

# Gas7 Cas9-KO Strategy

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**Reviewer:** 

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## **Project Overview**



**Project Name** 

Gas7

**Project type** 

Cas9-KO

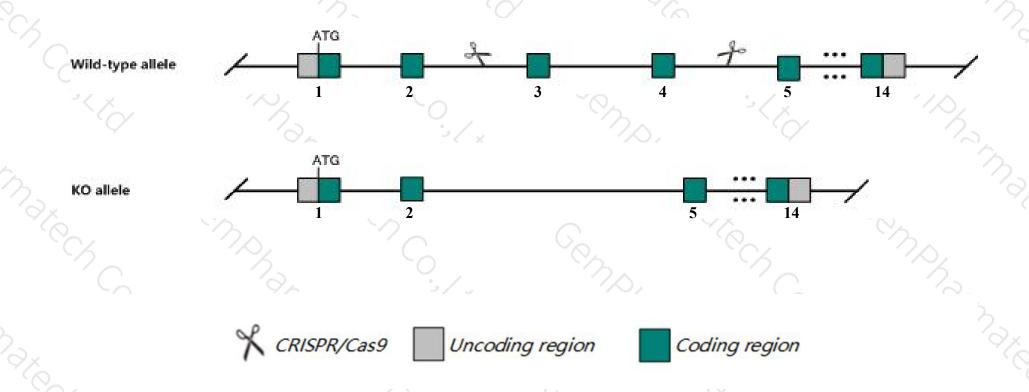
Strain background

C57BL/6JGpt

## **Knockout strategy**



This model will use CRISPR/Cas9 technology to edit the Gas7 gene. The schematic diagram is as follows:



### **Technical routes**



- ➤ The Gas7 gene has 11 transcripts. According to the structure of Gas7 gene, exon3-exon4 of Gas7-204

  (ENSMUST00000108682.8) transcript is recommended as the knockout region. The region contains 167bp coding sequence.

  Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify Gas7 gene. The brief process is as follows: CRISPR/Cas9 system

### **Notice**



- ➤ According to the existing MGI data, Aged mice homozygous for a hypomorphic allele show impaired coordination, decreased grip strength, decreased motor neuron number, muscle weakness, abnormal skeletal muscle fiber type ratio in the soleus muscle, and defects in motor neuron axon terminal sprouting.
- > The Gas7 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)



#### Gas7 growth arrest specific 7 [Mus musculus (house mouse)]

Gene ID: 14457, updated on 31-Jan-2019

#### Summary

☆ ?

Official Symbol Gas7 provided by MGI

Official Full Name growth arrest specific 7 provided by MGI

Primary source MGI:MGI:1202388

See related Ensembl:ENSMUSG00000033066

Gene type protein coding
RefSeq status 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 AW124766, B230343A10Rik, Gas7-cb

Expression Broad expression in cortex adult (RPKM 37.9), cerebellum adult (RPKM 29.9) and 15 other tissuesSee more

Orthologs human all

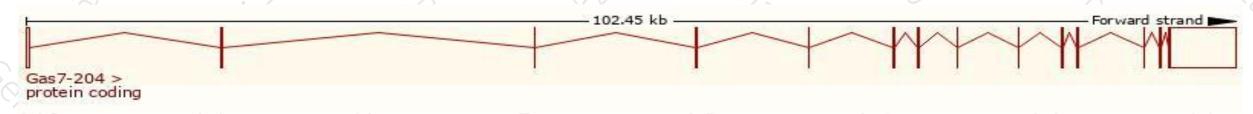
## Transcript information (Ensembl)



The gene has 11 transcripts, all transcripts are shown below:

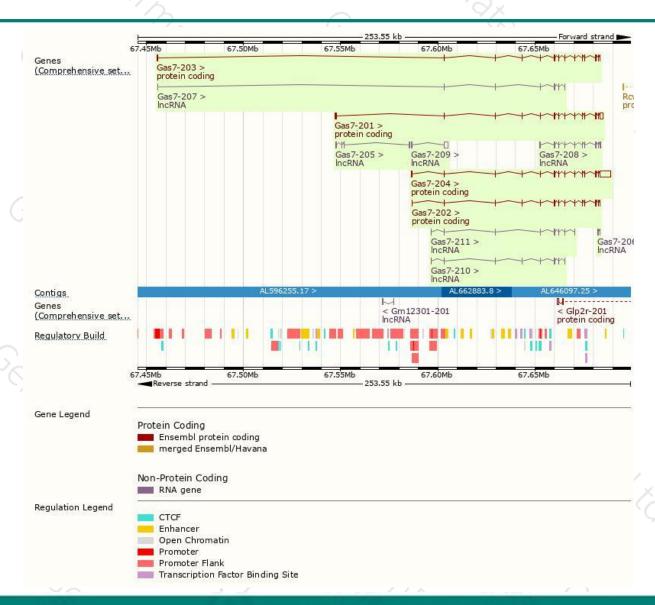
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags	
Gas7-204	ENSMUST00000108682.8	7067	<u>416aa</u>	Protein coding	CCDS48821	Q3U432	TSL:1 GENCODE basic APPRIS P3	
Gas7-201	ENSMUST00000041611.11	2968	<u>412aa</u>	Protein coding	CCDS70210	B1ATI9	TSL:1 GENCODE basic APPRIS ALT1	
Gas7-203	ENSMUST00000108681.7	1595	412aa	Protein coding	CCDS70210	B1ATI9	TSL:5 GENCODE basic APPRIS ALT1	
Gas7-202	ENSMUST00000108680.1	1466	412aa	Protein coding	CCDS70210	B1ATI9	TSL:1 GENCODE basic APPRIS ALT1	
Gas7-209	ENSMUST00000145496.1	2628	No protein	IncRNA		-	TSL:2	
Gas7-208	ENSMUST00000140863.1	1479	No protein	IncRNA	<del>,</del>	-	TSL:2	
Gas7-206	ENSMUST00000133367.1	789	No protein	IncRNA	20	-	TSL:2	
Gas7-205	ENSMUST00000129801.1	785	No protein	IncRNA	20	10	TSL:3	
Gas7-211	ENSMUST00000151784.7	750	No protein	IncRNA	- Tá	-	TSL:5	
Gas7-210	ENSMUST00000147276.7	721	No protein	IncRNA		*	TSL:3	
Gas7-207	ENSMUST00000136796.7	575	No protein	IncRNA	20	-	TSL:3	
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The strategy is based on the design of Gas7-204 transcript, The transcription is shown below



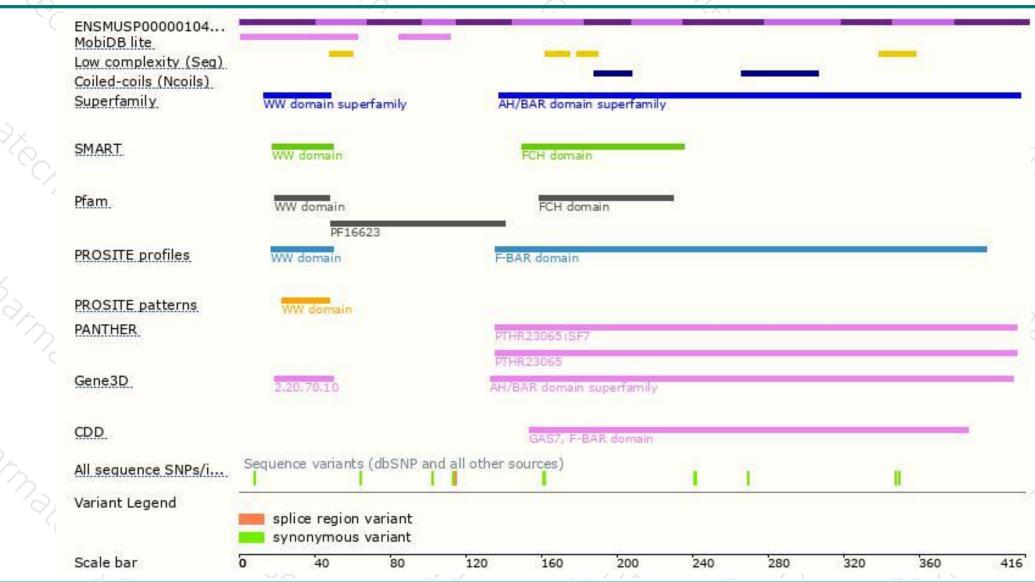
### Genomic location distribution





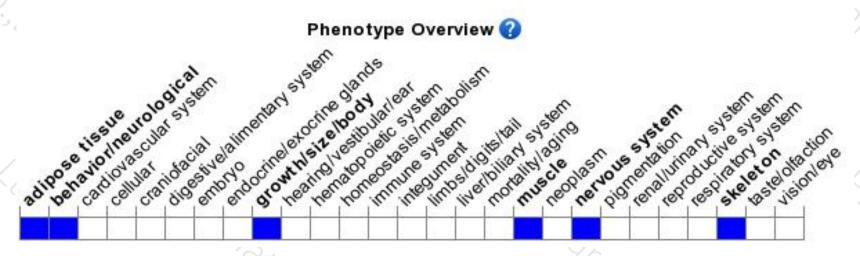
### Protein domain





## 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, Aged mice homozygous for a hypomorphic allele show impaired coordination, decreased grip strength, decreased motor neuron number, muscle weakness, abnormal skeletal muscle fiber type ratio in the soleus muscle, and defects in motor neuron axon terminal sprouting.



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





