

Was Cas9-CKO Strategy

Designer: Huimin Su

Reviewer: Ruirui Zhang

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Project Overview



Project Name

Project type

Cas9-CKO

Was

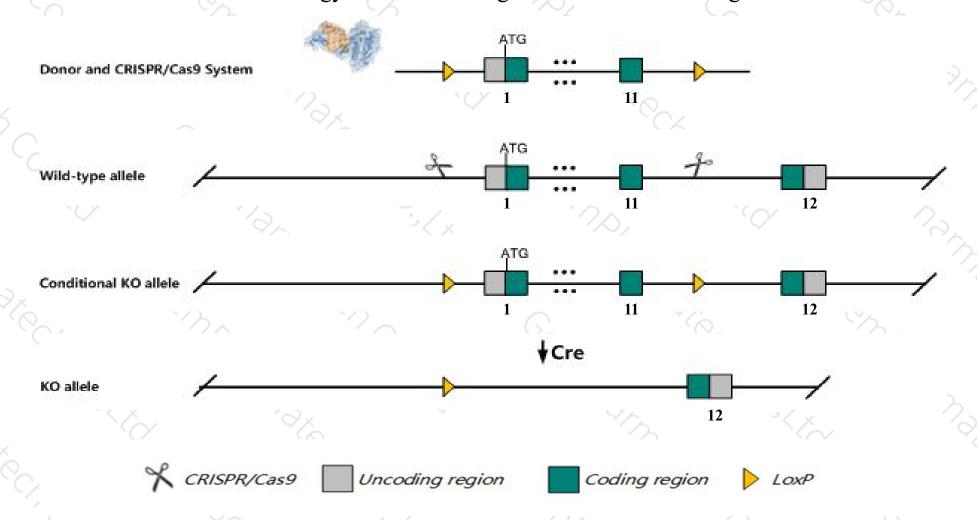
Strain background

C57BL/6JGpt

Conditional Knockout strategy



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



Technical routes



- The *Was* gene has 2 transcripts. According to the structure of *Was* gene, exon1-exon11 of *Was-201* (ENSMUST0000033505.6) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Was* gene. The brief process is as follows:CRISPR/Cas9 system and Donor 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.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

Notice



- > According to the existing MGI data, Homozygous mutant females and hemizygous mutant males exhibit reduced numbers of peripheral blood lymphocytes and platelets, but increased numbers of neutrophils.
- ➤ The 5 termial regulation of Gm6787 may be affected.
- The *Was* 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)



Was Wiskott-Aldrich syndrome [Mus musculus (house mouse)]

Gene ID: 22376, updated on 5-Nov-2019

Summary

Official Full Name Wiskott-Aldrich syndrome provided by MGI

Primary source MGI:MGI:105059

Official Symbol Was provided by MGI

See related Ensembl: ENSMUSG00000031165

Gene type protein coding
RefSeq status VALIDATED
Organism <u>Mus musculus</u>

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

Murinae; Mus; Mus

Also known as Wasp; U42471

Expression Biased expression in thymus adult (RPKM 28.4), spleen adult (RPKM 26.0) and 11 other tissues See more

Orthologs <u>human</u> all

Genomic context

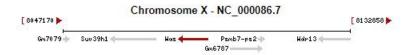
☆ ?

Location: X A1.1; X 3.65 cM

See Was in Genome Data Viewer

Exon count: 12

Annotation release	Status	Assembly	Chr	Location	
108	current	GRCm38.p6 (GCF_000001635.26)	X	NC_000086.7 (80814668090491, complement)	
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	X	NC_000086.6 (76585927667617, complement)	



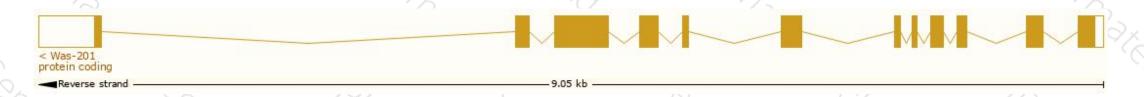
Transcript information (Ensembl)



The gene has 2 transcripts, all transcripts are shown below:

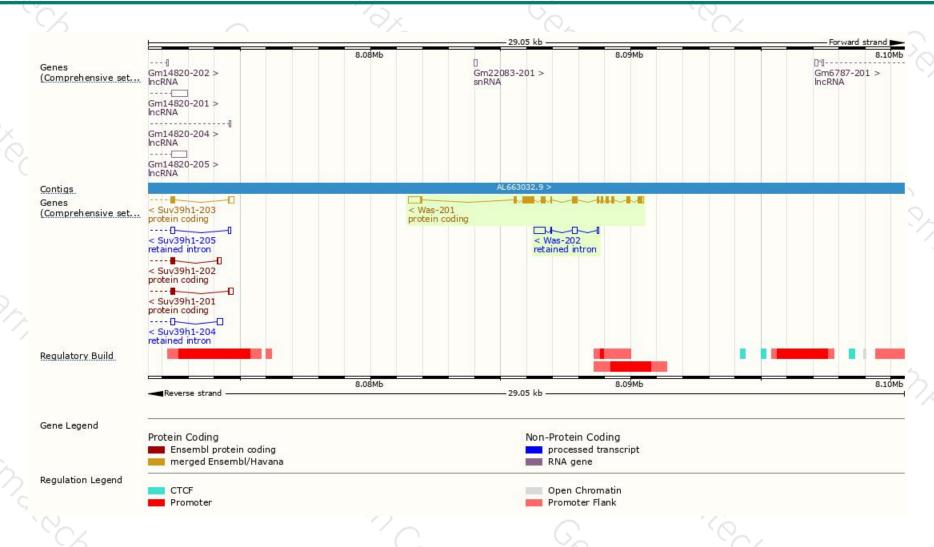
Name 🍦	Transcript ID	bp 🛊	Protein	Biotype	CCDS	UniProt	Flags		
Was-201	ENSMUST00000033505.6	2114	<u>520aa</u>	Protein coding	CCDS29984 €	P70315@Q53WY0@	TSL:1 GE	ENCODE basic	APPRIS P1
Was-202	ENSMUST00000146029.1	707	No protein	Retained intron	-	-	TSL:5		

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



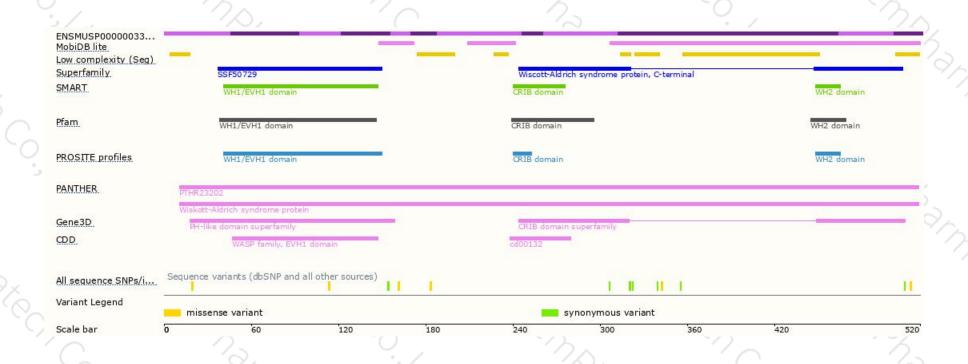
Genomic location distribution





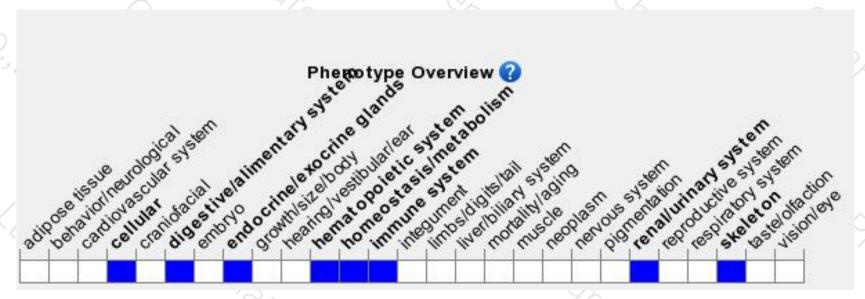
Protein domain





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, Homozygous mutant females and hemizygous mutant males exhibit reduced numbers of peripheral blood lymphocytes and platelets, but increased numbers of neutrophils.



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





