPKM 25.6) and 9 other tissues See more
Orthologs	human all

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## **Transcript information (Ensembl)**



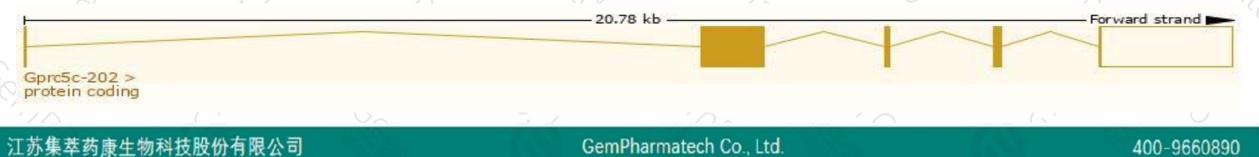
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### The gene has 8 transcripts, all transcripts are shown below:

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Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags		
Gprc5c-202	ENSMUST00000053361.11	3667	<u>441aa</u>	Protein coding	CCDS48978	I7HPW4	TSL:1 GENCODE basic APPRIS ALT2		
Gprc5c-201	ENSMUST00000021071.13	2200	<u>440aa</u>	Protein coding	CCDS25612	<u>J3JS84</u>	TSL:1 GENCODE basic APPRIS P3		
Gprc5c-208	ENSMUST00000177952.7	1809	<u>440aa</u>	Protein coding	CCDS25612	<u>J3JS84</u>	TSL:1 GENCODE basic APPRIS P3		
Gprc5c-203	ENSMUST00000122967.2	1709	<u>104aa</u>	Protein coding	20	E9PZP1	TSL:5 GENCODE basic		
Gprc5c-205	ENSMUST00000136785.1	988	<u>283aa</u>	Protein coding	₹4	B7FAT8	CDS 3' incomplete TSL:3		
Gprc5c-207	ENSMUST00000152314.1	645	<u>109aa</u>	Protein coding	-	B7FAT9	CDS 5' incomplete TSL:5		
Gprc5c-206	06 ENSMUST00000142262.1 481		<u>133aa</u>	Protein coding	10	B7FAT7	CDS 3' incomplete TSL:5		
Gprc5c-204	ENSMUST00000133245.1	359	<u>78aa</u>	Protein coding	17 <u>1</u> 3	B7FAU0	CDS 3' incomplete TSL:3		

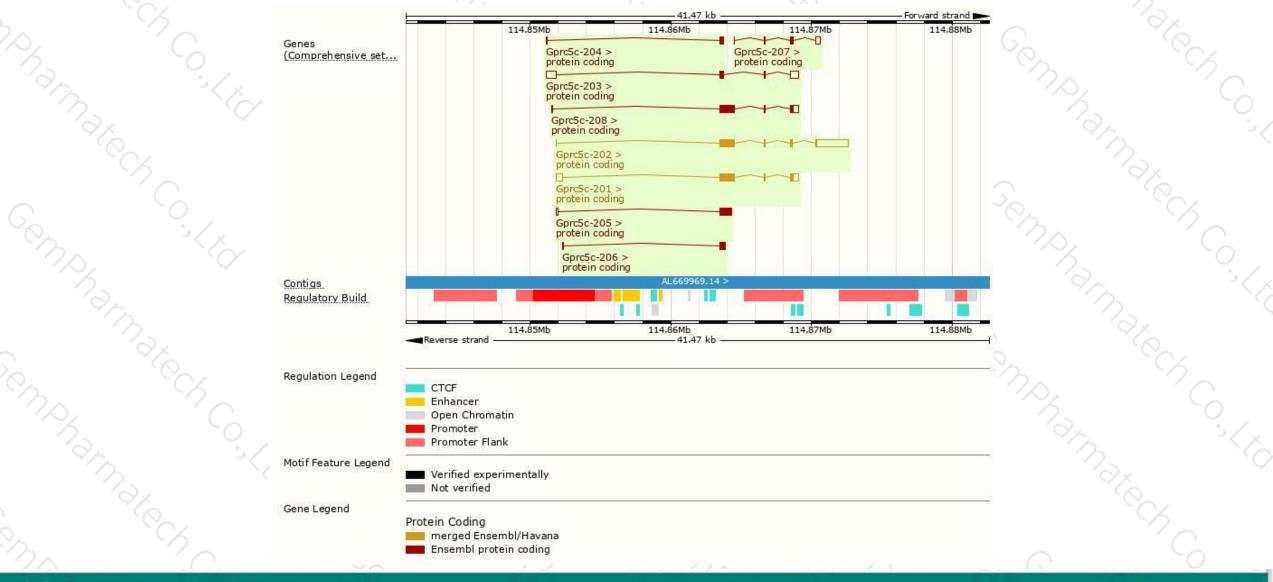
The strategy is based on the design of Gprc5c-202 transcript, The transcription is shown below



### **Genomic location distribution**



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### **Protein domain**

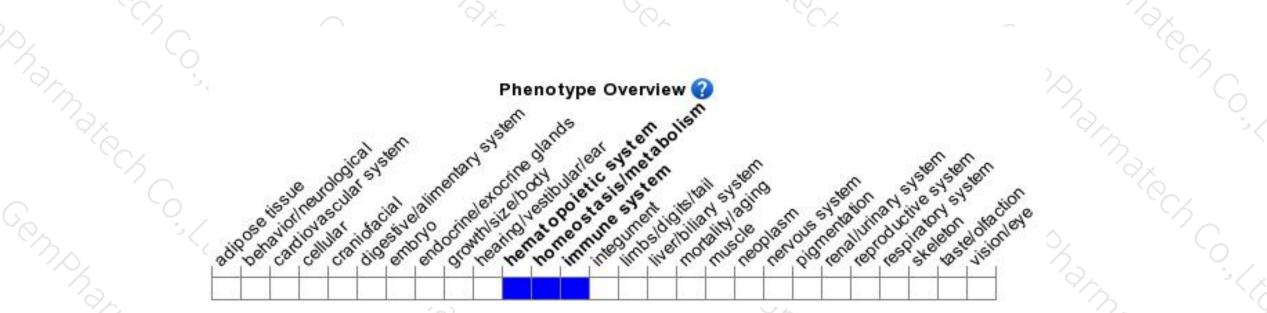
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### Mouse phenotype description(MGI)





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 homozygous for a reporter allele are behaviorally normal but exhibit reticulocytosis, increased mean corpuscular volume, increased percentage of basophils, decreased mean corpuscular hemoglo concentration, and increased alkaline phophatase and lactic dehydrogenase levels.

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



