

Wdr34 Cas9-CKO Strategy

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Reviewer:

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Design Date:

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Project Overview



Project Name

Wdr34

Project type

Cas9-CKO

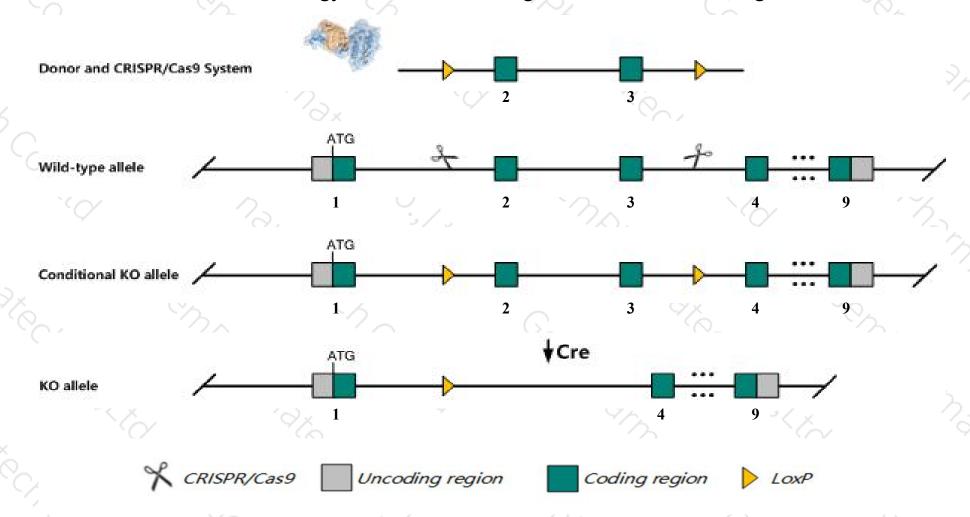
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Wdr34* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Wdr34* gene has 4 transcripts. According to the structure of *Wdr34* gene, exon2-exon3 of *Wdr34-201* (ENSMUST00000113711.2) transcript is recommended as the knockout region. The region contains 362bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Wdr34* 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



- The *Wdr34* gene is located on the Chr2. 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)



Wdr34 WD repeat domain 34 [Mus musculus (house mouse)]

Gene ID: 71820, updated on 31-Jan-2019

Summary

☆ ?

Official Symbol Wdr34 provided by MGI

Official Full Name WD repeat domain 34 provided by MGI

Primary source MGI:MGI:1919070

See related Ensembl: ENSMUSG00000039715

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 3200002l06Rik

Expression Ubiquitous expression in ovary adult (RPKM 29.0), testis adult (RPKM 27.5) and 28 other tissuesSee more

Orthologs <u>human</u> all

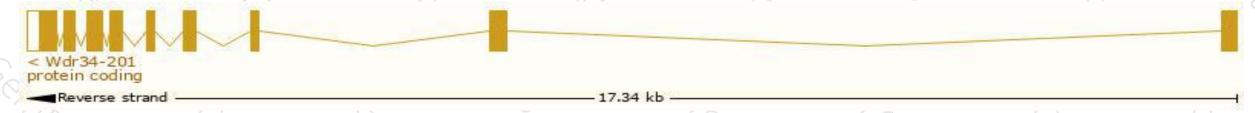
Transcript information (Ensembl)



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

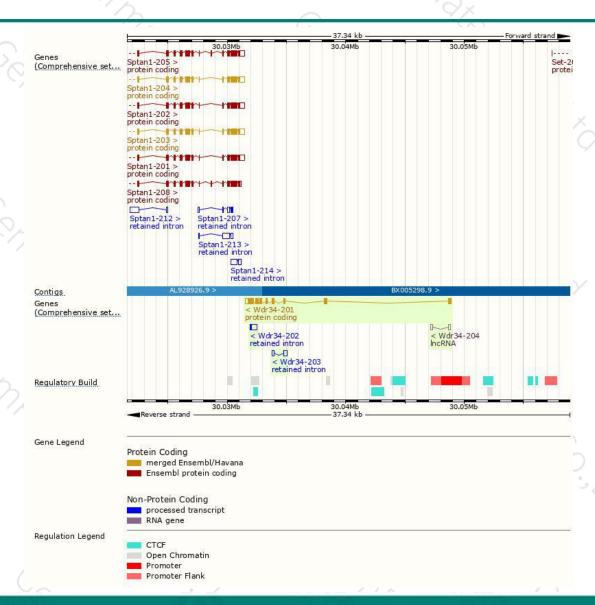
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Wdr34-201	ENSMUST00000113711.2	1814	<u>537aa</u>	Protein coding	CCDS15865	A2BE91	TSL:1 GENCODE basic APPRIS P1
Wdr34-202	ENSMUST00000139145.1	494	No protein	Retained intron	8-		TSL:2
Wdr34-203	ENSMUST00000143892.1	454	No protein	Retained intron	14	23	TSL:2
Wdr34-204	ENSMUST00000146275.1	366	No protein	IncRNA	92	25	TSL:2

The strategy is based on the design of *Wdr34-201* transcript, The transcription is shown below



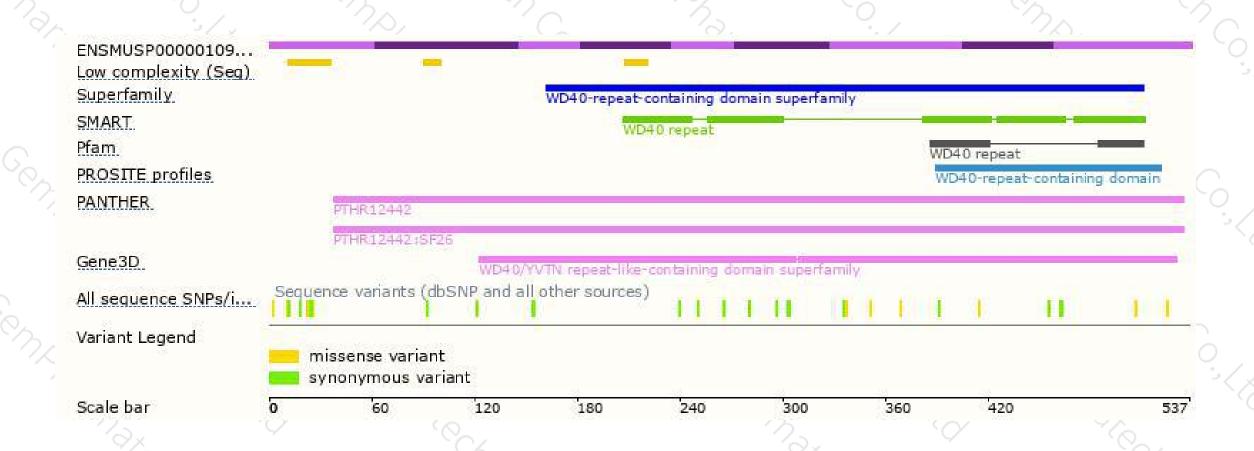
Genomic location distribution





Protein domain







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





