

# Dolar Day Co. Nudc Cas9-CKO Strategy To hall alto color color

Designer: Shilei Zhu

# **Project Overview**



**Project Name** 

Project type Cas9-CKO

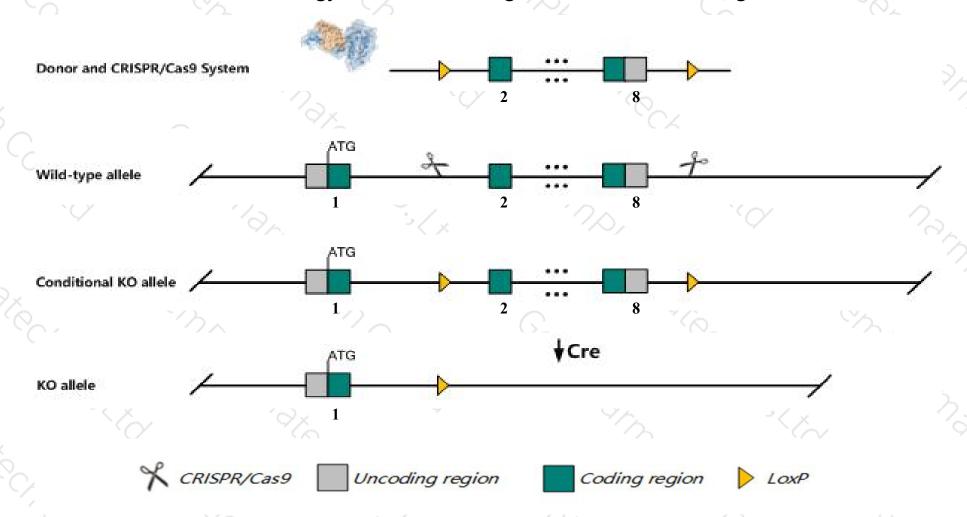
Strain background C57BL/6JGpt

Nudc

## Conditional Knockout strategy



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



## Technical routes



- The *Nudc* gene has 3 transcripts. According to the structure of *Nudc* gene, exon2-exon8 of *Nudc-201* (ENSMUST00000030665.6) transcript is recommended as the knockout region. The region contains most 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 *Nudc* 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 *Nudc* gene is located on the Chr4. 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)



#### Nudc nudC nuclear distribution protein [Mus musculus (house mouse)]

Gene ID: 18221, updated on 31-Jan-2019

#### Summary

☆ ?

Official Symbol Nudc provided by MGI

Official Full Name nudC nuclear distribution protein provided by MGI

Primary source MGI:MGI:106014

See related Ensembl: ENSMUSG00000028851

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 SIG-92, Silg92

Expression Ubiquitous expression in CNS E11.5 (RPKM 117.1), liver E14 (RPKM 77.3) and 26 other tissuesSee more

Orthologs human all

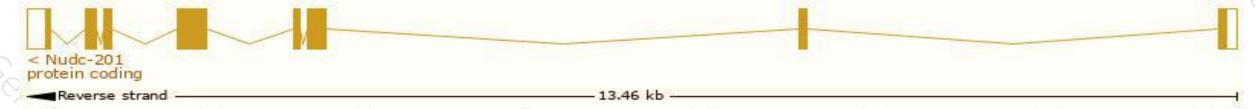
# Transcript information (Ensembl)



The gene has 3 transcripts, all transcripts are shown below:

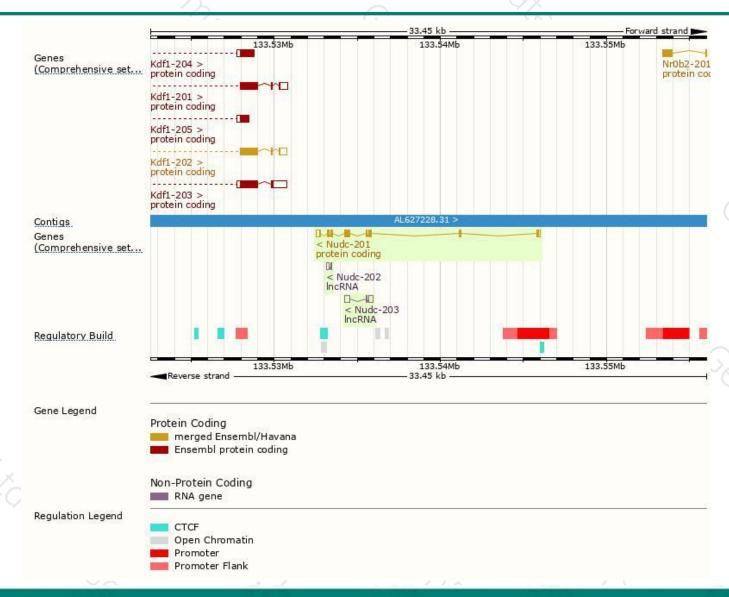
Name	Transcript ID	bp	Protein	Biotype	ccds	UniProt	Flags
Nudc-201	ENSMUST00000030665.6	1316	332aa	Protein coding	CCDS18750	<u>O35685</u>	TSL:1 GENCODE basic APPRIS P1
Nudc-203	ENSMUST00000148612.1	641	No protein	Processed transcript	<del>-</del> 8		TSL:2
Nudc-202	ENSMUST00000141860.1	204	No protein	Processed transcript	20	2	TSL:2

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



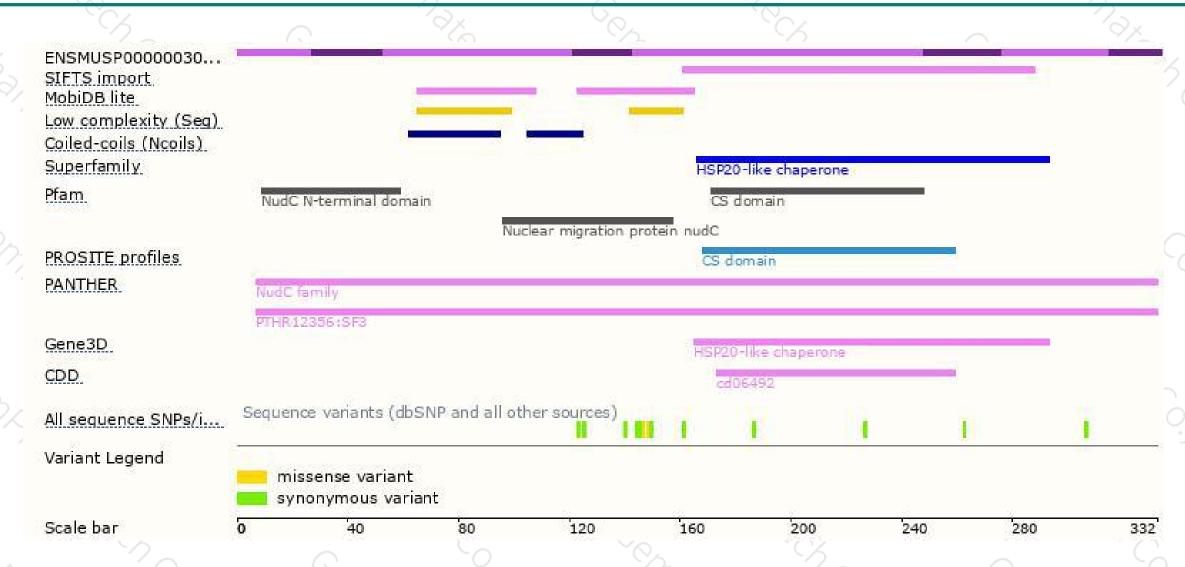
## Genomic location distribution





## Protein domain







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





