

# *Wdr60* Cas9-CKO Strategy

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# Project Overview

**Project Name**

***Wdr60***

**Project type**

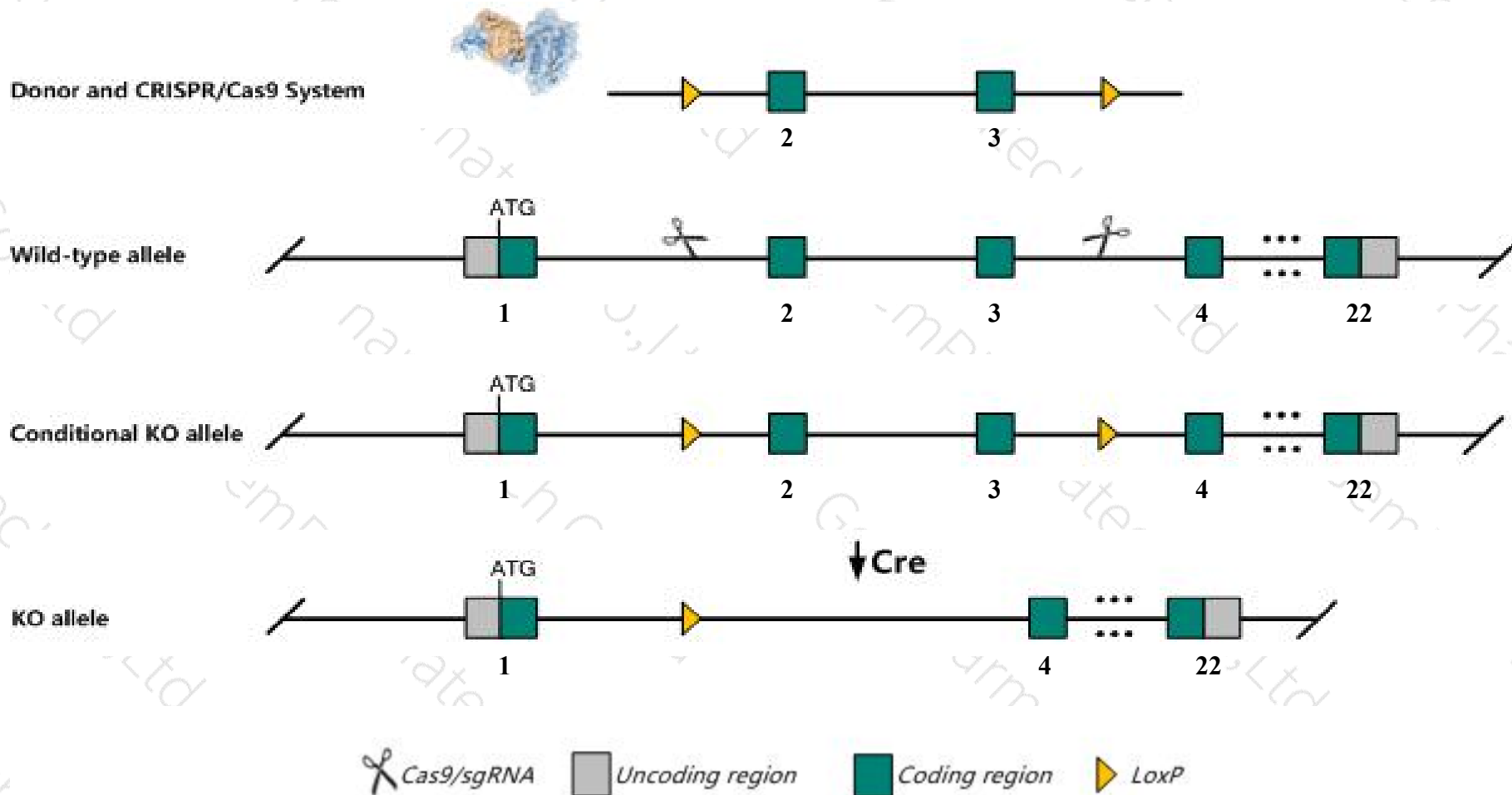
**Cas9-CKO**

**Strain background**

**C57BL/6JGpt**

# Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Wdr60* gene. The schematic diagram is as follows:



- The *Wdr60* gene has 8 transcripts. According to the structure of *Wdr60* gene, exon2-exon3 of *Wdr60-201* (ENSMUST00000039349.7) transcript is recommended as the knockout region. The region contains 475bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Wdr60* 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.

- The *Wdr60* gene is located on the Chr12. 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)

## Wdr60 WD repeat domain 60 [ *Mus musculus* (house mouse) ]

Gene ID: 217935, updated on 4-Jun-2020

### Summary



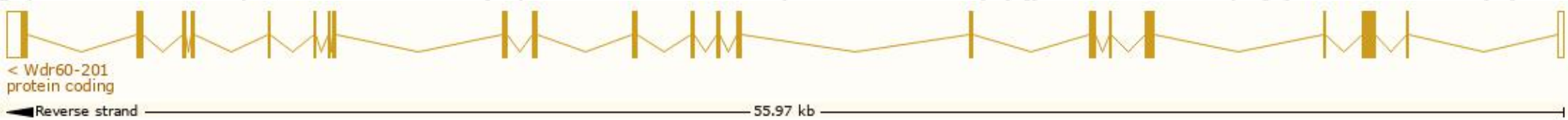
Official Symbol	Wdr60 provided by <a href="#">MGI</a>
Official Full Name	WD repeat domain 60 provided by <a href="#">MGI</a>
Primary source	<a href="#">MGI:MGI:2445085</a>
See related	<a href="#">Ensembl:ENSMUSG000000042050</a>
Gene type	protein coding
RefSeq status	VALIDATED
Organism	<a href="#">Mus musculus</a>
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	D430033N04Rik
Expression	Broad expression in testis adult (RPKM 6.3), frontal lobe adult (RPKM 3.7) and 21 other tissues <a href="#">See more</a>
Orthologs	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

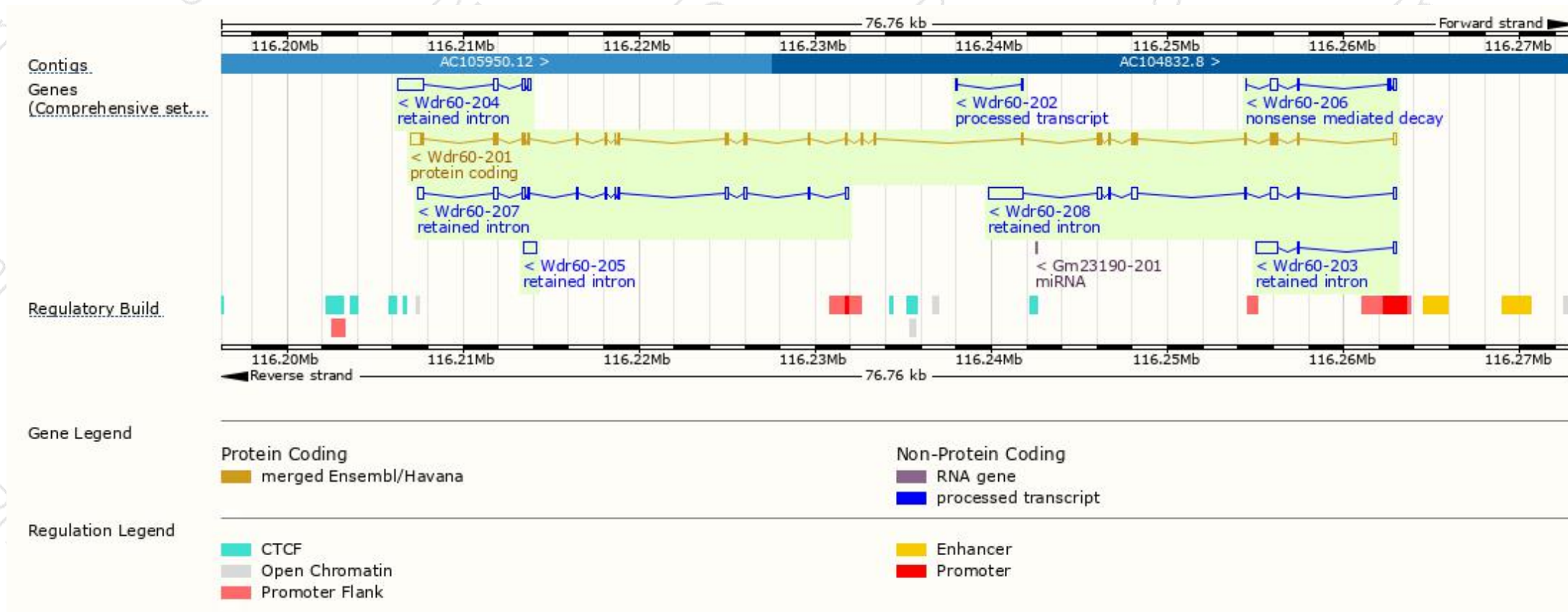
The gene has 8 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Wdr60-201	<a href="#">ENSMUST00000039349.7</a>	3721	<a href="#">999aa</a>	Protein coding	<a href="#">CCDS36575</a>	<a href="#">Q8C761</a>	TSL:1 GENCODE basic APPRIS P1
Wdr60-206	<a href="#">ENSMUST00000222679.1</a>	765	<a href="#">45aa</a>	Nonsense mediated decay	-	<a href="#">A0A1Y7VNH8</a>	TSL:1
Wdr60-202	<a href="#">ENSMUST00000220761.1</a>	194	No protein	Processed transcript	-	-	TSL:1
Wdr60-208	<a href="#">ENSMUST00000223039.1</a>	3256	No protein	Retained intron	-	-	TSL:1
Wdr60-204	<a href="#">ENSMUST00000221748.1</a>	2067	No protein	Retained intron	-	-	TSL:1
Wdr60-207	<a href="#">ENSMUST00000222764.1</a>	1747	No protein	Retained intron	-	-	TSL:1
Wdr60-203	<a href="#">ENSMUST00000221029.1</a>	1404	No protein	Retained intron	-	-	TSL:1
Wdr60-205	<a href="#">ENSMUST00000222539.1</a>	732	No protein	Retained intron	-	-	TSL:NA

The strategy is based on the design of *Wdr60-201* transcript,the transcription is shown below:

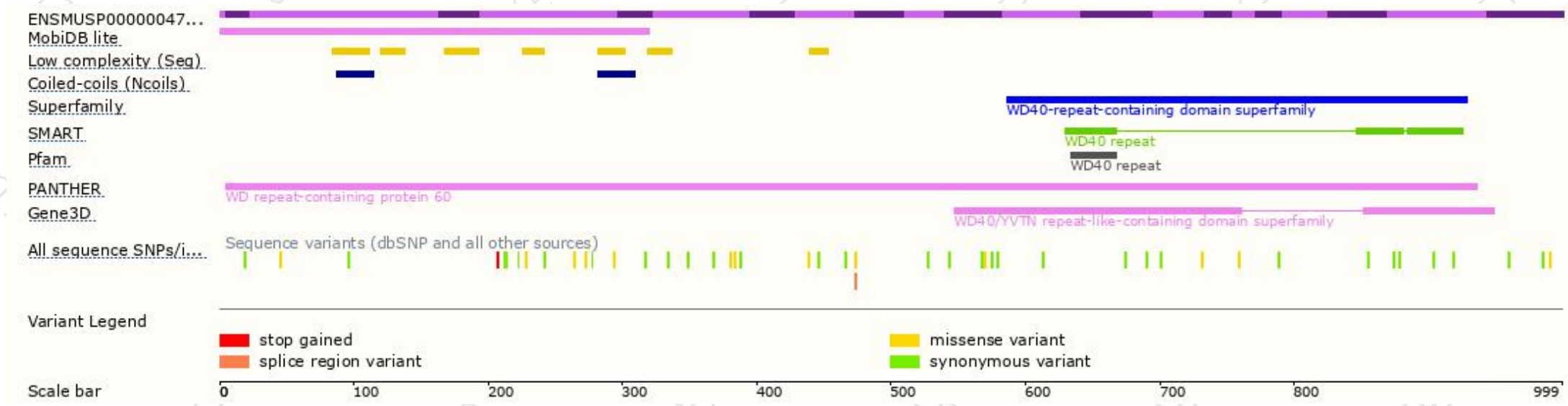


# Genomic location distribution

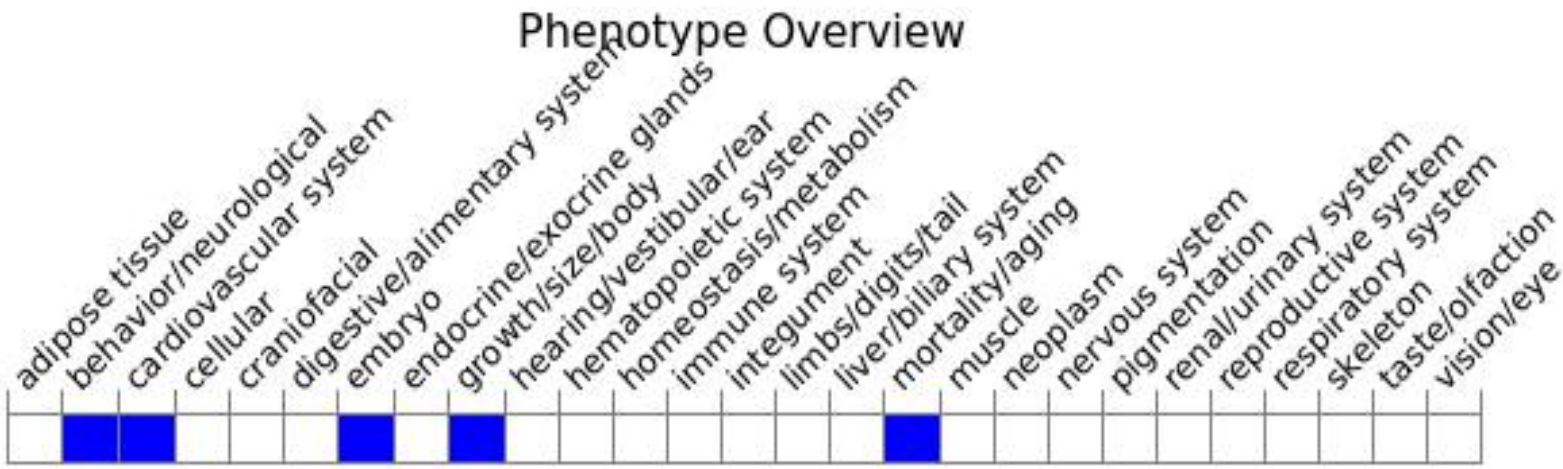




# Protein domain



# Mouse phenotype description(MGI)



*Phenotypes affected by the gene are marked in blue. Data quoted from MGI database(<http://www.informatics.jax.org/>).*

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

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