

Krtap27-1 Cas9-KO Strategy

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Project Overview



Project Name

Krtap27-1

Project type

Cas9-KO

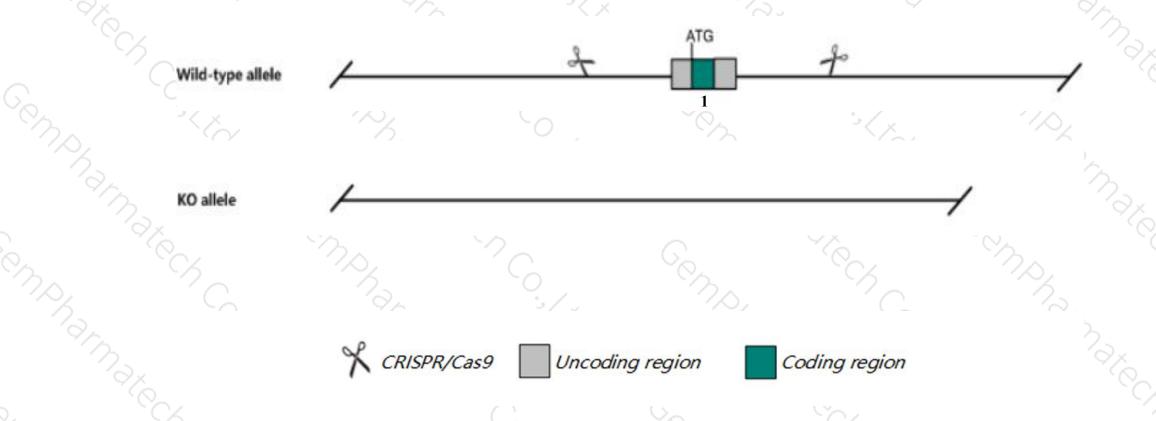
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Krtap27-1* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Krtap27-1* gene has 1 transcript. According to the structure of *Krtap27-1* gene, exon1 of *Krtap27-1-201*(ENSMUST00000165580.2) 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 *Krtap27-1* gene. The brief process is as follows: CRISPR/Cas9 system 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.

Notice



- > The *Krtap27-1* gene is located on the Chr16. 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)



Krtap27-1 keratin associated protein 27-1 [Mus musculus (house mouse)]

Gene ID: 239933, updated on 13-Mar-2020





Official Symbol Krtap27-1 provided by MGI

Official Full Name keratin associated protein 27-1 provided by MGI

Primary source MGI:MGI:3646229

See related Ensembl: ENSMUSG00000090515

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 EG239933

Orthologs <u>human all</u>

Transcript information (Ensembl)



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

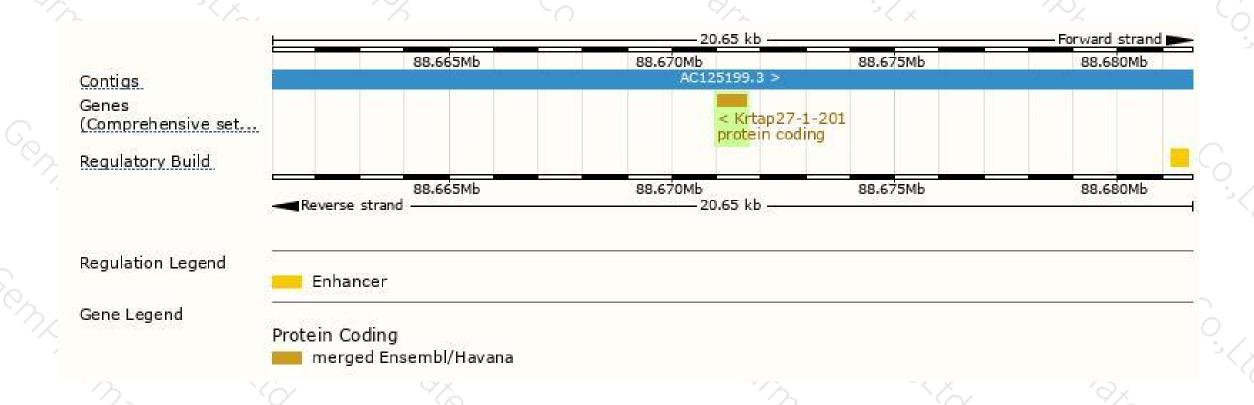
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags	
Krtap27-1-201	ENSMUST00000165580.2	646	202aa	Protein coding	CCDS49892	E9PX37	TSL:NA GENCODE basic APPRIS P1	Ľ

The strategy is based on the design of *Krtap27-1-201* transcript, the transcription is shown below:



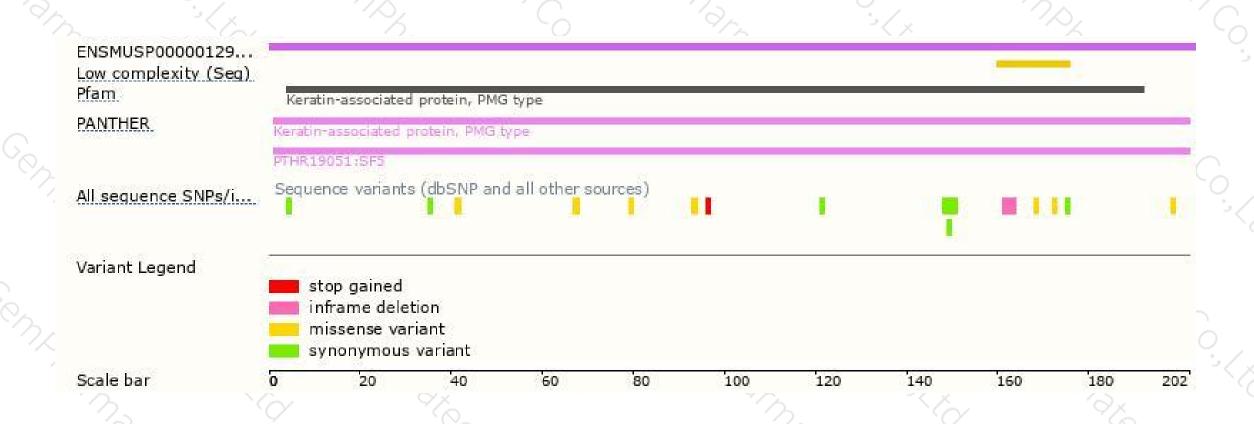
Genomic location distribution





Protein domain







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





