

# Ubqln4 Cas9-CKO Strategy

Designer: JiaYu

Reviewer: Xiaojing Li

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# **Project Overview**



**Project Name** 

Ubqln4

**Project type** 

Cas9-CKO

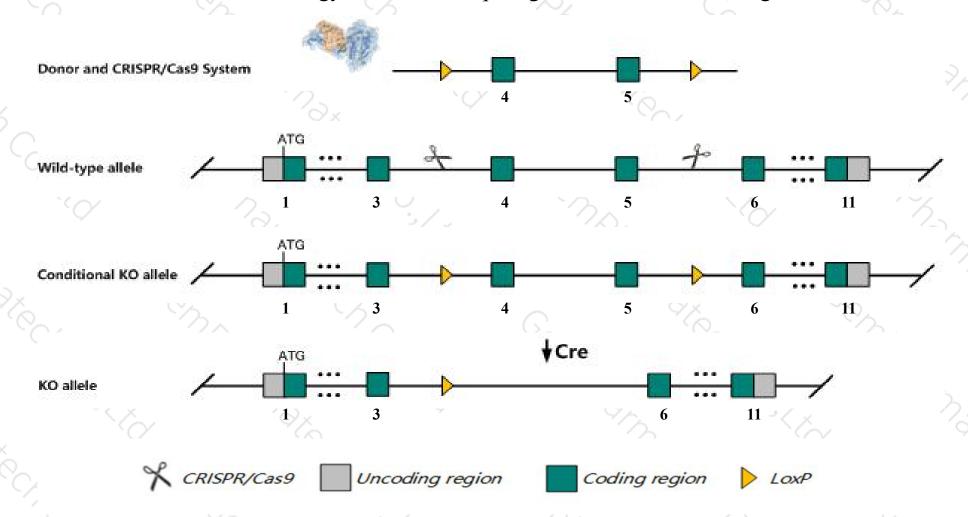
Strain background

C57BL/6JGpt

## Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Ubqln4* gene. The schematic diagram is as follows:



### Technical routes



- The *Ubqln4* gene has 5 transcripts. According to the structure of *Ubqln4* gene, exon4-exon5 of *Ubqln4-201* (ENSMUST00000008748.7) transcript is recommended as the knockout region. The region contains 422bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Ubqln4* 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 *Ubqln4* gene is located on the Chr3. 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.
- 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)



#### Ubqln4 ubiquilin 4 [Mus musculus (house mouse)]

Gene ID: 94232, updated on 31-Jan-2019

#### Summary

☆ ?

Official Symbol Ubqln4 provided by MGI

Official Full Name ubiquilin 4 provided by MGI

Primary source MGI:MGI:2150152

See related Ensembl:ENSMUSG00000008604

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 A1Up, A1u, Al663987, CIP75, UBIN

Expression Ubiquitous expression in adrenal adult (RPKM 35.7), ovary adult (RPKM 34.6) and 28 other tissuesSee more

Orthologs human all

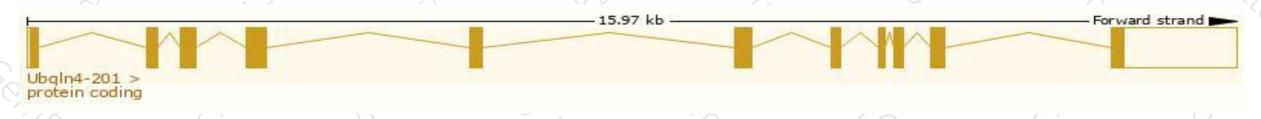
# Transcript information (Ensembl)



The gene has 5 transcripts, all transcripts are shown below:

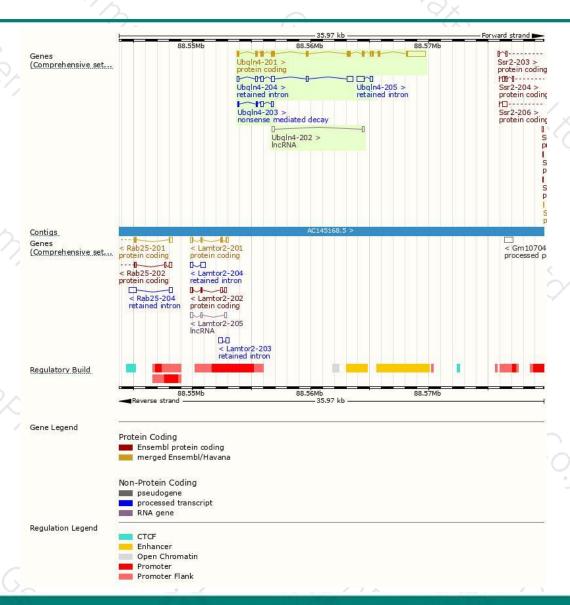
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ubqln4-201	ENSMUST00000008748.7	3330	596aa	Protein coding	CCDS17479	Q99NB8	TSL:1 GENCODE basic APPRIS P1
UbqIn4-203	ENSMUST00000192962.1	590	<u>45aa</u>	Nonsense mediated decay	571	A0A0A6YXU5	TSL:3
Ubqln4-204	ENSMUST00000194303.5	1379	No protein	Retained intron	(1 <u>4</u> )	ų.	TSL:1
Ubqln4-205	ENSMUST00000195498.1	871	No protein	Retained intron	120		TSL:3
Ubqln4-202	ENSMUST00000192687.1	362	No protein	IncRNA	1783	8	TSL:5

The strategy is based on the design of *Ubqln4-201* transcript, The transcription is shown below



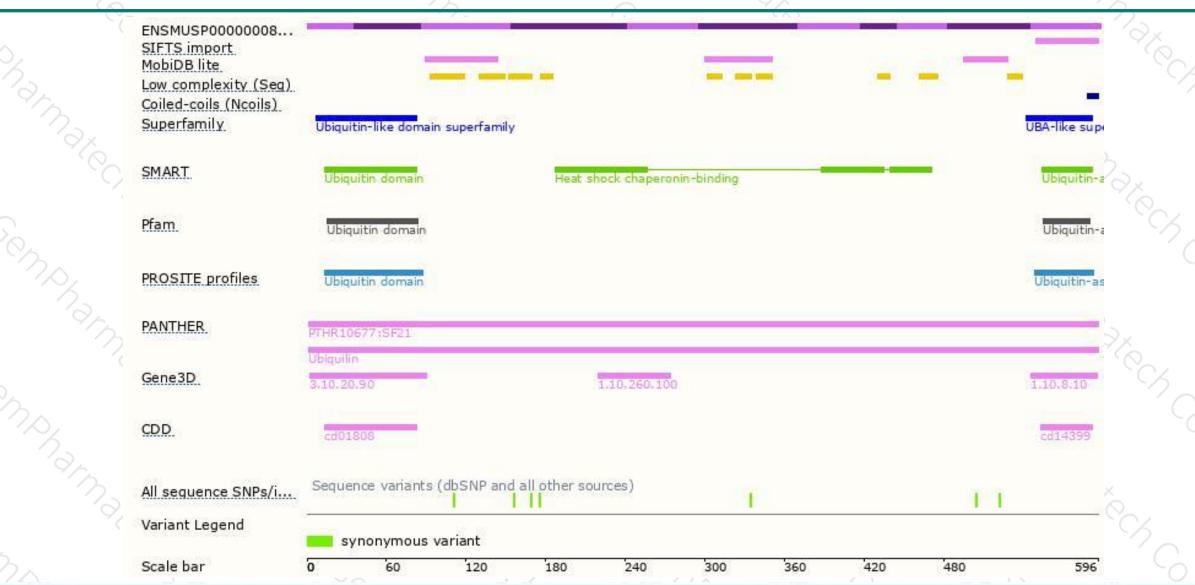
### Genomic location distribution





### Protein domain







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





