

Nfe2l2 Cas9-KO Strategy To hall alto color color

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Project Overview



Project Name

Project type

Strain background C57BL/6J

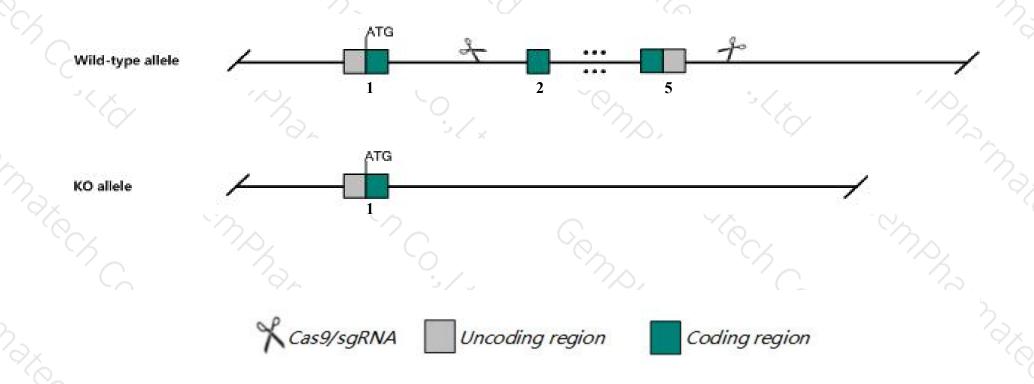
Nfe2l2

Cas9-KO

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Nfe2l2 gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Nfe2l2* gene has 2 transcripts. According to the structure of *Nfe2l2* gene, exon2-exon5 of *Nfe2l2-201* (ENSMUST00000102672.4) 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 *Nfe2l2* gene. The brief process is as follows: sgRNA was transcribed in vitro.Cas9 and sgRNA were microinjected into the fertilized eggs of C57BL/6J 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/6J mice.

Notice



- ➤ According to the existing MGI data, Mice homozygous for a knock-out allele exhibit increased sensitivity to oxidative stress in a variety of organs and cells including brain, liver, erythrocytes, and spleen, abnormal tooth enamel, and abnormal response to various injuries, chemical treatments, and induced inflammatory diseases.
- ➤ The *Nfe2l2* gene is located on the Chr2. 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)



Nfe2l2 nuclear factor, erythroid derived 2, like 2 [Mus musculus (house mouse)]

Gene ID: 18024, updated on 9-Apr-2019

Summary

↑ ?

Official Symbol Nfe2l2 provided by MGI

Official Full Name nuclear factor, erythroid derived 2, like 2 provided by MGI

Primary source MGI:MGI:108420

See related Ensembl:ENSMUSG00000015839

Gene type protein coding
RefSeq status REVIEWED
Organism Mus musculus

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;

Muroidea; Muridae; Murinae; Mus; Mus

Also known as Nrf2

Summary This gene encodes a transcription factor which is a member of a small family of basic leucine zipper (bZIP) proteins. The encoded

transcription factor regulates genes which contain antioxidant response elements (ARE) in their promoters; many of these genes encode proteins involved in response to injury and inflammation which includes the production of free radicals. [provided by RefSeq, Sep 2015]

Expression Ubiquitous expression in bladder adult (RPKM 57.1), duodenum adult (RPKM 26.8) and 26 other tissuesSee more

Orthologs <u>human all</u>

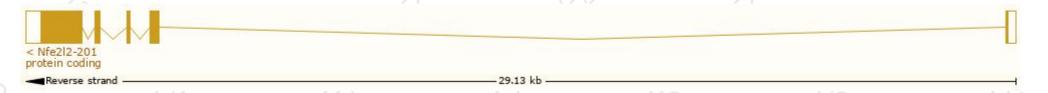
Transcript information (Ensembl)



The gene has 2 transcripts, all transcripts are shown below:

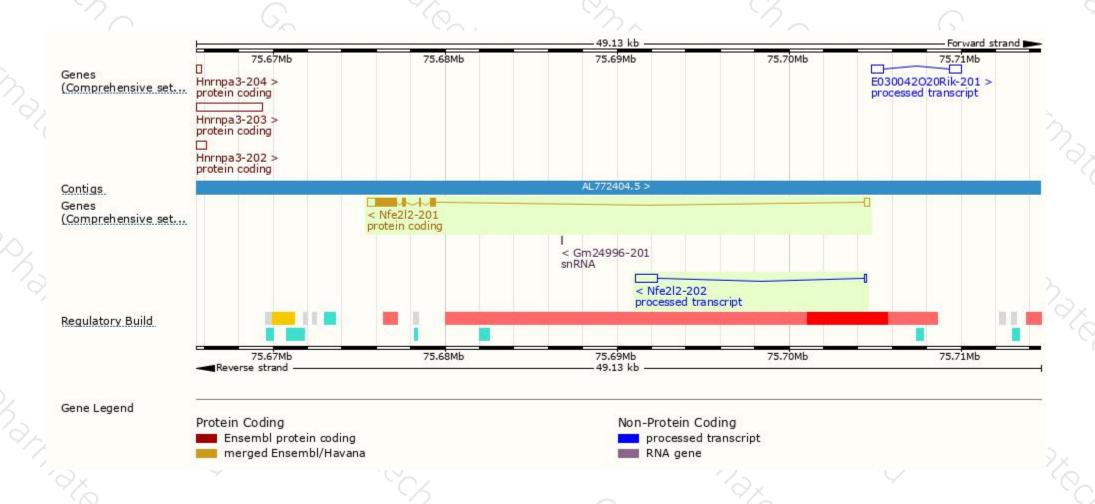
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Nfe2l2-201	ENSMUST00000102672.4	2475	<u>597aa</u>	Protein coding	CCDS16150	Q60795	TSL:1 GENCODE basic APPRIS P1
Nfe2l2-202	ENSMUST00000152371.1	1447	No protein	Processed transcript	·		TSL:1

The strategy is based on the design of Nfe2l2-201 transcript, The transcription is shown below



Genomic location distribution





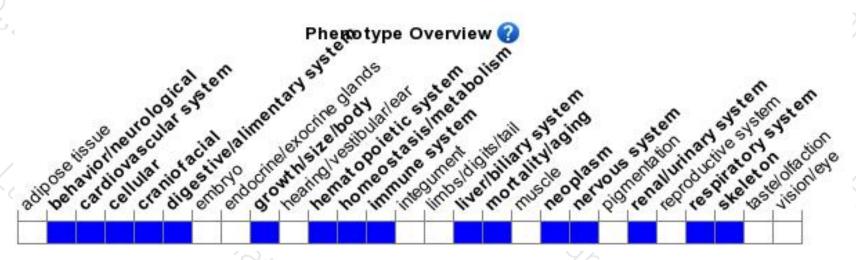
Protein domain





Mouse phenotype description(MGI)





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

According to the existing MGI data, Mice homozygous for a knock-out allele exhibit increased sensitivity to oxidative stress in a variety of organs and cells including brain, liver, erythrocytes, and spleen, abnormal tooth enamel, and abnormal response to various injuries, chemical treatments, and induced inflammatory diseases.



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

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