

# Spag4 Cas9-KO Strategy

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#### Overview

#### Target Gene Name

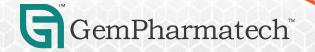
• Spag4

### Project Type

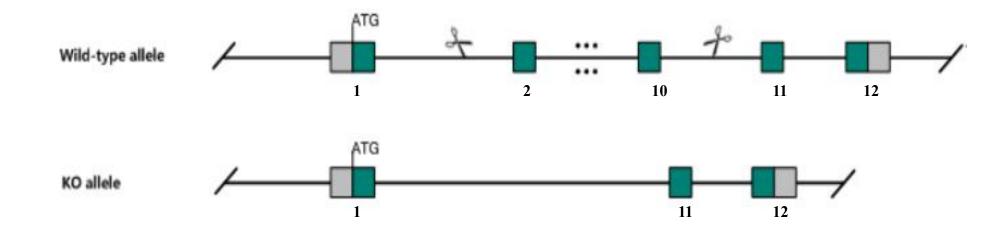
• Cas9-KO

#### Genetic Background

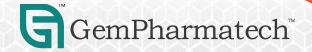
• C57BL/6JGpt



## Strain Strategy







#### **Technical Information**

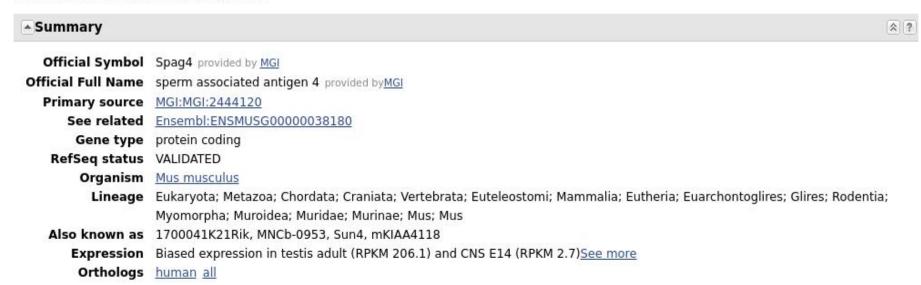
- The *Spag4* gene has 8 transcripts. According to the structure of *Spag4* gene, exon2-exon10 of *Spag4*-201 (ENSMUST00000038860.12) transcript is recommended as the knockout region. The region contains 773bp coding sequence. Knocking out the region will result in disruption of protein function.
- In this project we use CRISPR-Cas9 technology to modify *Spag4* 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 ontarget 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

#### Spag4 sperm associated antigen 4 [Mus musculus (house mouse)]

Gene ID: 245865, updated on 13-Mar-2020



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

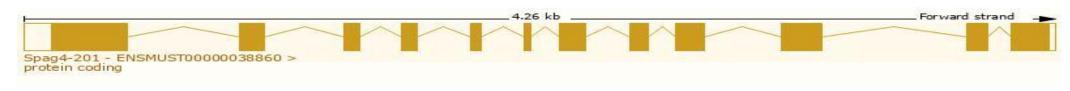


### Transcript Information

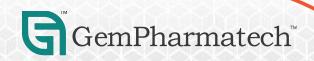
The gene has 8 transcripts, all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Spag4-201	ENSMUST00000038860.11	1473	443aa	Protein coding	CCDS50777	Q9JJF2	TSL:1 GENCODE basic APPRIS P1
Spag4-206	ENSMUST00000137966.1	747	210aa	Protein coding	-	B7ZCP3	CDS 3' incomplete TSL:3
Spag4-207	ENSMUST00000138178.7	438	<u>78aa</u>	Nonsense mediated decay	· ·	F65430	CDS 5' incomplete TSL:3
Spag4-203	ENSMUST00000131144.7	370	40aa	Nonsense mediated decay	-	F6XNA5	CDS 5' incomplete TSL:3
Spag4-208	ENSMUST00000149139.1	667	No protein	Processed transcript	¥	-	TSL:5
Spag4-202	ENSMUST00000125996.7	3892	No protein	Retained intron	-5	0.72	TSL:2
Spag4-205	ENSMUST00000136765.7	2203	No protein	Retained intron	- 1		TSL:1
Spag4-204	ENSMUST00000133511.7	2022	No protein	Retained intron	-	122	TSL:1

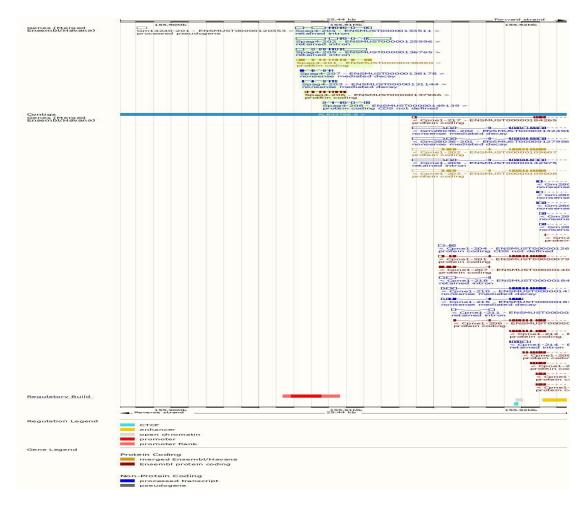
The strategy is based on the design of *Spag4*-201 transcript, the transcription is shown below:



Source: https://www.ensembl.org



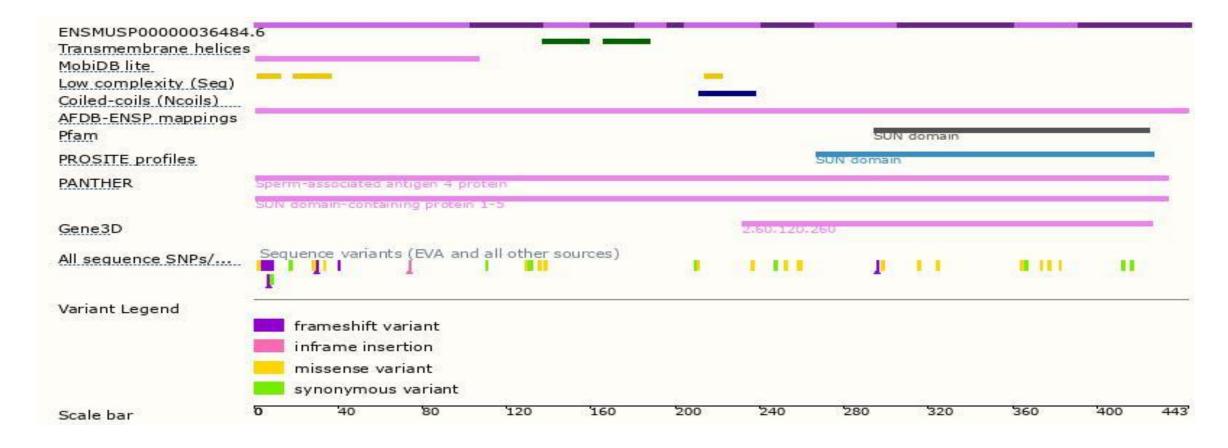
#### Genomic Information





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

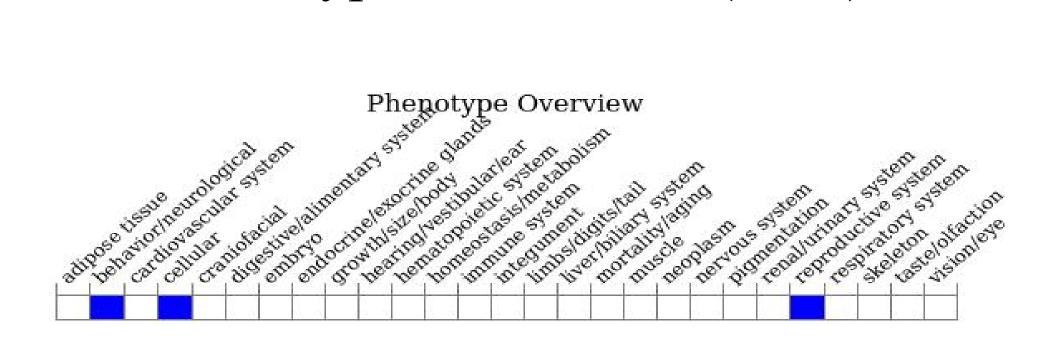
#### Protein Information





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

### Mouse Phenotype Information (MGI)



• Mice homozygous for a knock-out allele show disrupted spermiogenesis, severe defects in sperm head formation, abnormal manchette morphology, globozoospermia, and male infertility.



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

### Important Information

- *Spag4* is located on Chr2. 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.
- Intron 1-2(462 bp) and 10-11(598 bp), the effect of LOXP insertion on gene is unknown.

