

# *Wdr90* Cas9-CKO Strategy

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**Reviewer: Rui Xiong**

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# Project Overview

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**Project Name**

*Wdr90*

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**Project type**

**Cas9-CKO**

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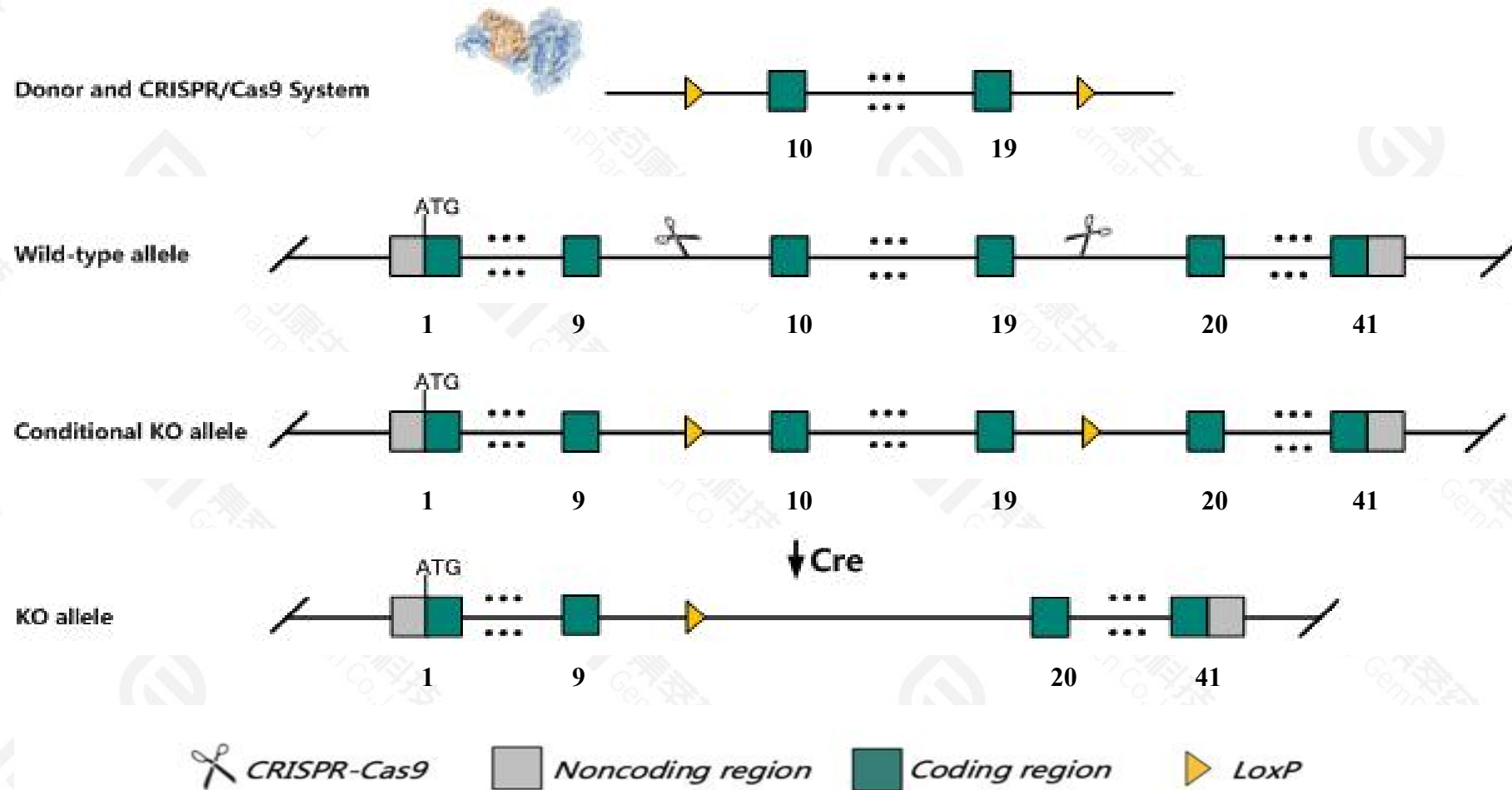
**Strain background**

**C57BL/6JGpt**

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# Conditional Knockout strategy

This model will use CRISPR-Cas9 technology to edit the *Wdr90* gene. The schematic diagram is as follows:



- The *Wdr90* gene has 12 transcripts. According to the structure of *Wdr90* gene, exon10-exon19 of *Wdr90-201*(ENSMUST00000079461.15) transcript is recommended as the knockout region. The region contains 1351bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR-Cas9 technology to modify *Wdr90* 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.



- The *Wdr90* gene is located on the Chr17. 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.
- Some amino acids will remain at the N-terminus and some functions may be retained.
- Transcript *Wdr90-212* may not be affected.
- 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)

## Wdr90 WD repeat domain 90 [Mus musculus (house mouse)]

Gene ID: 106618, updated on 25-Sep-2020

### Summary



**Official Symbol** Wdr90 provided by [MGI](#)

**Official Full Name** WD repeat domain 90 provided by [MGI](#)

**Primary source** [MGI:MGI:1921267](#)

**See related** [Ensembl:ENSMUSG00000073434](#)

**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** 3230401M21Rik, AI551153, mKIAA1924

**Expression** Ubiquitous expression in ovary adult (RPKM 17.6), thymus adult (RPKM 16.0) and 28 other tissues [See more](#)

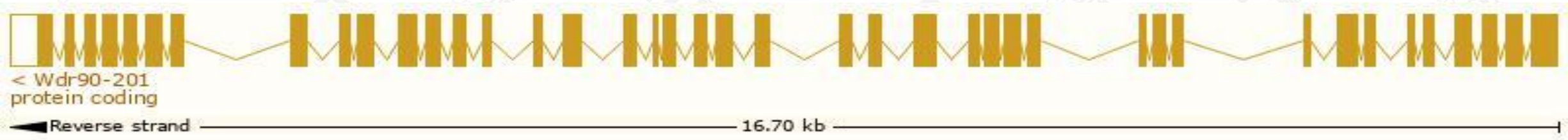
**Orthologs** [human](#) [all](#)

# Transcript information (Ensembl)

The gene has 12 transcripts,all transcripts are shown below:

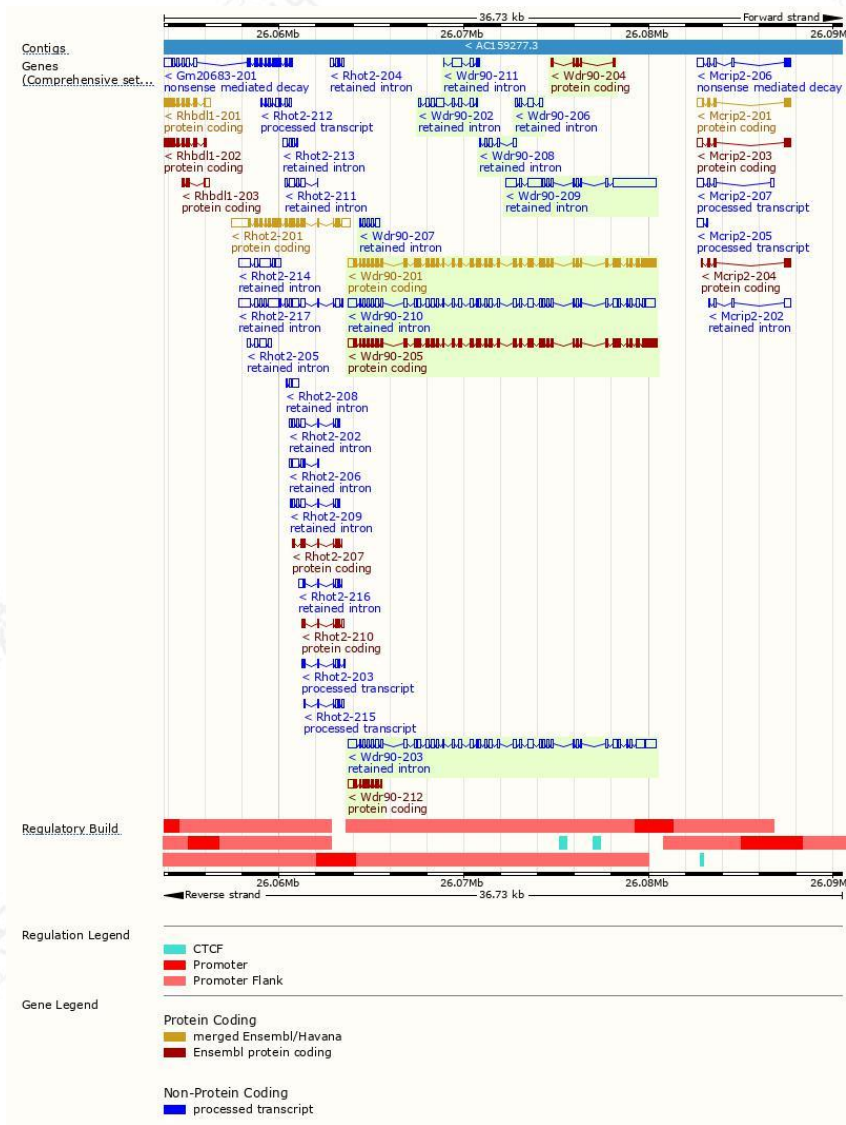
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Wdr90-201	<a href="#">ENSMUST00000079461.15</a>	5991	<a href="#">1892aa</a>	Protein coding	<a href="#">CCDS50037</a>		TSL:5 , GENCODE basic , APPRIS P2 ,
Wdr90-205	<a href="#">ENSMUST00000176923.9</a>	5962	<a href="#">1874aa</a>	Protein coding	-		TSL:5 , GENCODE basic , APPRIS ALT2 ,
Wdr90-212	<a href="#">ENSMUST00000235437.2</a>	1237	<a href="#">314aa</a>	Protein coding	-		CDS 5' incomplete ,
Wdr90-204	<a href="#">ENSMUST00000176678.2</a>	541	<a href="#">181aa</a>	Protein coding	-		CDS 5' and 3' incomplete , TSL:3 ,
Wdr90-203	<a href="#">ENSMUST00000176575.8</a>	6195	No protein	Retained intron	-		TSL:1 ,
Wdr90-210	<a href="#">ENSMUST00000177170.8</a>	5946	No protein	Retained intron	-		TSL:1 ,
Wdr90-209	<a href="#">ENSMUST00000177031.8</a>	4535	No protein	Retained intron	-		TSL:2 ,
Wdr90-202	<a href="#">ENSMUST00000176170.8</a>	1352	No protein	Retained intron	-		TSL:5 ,
Wdr90-211	<a href="#">ENSMUST00000177340.2</a>	748	No protein	Retained intron	-		TSL:5 ,
Wdr90-207	<a href="#">ENSMUST00000176983.2</a>	722	No protein	Retained intron	-		TSL:3 ,
Wdr90-206	<a href="#">ENSMUST00000176963.2</a>	667	No protein	Retained intron	-		TSL:3 ,
Wdr90-208	<a href="#">ENSMUST00000177017.2</a>	628	No protein	Retained intron	-		TSL:5 ,

The strategy is based on the design of *Wdr90-201* transcript,the transcription is shown below:



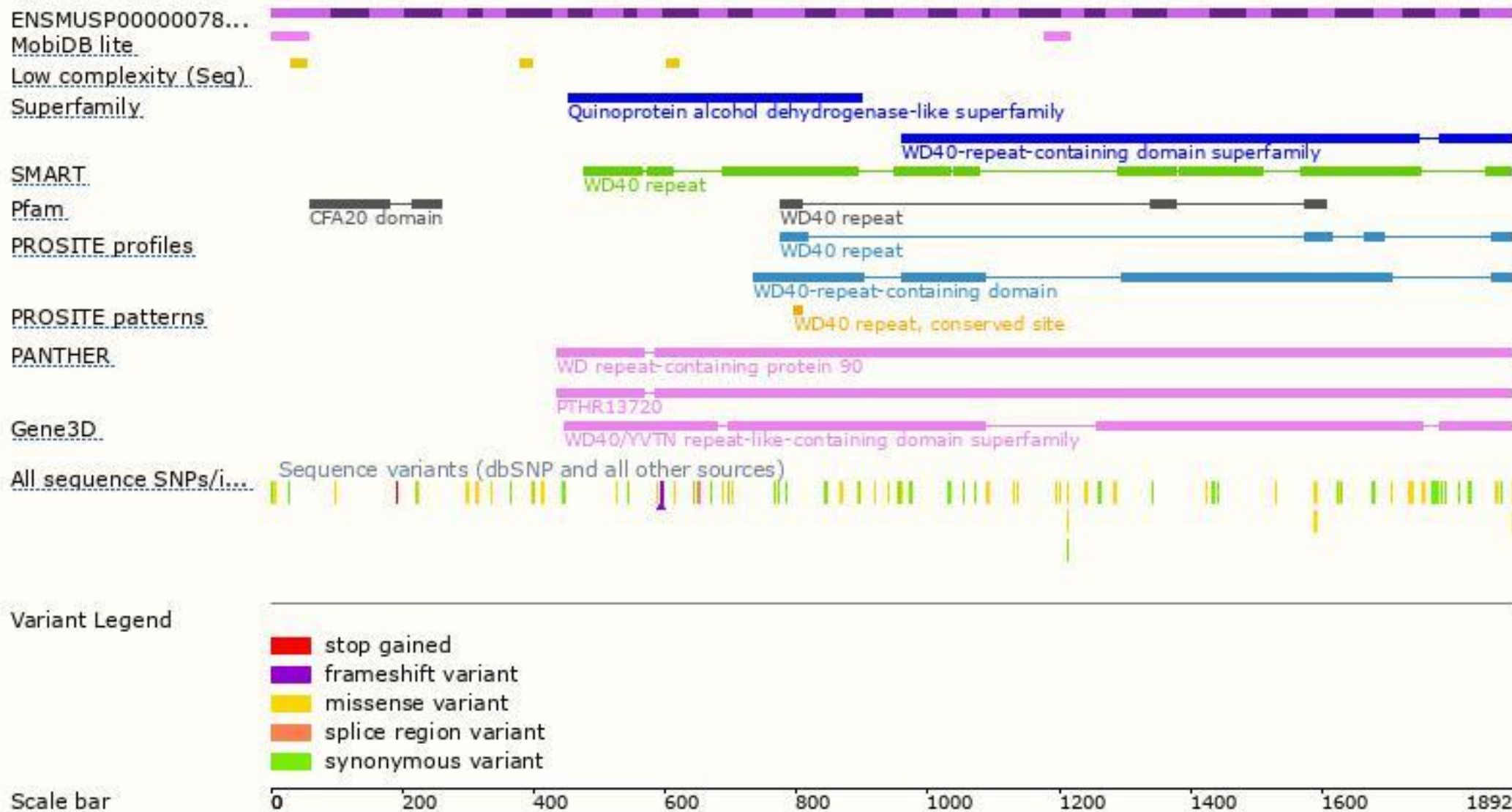


# Genomic location distribution





# Protein domain



If you have any questions, you are welcome to inquire.  
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