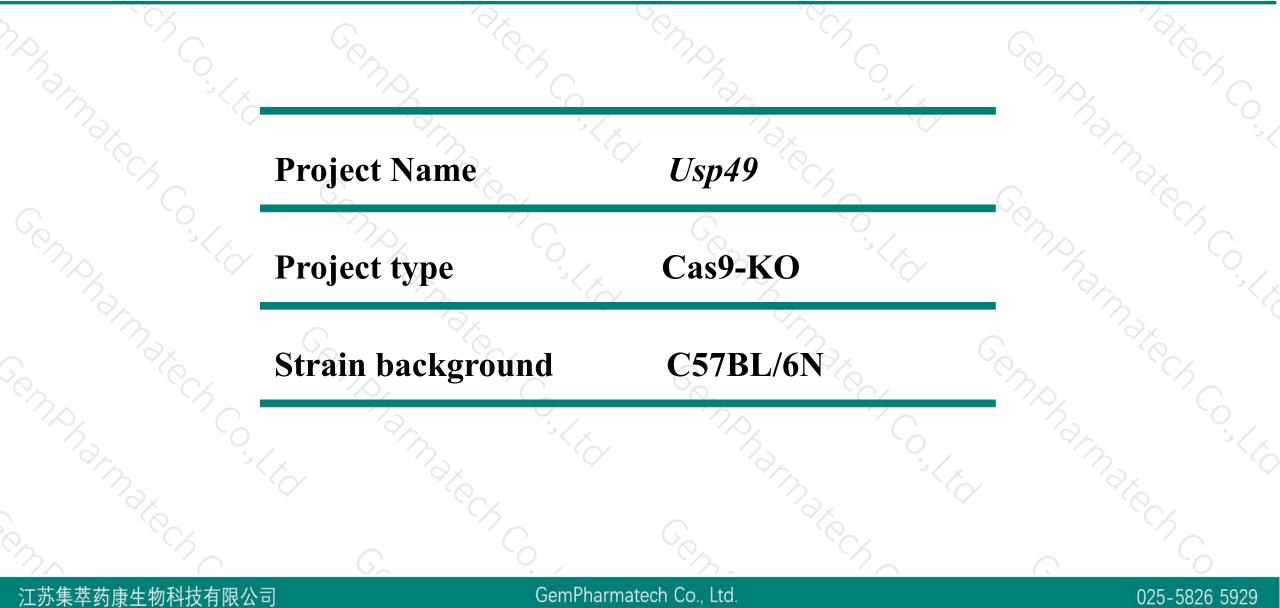


# Usp49 Cas9-KO Strategy

Designer: Daohua Xu Reviewer: Huimin Su Date: 2019/9/29

### **Project Overview**



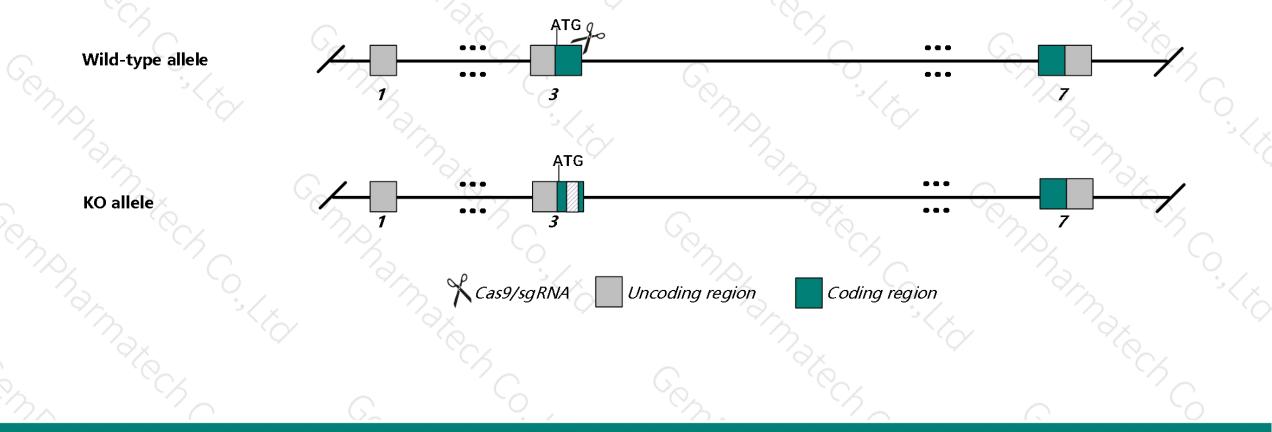


### **Knockout** strategy



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This model will use CRISPR/Cas9 technology to edit the Usp49 gene. The schematic diagram is as follows:



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In this project we use CRISPR/Cas9 technology to modify Usp49 gene. The brief process is as follows: sgRNA was transcribed in vitro.Cas9 and sgRNA were microinjected into the fertilized eggs of C57BL/6N 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/6N mice.



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- The Usp49 gene is located on the Chr17. 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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#### Usp49 ubiquitin specific peptidase 49 [ Mus musculus (house mouse) ]

Gene ID:	224836,	updated on	12-Aug-2019
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Summary

 Official Symbol
 Usp49 provided by MGI

 Official Full Name
 ubiquitin specific peptidase 49 provided by MGI

 Primary source
 MGI:MGI:2685391

 See related
 Ensembl:ENSMUSG0000090115

 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
 Gm545; C330046L10Rik

 Expression
 Ubiquitous expression in testis adult (RPKM 16.6), CNS E14 (RPKM 9.6) and 27 other tissues See more

 Orthologs
 human all

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## **Transcript information (Ensembl)**



### The gene has 4 transcripts, all transcripts are shown below:

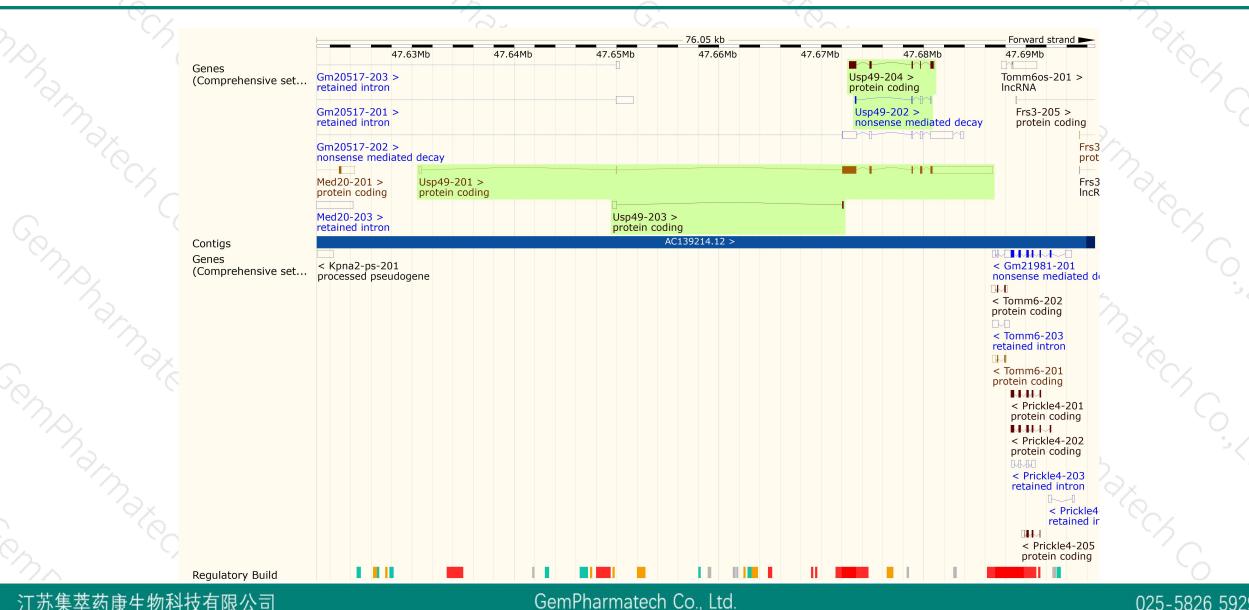
-				0000		
Transcript ID	bp 🍦	Protein 🝦	Biotype	CCDS 🍦	UniProt 🍦	Flags
ENSMUST0000024779.14	8252	<u>685aa</u>	Protein coding	<u>CCDS28852</u> &	<u>Q6P9L4</u> &	TSL:1 GENCODE basic APPRIS P1
ENSMUST00000152724.1	1443	<u>453aa</u>	Protein coding	-	<u> </u>	CDS 5' incomplete TSL:1
ENSMUST00000145314.1	544	<u>37aa</u>	Protein coding	-	<u>D3YZI5</u> @	CDS 3' incomplete TSL:2
ENSMUST00000131550.1	480	<u>52aa</u>	Nonsense mediated decay	-	<u>F6VK47</u> @	CDS 5' incomplete TSL:5
	~	3	<u>`0</u>	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~		$\times$ $1/O_{L}$
		Qr.	"3 ( x	2		$q \sim \gamma_{2}$
			$\sim$	25		
	ENSMUST00000152724.1 ENSMUST00000145314.1	ENSMUST0000024779.148252ENSMUST00000152724.11443ENSMUST00000145314.1544ENSMUST00000131550.1480	ENSMUST0000024779.14         8252         685aa           ENSMUST00000152724.1         1443         453aa           ENSMUST00000145314.1         544         37aa	ENSMUST0000024779.148252685aaProtein codingENSMUST00000152724.11443453aaProtein codingENSMUST00000145314.154437aaProtein codingENSMUST00000131550.148052aaNonsense mediated decay	ENSMUST0000024779.14         8252         685aa         Protein coding         CCDS28852 Image: CCDS28851 Image: CCDS28851 Image: CCDS28851 Image: CCDS28851 Image: CCDS	ENSMUST0000024779.14         8252         685aa         Protein coding         CCDS28852@         Q6P9L4@           ENSMUST00000152724.1         1443         453aa         Protein coding         -         Q8BUQ6@           ENSMUST00000145314.1         544         37aa         Protein coding         -         D3YZI5@           ENSMUST00000131550.1         480         52aa         Nonsense mediated decay         -         F6VK47@

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### **Genomic location distribution**



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



