

Mcm7 Cas9-KO Strategy

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Overview

Target Gene Name

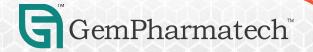
• Mcm7

Project Type

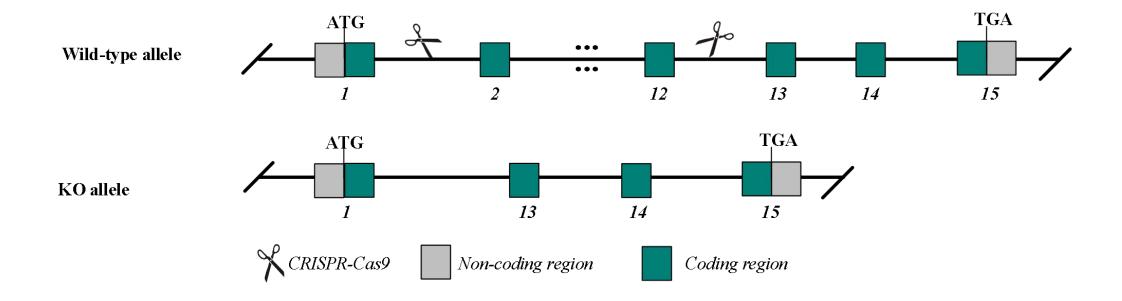
• Cas9-KO

Genetic Background

• C57BL/6JGpt



Strain Strategy



Technical Information

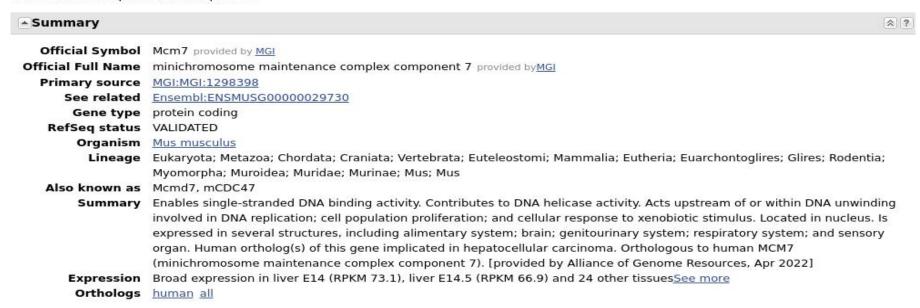
- The *Mcm7* gene has 10 transcripts. According to the structure of *Mcm7* gene, exon 2-exon 12 of *Mcm7*-201 (ENSMUST0000000505.16) transcript is recommended as the knockout region. The region contains 1648 bp coding sequence. Knocking out the region will result in disruption of protein function.
- In this project we use CRISPR-Cas9 technology to modify *Mcm7* gene. The brief process is as follows: gRNAs were transcribed in vitro. Cas9 and gRNAs were microinjected into the fertilized eggs of C57BL/6JGpt mice. Fertilized eggs were transplanted to obtain positive F0 mice which were confirmed by PCR and on-target amplicon sequencing. A stable F1-generation mouse strain was obtained by mating positive F0-generation mice with C57BL/6JGpt mice and confirmation of the desired mutant allele was carried out by PCR and on-target amplicon sequencing.



Gene Information

Mcm7 minichromosome maintenance complex component 7 [Mus musculus (house mouse)]

Gene ID: 17220, updated on 12-Apr-2023



Source: https://www.ncbi.nlm.nih.gov/

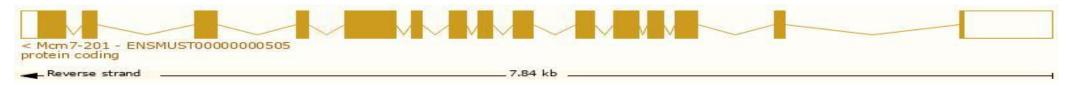


Transcript Information

The gene has 10 transcripts, all transcripts are shown below:

Transcript ID	Name A	bp ≜	Protein A	Biotype	CCDS A	UniProt Match A	Flags			
ENSMUST00000000505.16	100 Table (100 Table (2962	719aa	Protein coding	CCDS19793 ₽	Q3U4T8₽ Q61881₽	Ensembl Canonical		APPRIS P1	TSL:1
ENSMUST00000148879.8	Mcm7-208	1351	389aa	Protein coding		D3Z6N3₽	GENCODE basic TSL:5			
ENSMUST00000139983.8	Mcm7-204	829	<u>110aa</u>	Protein coding		<u>D3Z0J6</u> ₽	TSL:3 CDS 3' incomplete			
ENSMUST00000153867.8	Mcm7-209	795	<u>142aa</u>	Protein coding		<u>D3Z2K9</u> ₽	TSL:5 CDS 3' incomplete			
ENSMUST00000155902.8	Mcm7-210	736	<u>110aa</u>	Protein coding		<u>D3Z0J6</u> ₽	TSL:5 CDS 3' incomplete			
ENSMUST00000148094.8	Mcm7-207	705	<u>50aa</u>	Protein coding		D3Z335₽	TSL:2 CDS 3' incomplete			
ENSMUST00000147920.2	Mcm7-206	660	6aa	Protein coding		7.7	TSL:3 CDS 3' incomplete			
ENSMUST00000142254.8	Mcm7-205	1023	No protein	Retained intron		199	TSL:5			
ENSMUST00000125316.2	Mcm7-202	986	No protein	Retained intron) (*)	TSL:2			
ENSMUST00000139223.2	Mcm7-203	629	No protein	Retained intron		-	TSL:2			

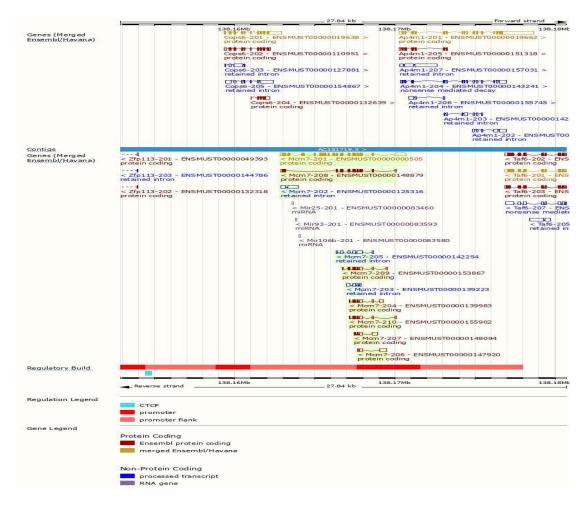
The strategy is based on the design of *Mcm7*-201 transcript, the transcription is shown below:



Source: https://www.ensembl.org

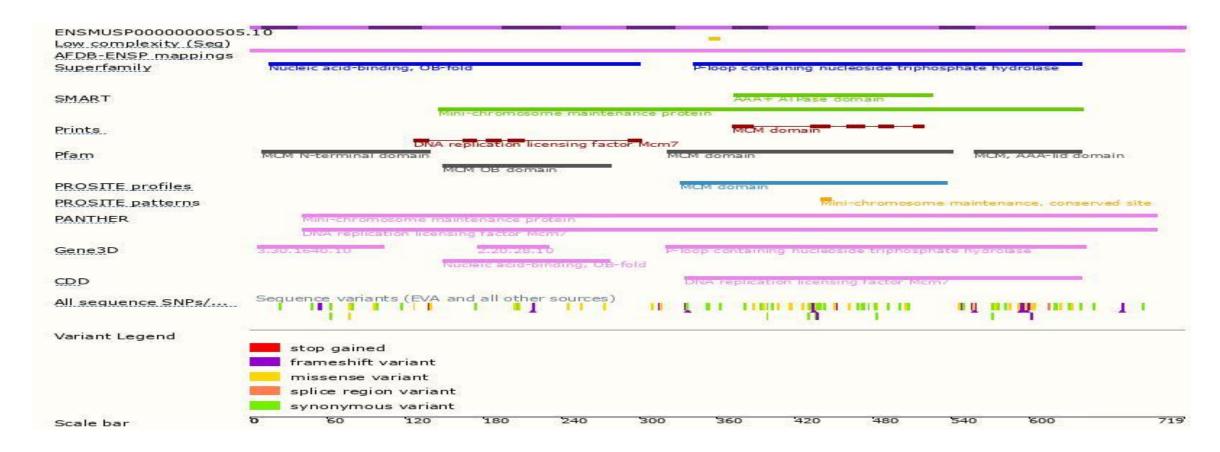


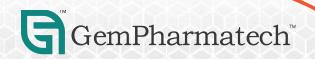
Genomic Information





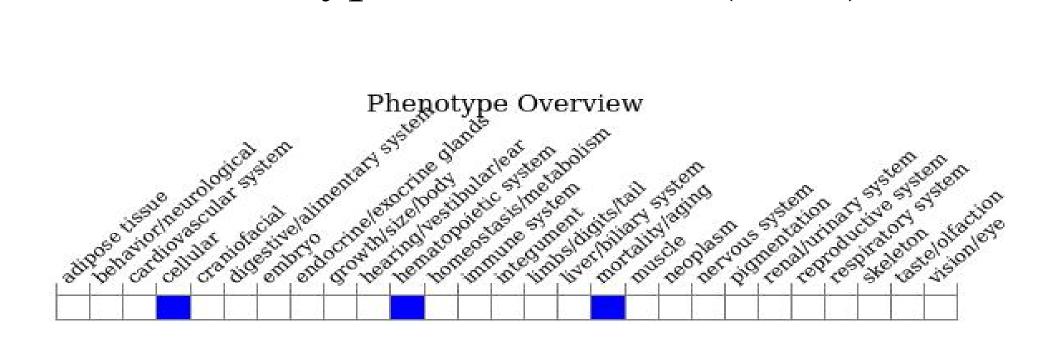
Protein Information





Source: : https://www.ensembl.org

Mouse Phenotype Information (MGI)



• Mice homozygous for a gene trapped allele exhibit prenatal lethality. Mice heterozygous for this allele exhibit increased micronulei-containing red blood cells.



Source: https://www.informatics.jax.org

Important Information

- According to the existing MGI data, Mice homozygous for a gene trapped allele exhibit prenatal lethality. Mice heterozygous for this allele exhibit increased micronulei-containing red blood cells.
- The influence of the transcript Mcm7-204&Mcm7-206&Mcm7-207&Mcm7-209&Mcm7-210 is unknown.
- The KO region may affect the function of Ap4m1&Cops6 &Mir25 gene.
- *Mcm7* is located on Chr5. If the knockout mice are crossed with other mouse strains to obtain double homozygous mutant offspring, please avoid the situation that the second gene is on the same chromosome.
- This Strategy is designed based on genetic information in existing databases. Due to the complexity of biological processes, all risks of the mutation on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

