

# Wdr91 Cas9-CKO Strategy

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## **Project Overview**



Project Name Wdr91

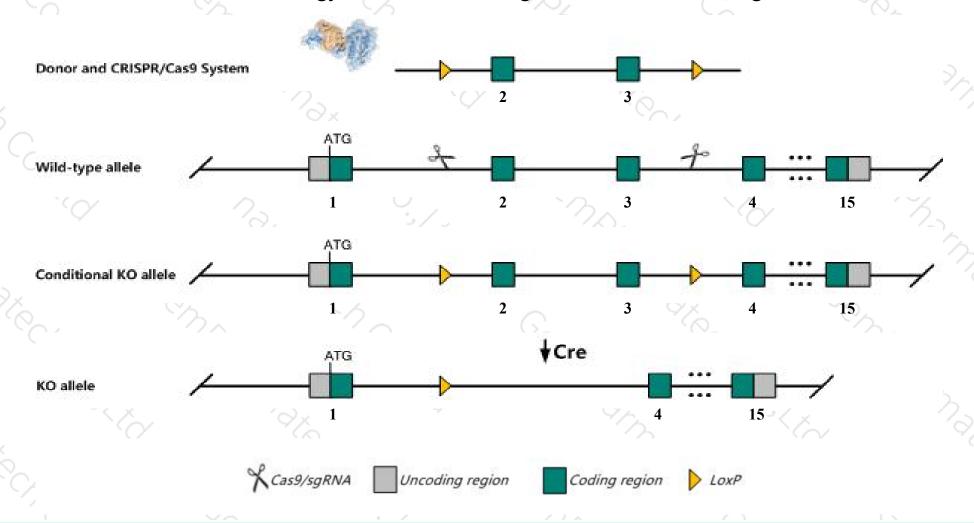
Project type Cas9-CKO

Strain background C57BL/6JGpt

## Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Wdr91* gene. The schematic diagram is as follows:



### Technical routes



- The *Wdr91* gene has 8 transcripts. According to the structure of *Wdr91* gene, exon2-exon3 of *Wdr91-201*(ENSMUST00000081214.11) transcript is recommended as the knockout region. The region contains 388bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Wdr91* gene. The brief process is as follows:sgRNA was transcribed in vitro, donor vector was constructed.Cas9, sgRNA 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 was 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,mice homozygous for a null alle exhibit neonatal lethality associated with intraabdominal bleeding. Mice homozygous for a conditional allele activated in neurons exhibit premature death, decreased brain size and weight, neuron apoptosis and reduced complexity and length of neurites.
- > The *Wdr91* gene is located on the Chr6. 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)



#### Wdr91 WD repeat domain 91 [Mus musculus (house mouse)]

Gene ID: 101240, updated on 13-Mar-2020

#### Summary

☆ ?

Official Symbol Wdr91 provided by MGI

Official Full Name WD repeat domain 91 provided by MGI

Primary source MGI:MGI:2141558

See related Ensembl:ENSMUSG00000058486

Gene type protein coding
RefSeq status PROVISIONAL
Organism Mus musculus

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

Muroidea; Muridae; Murinae; Mus; Mus

Also known as 9530020G05Rik, Al987683, AU018665

Expression Ubiquitous expression in kidney adult (RPKM 7.5), thymus adult (RPKM 7.0) and 28 other tissuesSee more

Orthologs <u>human</u> all

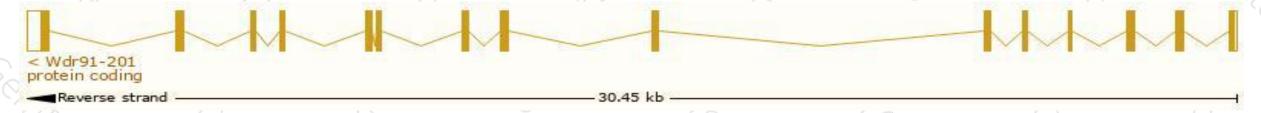
## Transcript information (Ensembl)



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

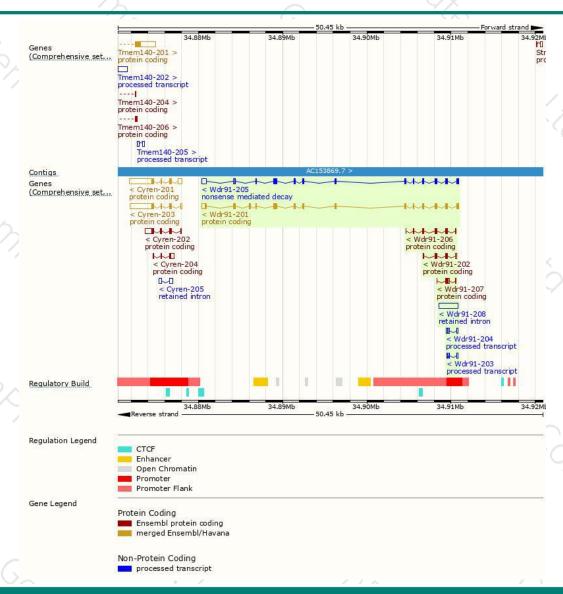
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Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Wdr91-201	ENSMUST00000081214.11	2682	748aa	Protein coding	CCDS39457	Q7TMQ7	TSL:1 GENCODE basic APPRIS P
Wdr91-206	ENSMUST00000149448.7	711	<u>196aa</u>	Protein coding	-	D3Z101	CDS 3' incomplete TSL:5
Wdr91-202	ENSMUST00000133336.7	602	<u>138aa</u>	Protein coding	ų.	D3Z0V8	CDS 3' incomplete TSL:3
Wdr91-207	ENSMUST00000152488.7	412	<u>58aa</u>	Protein coding	2	D3Z4Z6	CDS 3' incomplete TSL:5
Wdr91-205	ENSMUST00000146968.7	2569	604aa	Nonsense mediated decay	ā	S4R1X1	TSL:5
Wdr91-204	ENSMUST00000145765.1	385	No protein	Processed transcript	-		TSL:2
Wdr91-203	ENSMUST00000134830.1	340	No protein	Processed transcript	ų.	ų.	TSL:3
Wdr91-208	ENSMUST00000201569.1	2287	No protein	Retained intron		-	TSL:NA

The strategy is based on the design of *Wdr91-201* transcript, the transcription is shown below:



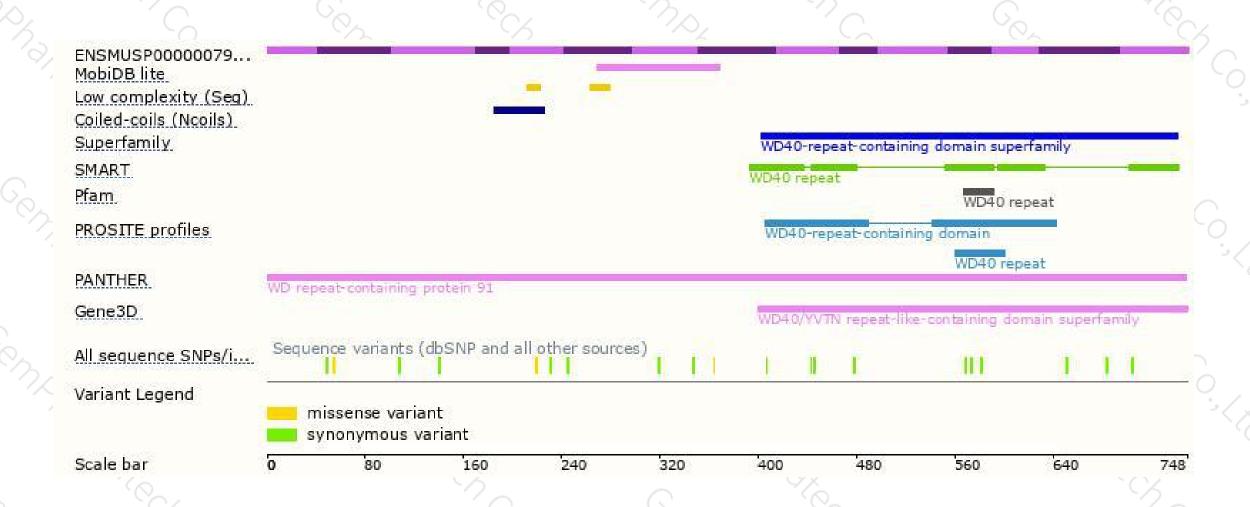
### Genomic location distribution





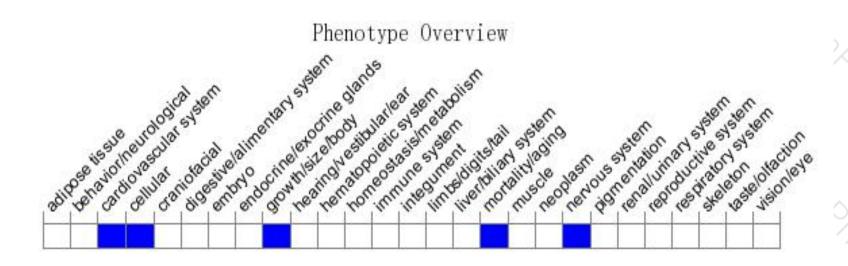
### Protein domain





## 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 null alle exhibit neonatal lethality associated with intraabdominal bleeding. Mice homozygous for a conditional allele activated in neurons exhibit premature death, decreased brain size and weight, neuron apoptosis and reduced complexity and length of neurites.



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

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