

# Nfic Cas9-CKO Strategy

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**Design Date:** 

2019-9-16

# **Project Overview**



Project Name Nfic

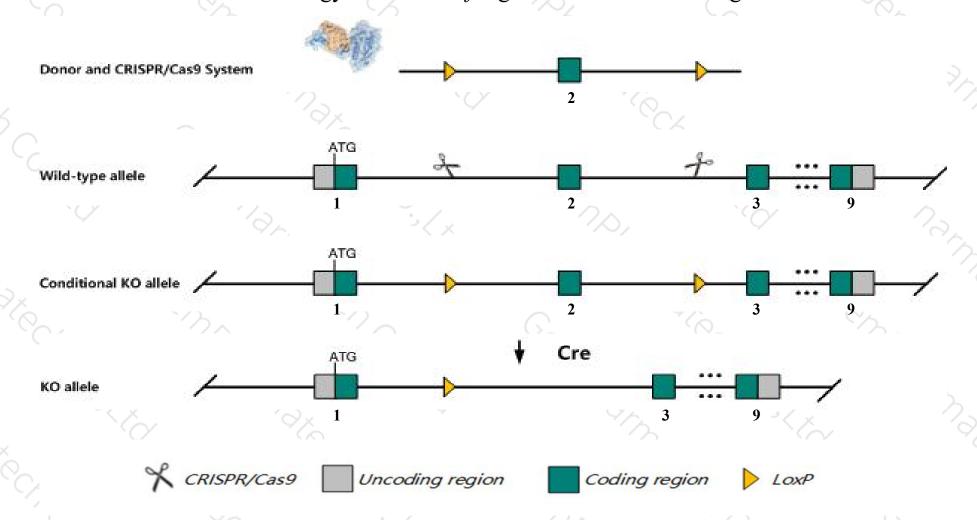
Project type Cas9-CKO

Strain background C57BL/6JGpt

## Conditional Knockout strategy



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



### Technical routes



- The *Nfic* gene has 7 transcripts. According to the structure of *Nfic* gene, exon2 of *Nfic-203*(ENSMUST00000105321.9) transcript is recommended as the knockout region. The region contains 532bp coding sequence.

  Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Nfic* 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 for a targeted null allele have abnormal incisor and molar root development, show reduced alveolar bone formation, and exhibit impaired feeding leading to severe runting and premature death when reared on standard laboratory chow.
- > The *Nfic* gene is located on the Chr10. 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)



#### Nfic nuclear factor I/C [Mus musculus (house mouse)]

Gene ID: 18029, updated on 31-Jan-2019

#### Summary

☆ ?

Official Symbol Nfic provided by MGI

Official Full Name nuclear factor I/C provided by MGI

Primary source MGI:MGI:109591

See related Ensembl:ENSMUSG00000055053

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 1110019L22Rik, 1500041O16Rik, AA589446, AI746521, NF1-C

Expression Broad expression in adrenal adult (RPKM 27.2), lung adult (RPKM 13.1) and 19 other tissuesSee more

Orthologs <u>human</u> all

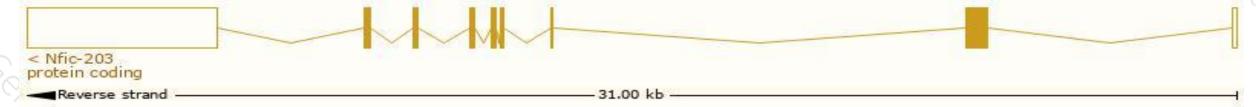
# Transcript information (Ensembl)



The gene has 7 transcripts, all transcripts are shown below:

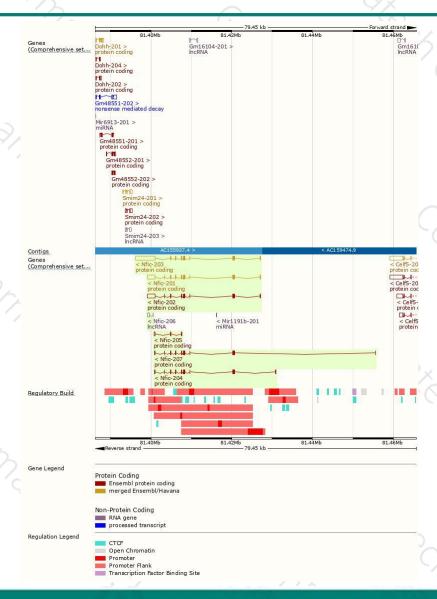
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Nfic-203	ENSMUST00000105321.9	6237	428aa	Protein coding	CCDS35999	P70255 Q3UHA6	TSL:1 GENCODE basic APPRIS ALT1
Nfic-201	ENSMUST00000020461.14	3348	<u>439aa</u>	Protein coding	CCDS24058	P70255	TSL:1 GENCODE basic APPRIS P3
Nfic-202	ENSMUST00000078185.13	3163	397aa	Protein coding	ų.	P70255	TSL:5 GENCODE basic
Nfic-204	ENSMUST00000117966.1	1655	430aa	Protein coding	2	P70255	TSL:1 GENCODE basic
Nfic-207	ENSMUST00000221817.1	1531	<u>461aa</u>	Protein coding	ā	A0A1Y7VK55	TSL:5 GENCODE basic
Nfic-205	ENSMUST00000140916.7	575	<u>96aa</u>	Protein coding		F6XYU7	CDS 5' incomplete TSL:3
Nfic-206	ENSMUST00000152670.1	476	No protein	IncRNA	ų.	V4-0	TSL:1

The strategy is based on the design of *Nfic-203* transcript, The transcription is shown below



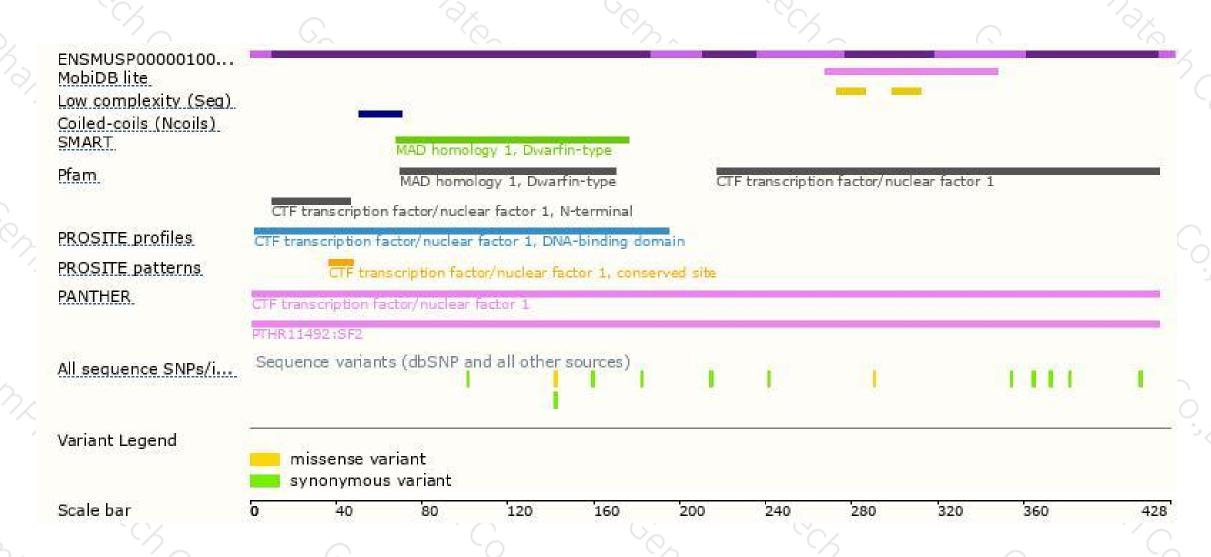
### Genomic location distribution





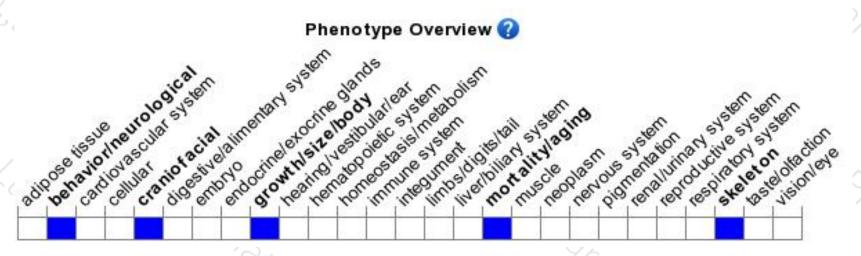
### Protein domain





# 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 targeted null allele have abnormal incisor and molar root development, show reduced alveolar bone formation, and exhibit impaired feeding leading to severe runting and premature death when reared on standard laboratory chow.



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





