

Krt24 Cas9-KO Strategy

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Project Overview



Project Name

Krt24

Project type

Cas9-KO

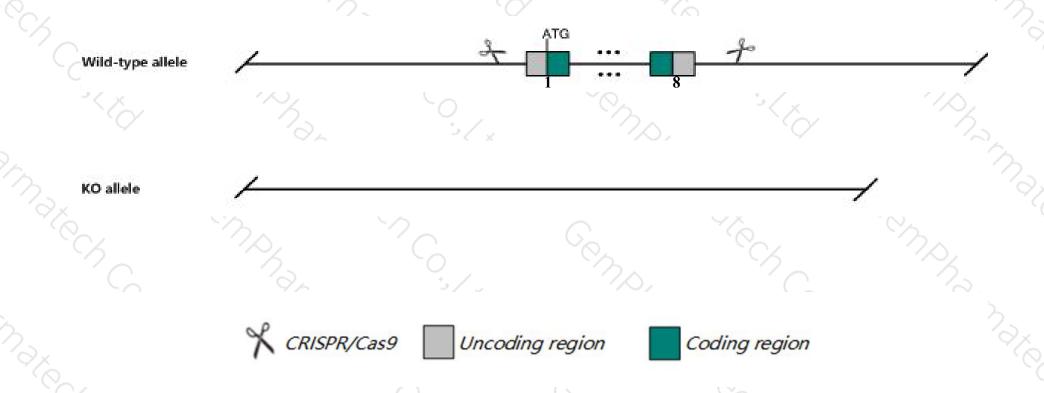
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Krt24* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Krt24* gene has 1 transcript. According to the structure of *Krt24* gene, exon1-exon8 of *Krt24-201* (ENSMUST00000017255.3) 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 *Krt24* gene. The brief process is as follows: CRISPR/Cas9 system

Notice



- ➤ The *Krt24* gene is located on the Chr11. 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)



Krt24 keratin 24 [Mus musculus (house mouse)]

Gene ID: 75706, updated on 12-Aug-2019

Summary

☆ ?

Official Symbol Krt24 provided by MGI

Official Full Name keratin 24 provided by MGI

Primary source MGI:MGI:1922956

See related Ensembl: ENSMUSG00000020913

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 K24; CK-24; 2310058N18Rik; 2310075C18Rik

Expression Low expression observed in reference dataset See more

Orthologs human all

Genomic context

☆ ?

Location: 11; 11 D

See Krt24 in Genome Data Viewer

Exon count: 8

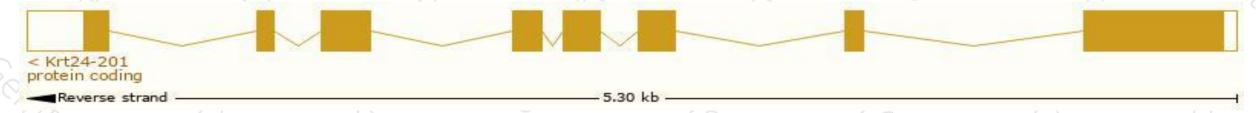
Transcript information (Ensembl)



The gene has 1 transcript, and the transcript is shown below:

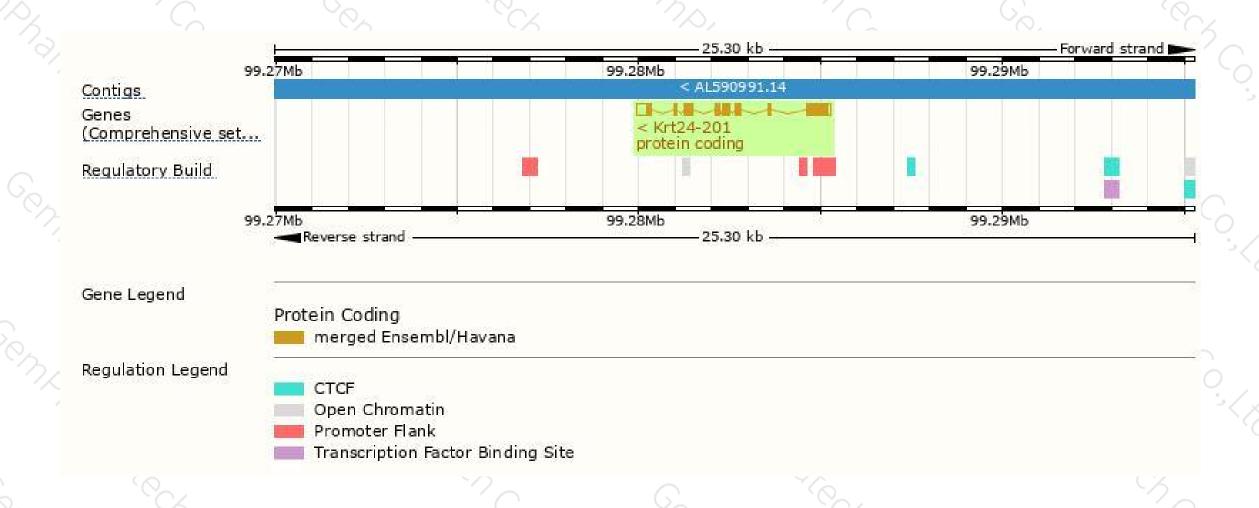
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags	l
Krt24-201	ENSMUST00000017255.3	1847	512aa	Protein coding	CCDS36305	A1L317	TSL:1 GENCODE basic APPRIS P1	Ľ

The strategy is based on the design of *Krt24-201* transcript, The transcription is shown below



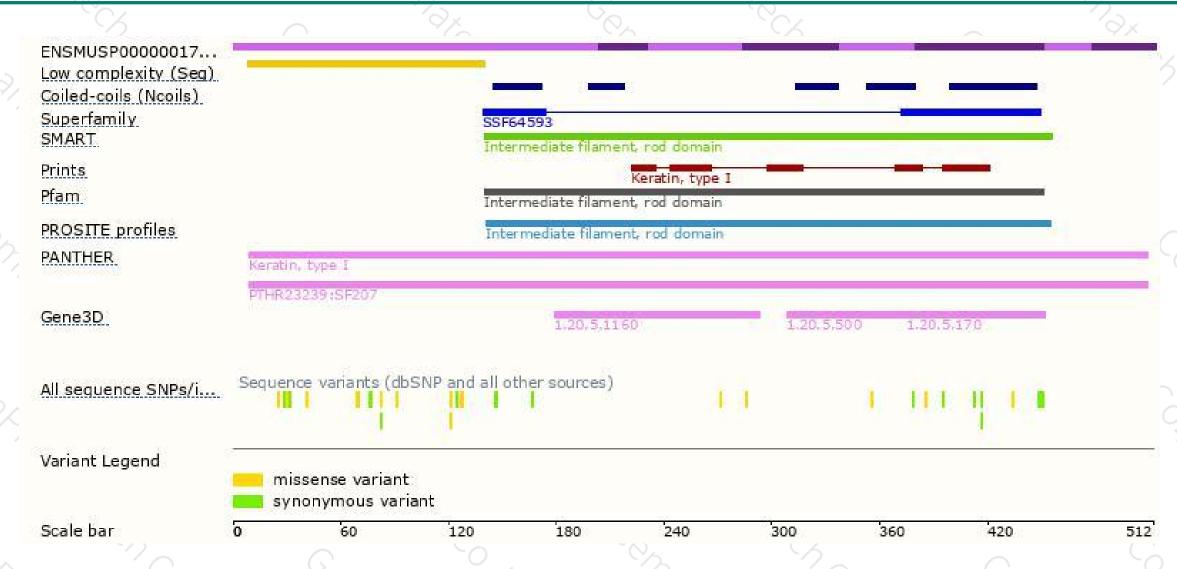
Genomic location distribution





Protein domain







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





