

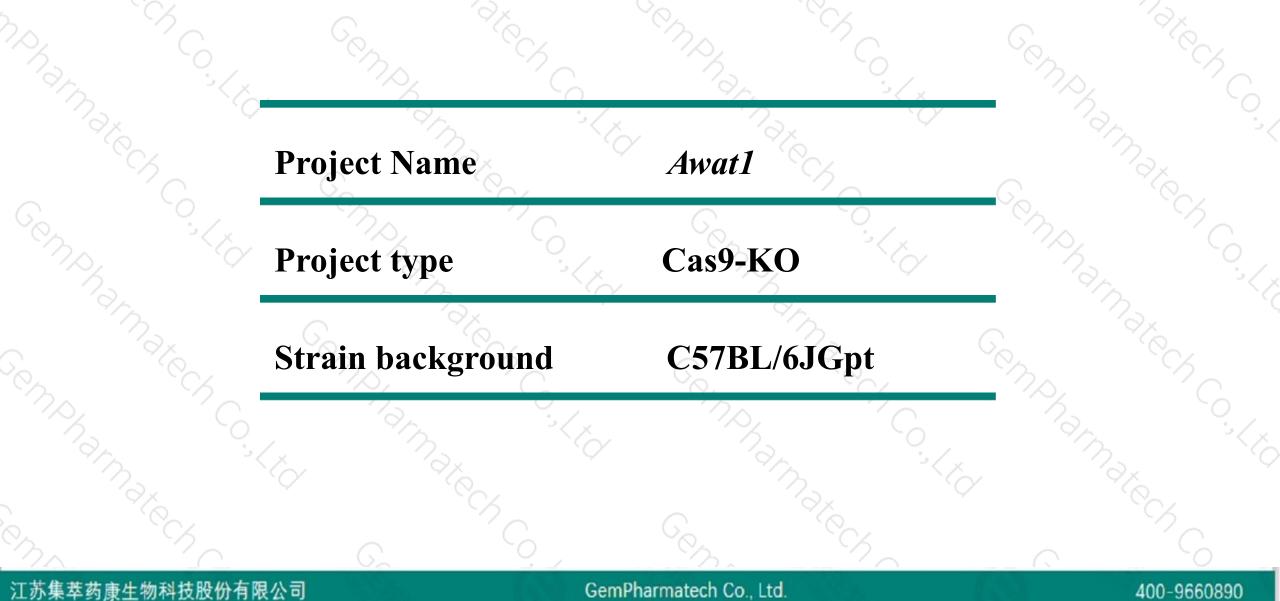
# Awat1 Cas9-KO Strategy

Designer:Xiaojing Li Reviewer:JiaYu Design Date:2020-4-7

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## **Project Overview**

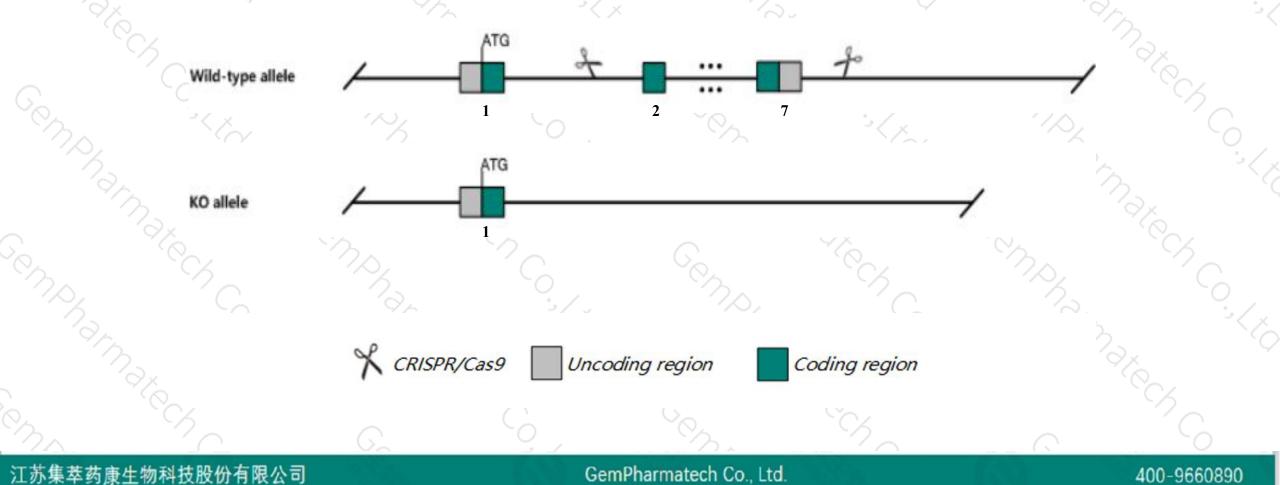




# **Knockout** strategy



This model will use CRISPR/Cas9 technology to edit the Awat1 gene. The schematic diagram is as follows:





- The Awat1 gene has 1 transcript. According to the structure of Awat1 gene, exon2-exon7 of Awat1-201 (ENSMUST0000096361.4) transcript is recommended as the knockout region. The region contains 911bp coding sequence. Knock out the region will result in disruption of protein function.
- > In this project we use CRISPR/Cas9 technology to modify Awat1 gene. The brief process is as follows: CRISPR/Cas9 system



- The Awat1 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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### Awat1 acyl-CoA wax alcohol acyltransferase 1 [Mus musculus (house mouse)]

Gene ID: 245533, updated on 13-Mar-2020

### Summary

Official Symbol	Awat1 provided by MGI						
Official Full Name	acyl-CoA wax alcohol acyltransferase 1 provided by <u>MGI</u>						
Primary source	<u>GI:MGI:3588200</u>						
See related	nsembl:ENSMUSG0000015665						
Gene type	protein coding						
RefSeq status	VALIDATED						
Organism	Mus musculus						
Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;							
	Muroidea; Muridae; Murinae; Mus; Mus						
Also known as	Dgat2I3						
Expression	Low expression observed in reference datasetSee more						
Orthologs	human all						

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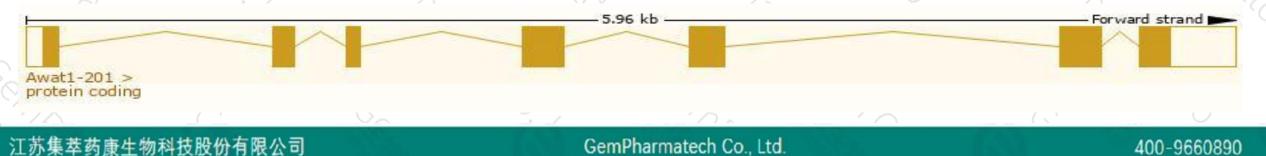
# **Transcript information (Ensembl)**



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

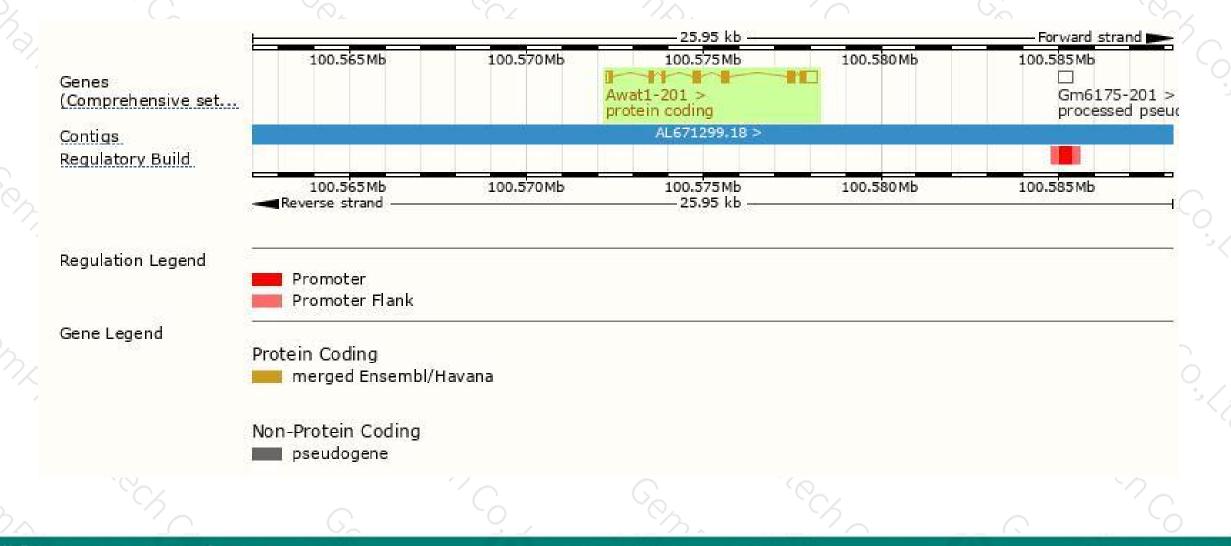
Name 🝦	Transcript ID	bp 👙	Protein 🝦	Biotype 🍦	CCDS 🝦	UniProt 🝦	Flags 🍦		
Awat1-201	ENSMUST0000096361.4	1390	<u>328aa</u>	Protein coding	<u>CCDS41074</u> &	<u>A2ADU9</u> &	TSL:1	GENCODE basic	APPRIS P1

The strategy is based on the design of Awat1-201 transcript, The transcription is shown below



### **Genomic location distribution**





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## **Protein domain**



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	ENSMUSP00000094 Transmembrane heli Low complexity (Seg) Pfam	Disculatures	ol acyltransferase		•	716 716	
Con.	PANTHER	PTHR12317 PTHR12317:SF16	or acyltransierase				- C
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	All sequence SNPs/i	Sequence variants (db:	SNP and all other so	ources)	0	11.1	
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If you have any questions, you are welcome to inquire. Tel: 400-9660890



