

Muc4 Cas9-KO Strategy

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Design Date: 2020-7-24

Project Overview



Project Name

Muc4

Project type

Cas9-KO

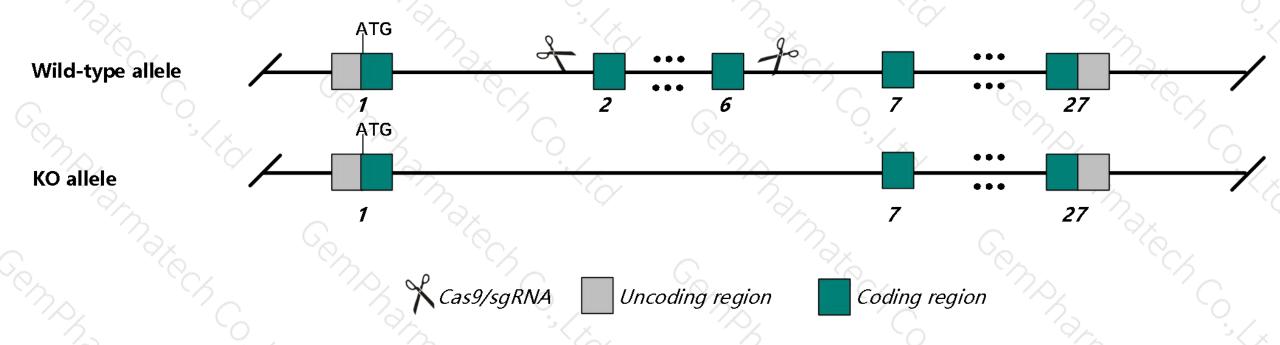
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Muc4* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Muc4* gene has 8 transcripts. According to the structure of *Muc4* gene, exon2-exon6 of *Muc4-201*(ENSMUST00000096106.9) transcript is recommended as the knockout region. The region contains 7187bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Muc4* gene. The brief process is as follows: gRNA was transcribed in vitro.Cas9 and gRNA 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 sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6JGpt mice.

Notice



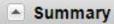
- > According to the existing MGI data, mice homozygous for a knock-out allele exhibit resistance to DSS-treated colitis and colitis-associated colorectal cancer.
- The *Muc4* gene is located on the Chr16. If the knockout mice are crossed with other mice strains to obtain double gene positive homozygous 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)



Muc4 mucin 4 [Mus musculus (house mouse)]

Gene ID: 140474, updated on 22-Jul-2020



☆ ?

Official Symbol Muc4 provided by MGI

Official Full Name mucin 4 provided by MGI

Primary source MGI:MGI:2153525

See related Ensembl: ENSMUSG00000079620

RefSeq status REVIEWED
Organism Mus musculus

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;

Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus

Also known as Asgp; 4933405l11Rik

Summary The major constituents of mucus, the viscous secretion that covers epithelial surfaces such as those in the trachea, colon, and

cervix, are highly glycosylated proteins called mucins. These glycoproteins play important roles in the protection of the epithelial cells and have been implicated in epithelial renewal and differentiation. This gene encodes an integral membrane glycoprotein found on the cell surface. A large 5' exon encodes at least 15 tandem repeats of 124-126 amino acids. [provided by RefSeq, Jul

2008]

Expression Biased expression in colon adult (RPKM 15.2), large intestine adult (RPKM 5.6) and 6 other tissues See more

Orthologs human all

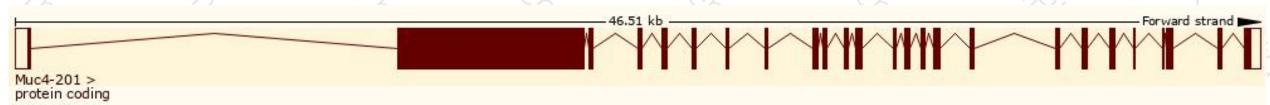
Transcript information (Ensembl)



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

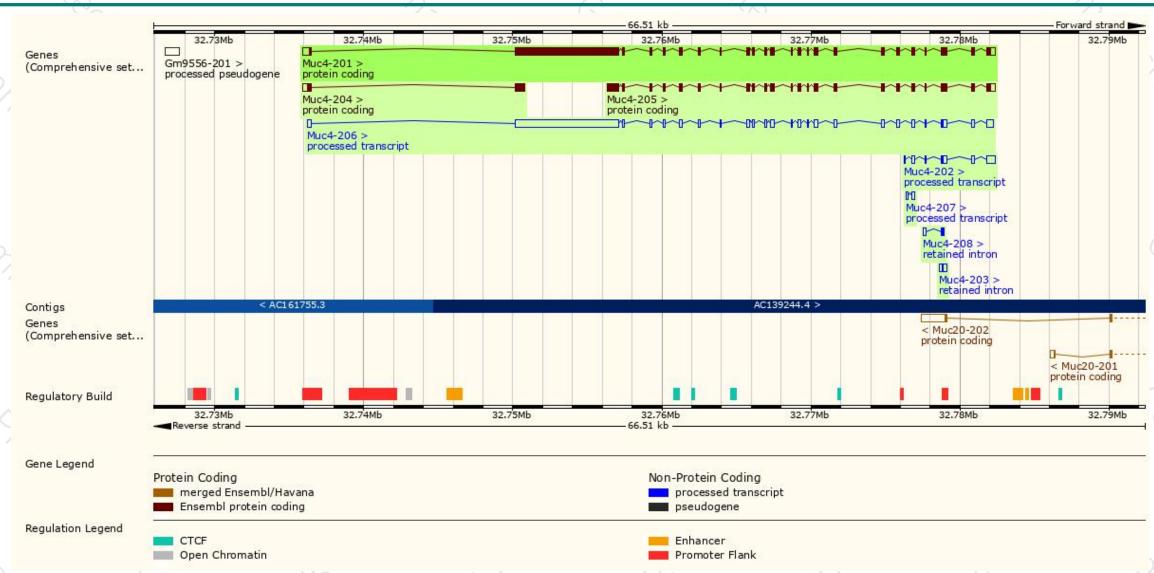
No. of Control of Cont						
Transcript ID	bp 🌲	Protein ▼	Biotype	CCDS	UniProt	Flags
ENSMUST00000096106.9	11277	3470aa	Protein coding	-	<u>E9Q7Q0</u> ₽	TSL:5 GENCODE basic APPRIS P1
ENSMUST00000135753.1	4575	<u>1401aa</u>	Protein coding	-	F6R2G3 ₺	CDS 5' incomplete TSL:1
ENSMUST00000132475.6	1190	293aa	Protein coding	-	E9Q9U1 ₽	CDS 3' incomplete TSL:1
ENSMUST00000142355.8	10805	No protein	Processed transcript	-	-	TSL:5
ENSMUST00000115119.7	1348	No protein	Processed transcript	-	-	TSL:1
ENSMUST00000143297.1	330	No protein	Processed transcript	-	-	TSL:5
ENSMUST00000131419.1	434	No protein	Retained intron	-	-	TSL:3
ENSMUST00000149773.1	297	No protein	Retained intron	-	-	TSL:3
	ENSMUST000000135753.1 ENSMUST00000132475.6 ENSMUST00000142355.8 ENSMUST00000115119.7 ENSMUST00000143297.1 ENSMUST00000131419.1	ENSMUST00000135753.1 4575 ENSMUST00000135753.1 4575 ENSMUST00000132475.6 1190 ENSMUST00000142355.8 10805 ENSMUST00000115119.7 1348 ENSMUST00000143297.1 330 ENSMUST00000131419.1 434	ENSMUST00000096106.9 11277 3470aa ENSMUST00000135753.1 4575 1401aa ENSMUST00000132475.6 1190 293aa ENSMUST00000142355.8 10805 No protein ENSMUST00000115119.7 1348 No protein ENSMUST00000143297.1 330 No protein ENSMUST00000131419.1 434 No protein	ENSMUST00000135753.1 4575 1401aa Protein coding ENSMUST00000132475.6 1190 293aa Protein coding ENSMUST00000142355.8 10805 No protein Processed transcript ENSMUST00000115119.7 1348 No protein Processed transcript ENSMUST00000143297.1 330 No protein Processed transcript ENSMUST00000131419.1 434 No protein Retained intron	ENSMUST00000096106.9 11277 3470aa Protein coding - ENSMUST00000135753.1 4575 1401aa Protein coding - ENSMUST00000132475.6 1190 293aa Protein coding - ENSMUST00000142355.8 10805 No protein Processed transcript - ENSMUST00000115119.7 1348 No protein Processed transcript - ENSMUST00000143297.1 330 No protein Processed transcript - ENSMUST00000131419.1 434 No protein Retained intron -	ENSMUST00000096106.9 11277 3470aa Protein coding - E9Q7Q0 ENSMUST00000135753.1 4575 1401aa Protein coding - F6R2G3 ENSMUST00000132475.6 1190 293aa Protein coding - E9Q9U1 ENSMUST00000142355.8 10805 No protein Processed transcript - - ENSMUST00000115119.7 1348 No protein Processed transcript - - ENSMUST00000143297.1 330 No protein Processed transcript - - ENSMUST00000131419.1 434 No protein Retained intron - -

The strategy is based on the design of *Muc4-201* transcript, the transcription is shown below:



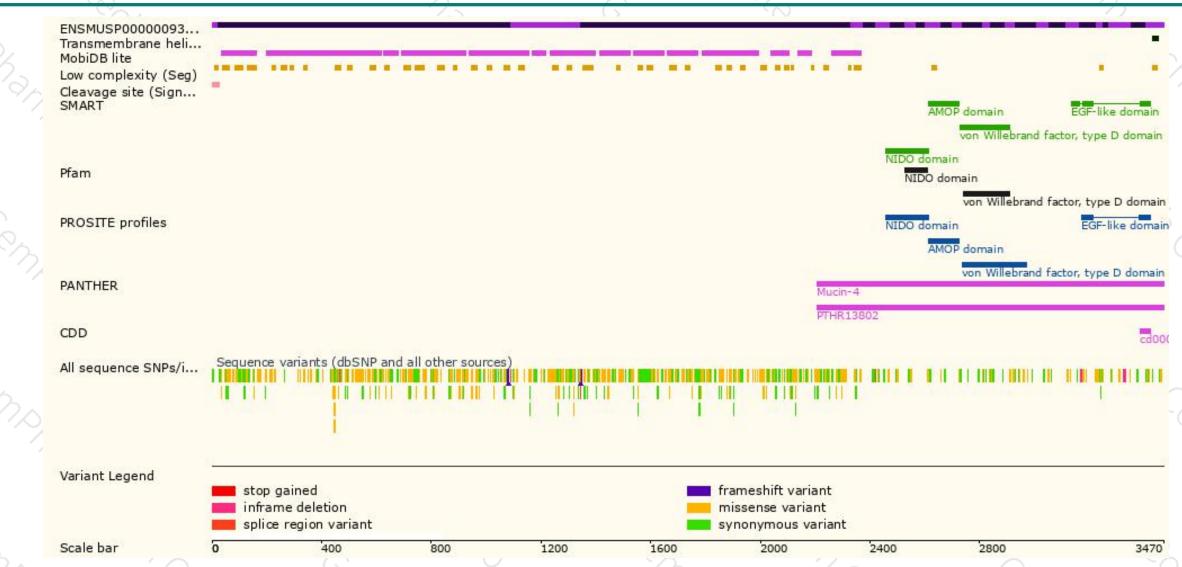
Genomic location distribution





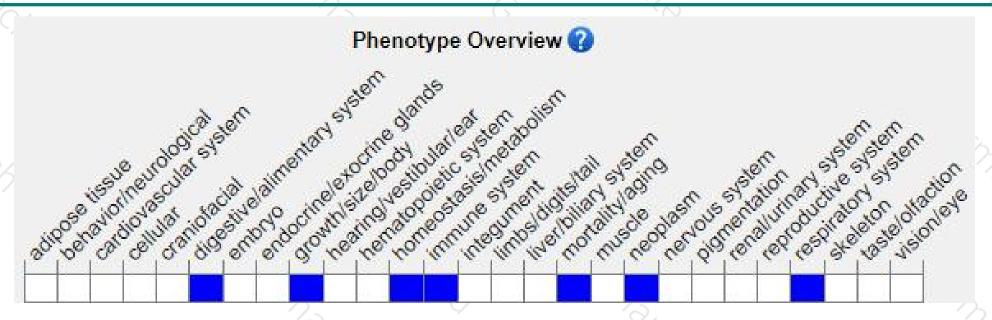
Protein domain





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, mice homozygous for a knock-out allele exhibit resistance to DSS-treated colitis

and colitis-associated colorectal cancer.



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





