

# Kpna7 Cas9-CKO Strategy

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Reviewer: Xueting Zhang

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## **Project Overview**



**Project Name** 

Kpna7

**Project type** 

Cas9-CKO

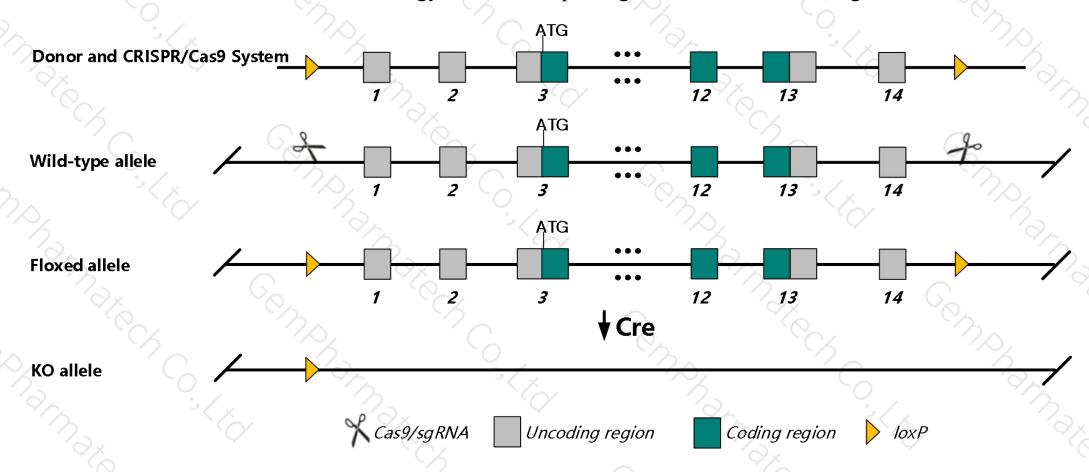
Strain background

C57BL/6JGpt

## Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Kpna7* gene. The schematic diagram is as follows:



### Technical routes



- The *Kpna7* gene has 6 transcripts. According to the structure of *Kpna7* gene, exon1-exon14 of *Kpna7-202* (ENSMUST00000110673.7) transcript is recommended as the knockout region. The region contains all of the coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Kpna7* 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**



- ➤ According to the existing MGI data, Mice homozygous or heterozygous for a null mutation display smaller litter sizes with preferential loss of females and accelerated cell cycles post fertilization resulting in loss of embryos.
- > The effect of transcripts 204,206 is unknown.
- > The *Kpna7* gene is located on the Chr5. 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)



#### Kpna7 karyopherin alpha 7 (importin alpha 8) [ Mus musculus (house mouse) ]

Gene ID: 381686, updated on 10-Oct-2019

#### Summary

☆ ?

Official Symbol Kpna7 provided by MGI

Official Full Name karyopherin alpha 7 (importin alpha 8) provided by MGI

Primary source MGI:MGI:2141165

See related Ensembl: ENSMUSG00000038770

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 Gm1055: AW146299

Expression Biased expression in ovary adult (RPKM 5.4) and testis adult (RPKM 2.0) See more

Orthologs human all

#### Genomic context

Location: 5; 5 G2

Exon count: 14

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See Kpna7 in Genome Data Viewer

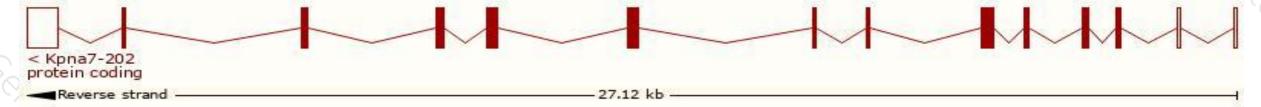
## Transcript information (Ensembl)



The gene has 6 transcripts, all transcripts are shown below:

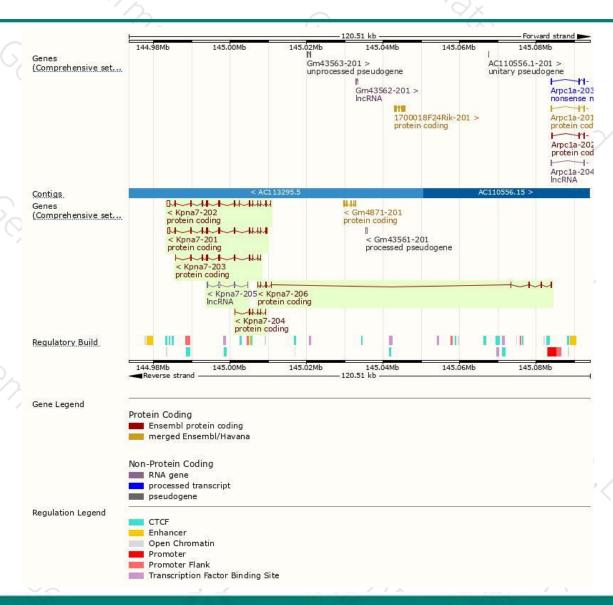
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Kpna7-202	ENSMUST00000110673.7	2453	520aa	Protein coding	CCDS84996	COLLJ0	TSL:5 GENCODE basic APPRIS ALT2
Kpna7-201	ENSMUST00000110672.7	2262	<u>499aa</u>	Protein coding	CCDS19852	COLLJ0	TSL:1 GENCODE basic APPRIS P3
Kpna7-203	ENSMUST00000116454.9	1500	<u>499aa</u>	Protein coding	CCDS19852	COLLJO	TSL:1 GENCODE basic APPRIS P3
Kpna7-206	ENSMUST00000151196.1	676	<u>38aa</u>	Protein coding	720	D3Z0I2	CDS 3' incomplete TSL:5
Kpna7-204	ENSMUST00000139024.7	598	<u>166aa</u>	Protein coding	127	D3Z2P7	CDS 3' incomplete TSL:5
Kpna7-205	ENSMUST00000142866.1	538	No protein	IncRNA		- 1	TSL:2

The strategy is based on the design of *Kpna7-202* transcript, The transcription is shown below



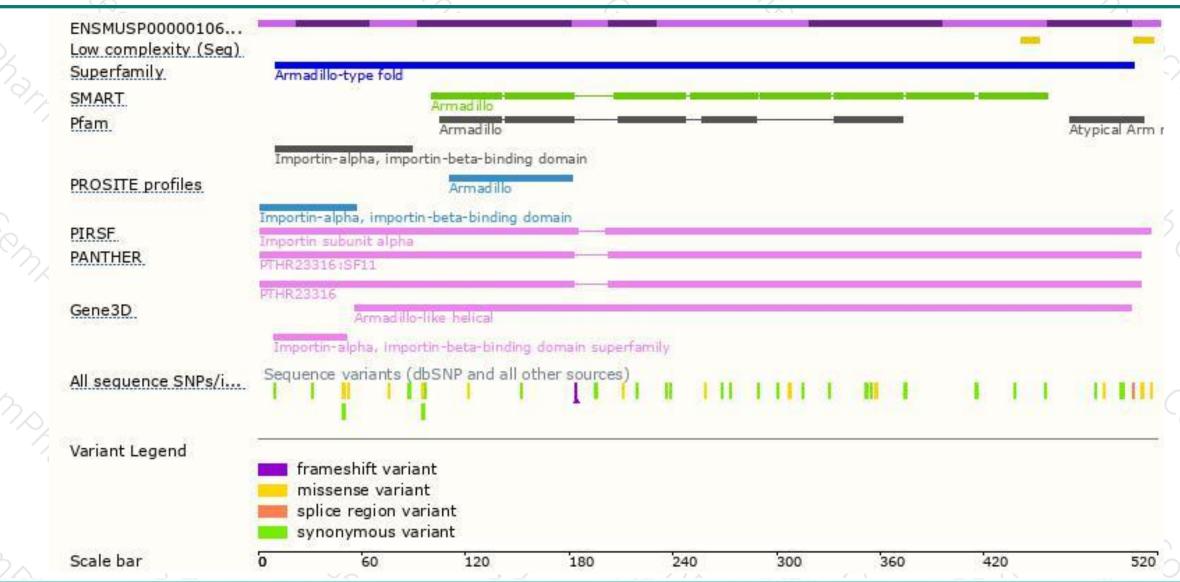
### Genomic location distribution





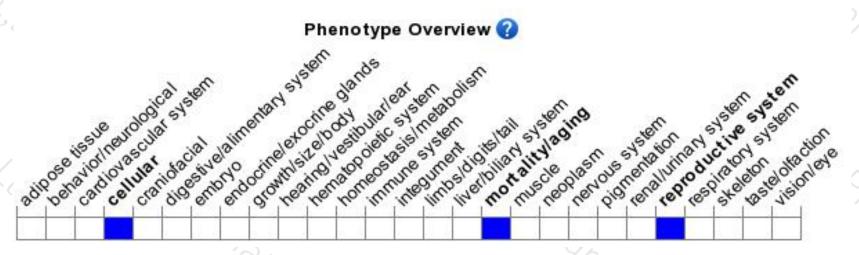
### Protein domain





## 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 or heterozygous for a null mutation display smaller litter sizes with preferential loss of females and accelerated cell cycles post fertilization resulting in loss of embryos.



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





