

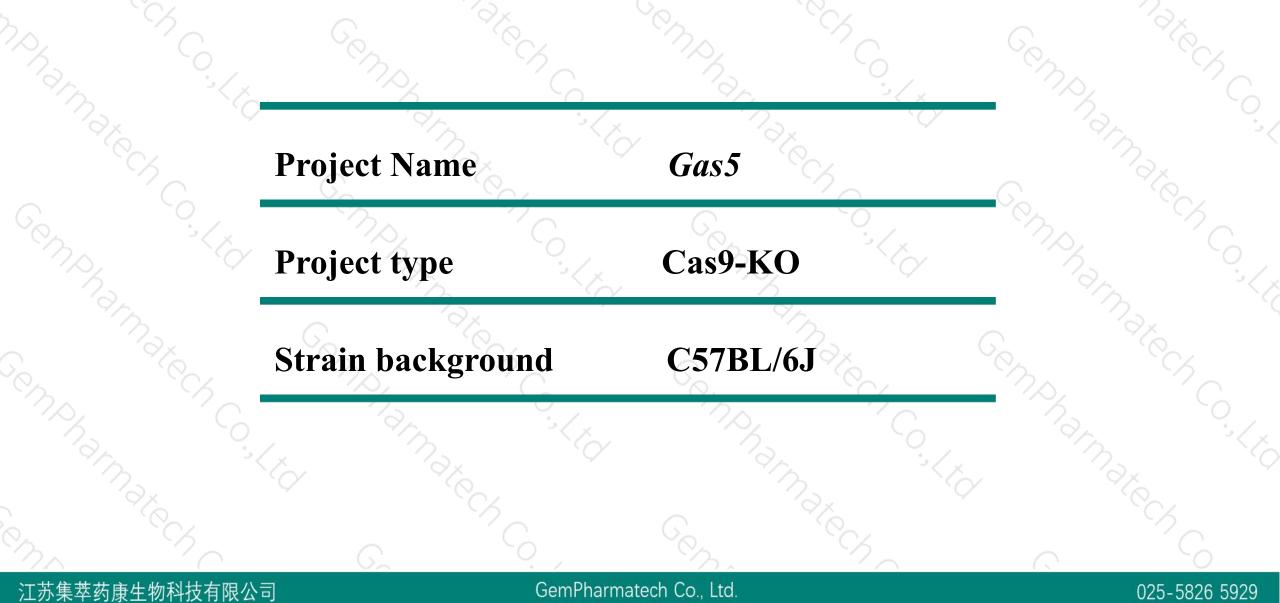
Gas5 Cas9-KO Strategy Andraker Costy

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Project Overview

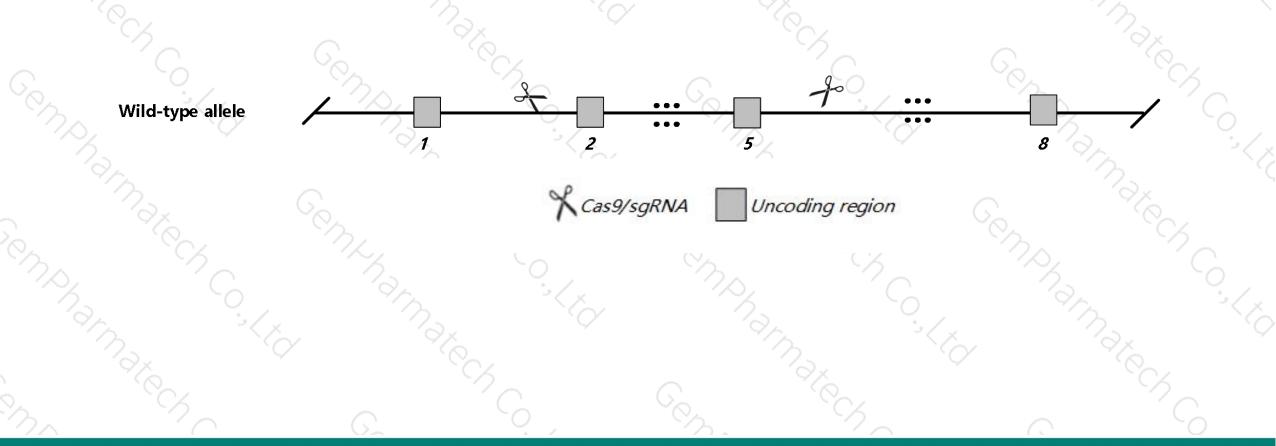






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This model will use CRISPR/Cas9 technology to edit the Gas5 gene. The schematic diagram is as follows:





The Gas5 gene has 25 transcripts. According to the structure of Gas5 gene, exon2-exon5 of Gas5-217
(ENSMUST00000161005.7) transcript is recommended as the knockout region. The region contains most sequence.
Knock out the region will result in disruption of gene function.

In this project we use CRISPR/Cas9 technology to modify Gas5 gene. The brief process is as follows: sgRNA was transcribed in vitro.Cas9 and sgRNA were microinjected into the fertilized eggs of C57BL/6J 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/6J mice.

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➤ The KO region contains functional region of the Snord80, Gm25789, Snord47, Gm26224, Gm22489, Gm22357, Gm23212, Snord78, Mir5117, Zbtb37 gene. Knockout the region may affect the function of Snord80, Gm25789, Snord47, Gm26224, Gm22489, Gm22357, Gm23212, Snord78, Mir5117, Zbtb37 gene.

The Gas5 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 gene transcription and translation processes, all risks cannot be predicted under existing information.

Notice

Gene information (NCBI)



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Gas5 growth arrest specific 5 [Mus musculus (house mouse)]

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Gene ID: 14455, updated on 31-Jan-2019

Summary

Official SymbolGas5 provided by MGIOfficial Full Namegrowth arrest specific 5 provided by MGIPrimary sourceMGI:MGI:95659See relatedEnsembl:ENSMUSG0000053332Gene typencRNARefSeq statusVALIDATEDOrganismMus musculusLineageEukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;
Muroidea; Muriae; Musinae; MusAlso known asGas-5; Snhg2; Mir5117ExpressionBroad expression in liver E14 (RPKM 95.8), CNS E11.5 (RPKM 88.1) and 16 other tissues See more
human all

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Transcript information (Ensembl)

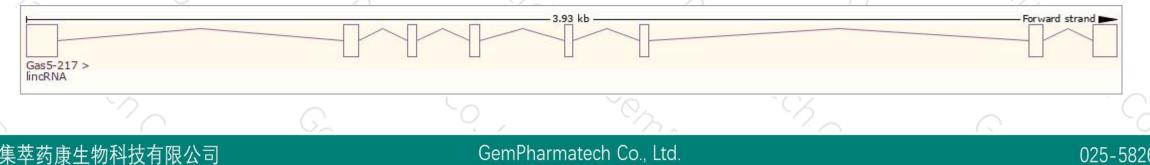


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ne 🝦	Transcript ID 💧	bp 🖕	Protein 🖕	Biotype 🖕	CCDS 🖕	Flags 🍦	\$
as5-218	ENSMUST00000161229.7	2556	No protein	Retained intron		TSL:1	
as5-222	ENSMUST00000162163.7	1614	No protein	Retained intron	(*)	TSL:1	
as5-211	ENSMUST00000159890.7	1066	No protein	Retained intron	1	TSL:5	
as5-223	ENSMUST00000162289.7	904	No protein	Retained intron	-	TSL:5	
as5-204	ENSMUST00000159153.7	872	No protein	Retained intron	- 22	TSL:5	
as5-220	ENSMUST00000161461.7	793	No protein	Retained intron	220	TSL:1	
as5-208	ENSMUST00000159438.7	779	No protein	Retained intron	1.50	TSL:5	
as5-213	ENSMUST00000160429.7	759	No protein	Retained intron		TSL:3	
as5-205	ENSMUST00000159157.7	723	No protein	Retained intron		TSL:3	
as5-212	ENSMUST00000160152.7	712	No protein	Retained intron		TSL:3	
as5-202	ENSMUST00000159037.1	636	No protein	Retained intron	-	TSL:2	
as5-214	ENSMUST00000160497.1	629	No protein	Retained intron	-	TSL:2	
as5-219	ENSMUST00000161380.7	600	No protein	Retained intron		TSL:5	
as5-215	ENSMUST00000160516.1	572	No protein	Retained intron	76 2 8	TSL:2	
as5-216	ENSMUST00000160551.1	549	No protein	Retained intron	1.5	TSL:3	
as5-207	ENSMUST00000159404.7	539	No protein	Retained intron	-	TSL:5	
as5-201 🗄	NSMUST0000065709.11	530	No protein	Retained intron		TSL:2	1
as5-206	ENSMUST00000159399.7	422	No protein	Retained intron	(19 1	TSL:2	
as5-221	ENSMUST00000161623.7	403	No protein	Retained intron	19 4 9	TSL:2	
as5-225	ENSMUST00000163081.1	370	No protein	Retained intron	-	TSL:2	
as5-203	ENSMUST00000159119.7	665	No protein	lincRNA		TSL:5 GENCODE basic	C
as5-210	ENSMUST00000159706.7	557	No protein	IincRNA	1929	TSL:2 GENCODE basic	C
as5-224	ENSMUST00000162558.7	502	No protein	IincRNA	9579	TSL:1 GENCODE basic	C
as5-217 E	ENSMUST00000161005.7	430	No protein	lincRNA	-	TSL:5 GENCODE basic	C
as5-209	ENSMUST00000159663.7	346	No protein	lincRNA		TSL:3 GENCODE basic	C

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The strategy is based on the design of *Gas5-217* transcript, The transcription is shown below



Genomic location distribution





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If you have any questions, you are welcome to inquire. Tel: 025-5864 1534



