

Taar5 Cas9-CKO Strategy

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Project Overview



Project Name

Taar5

Project type

Cas9-CKO

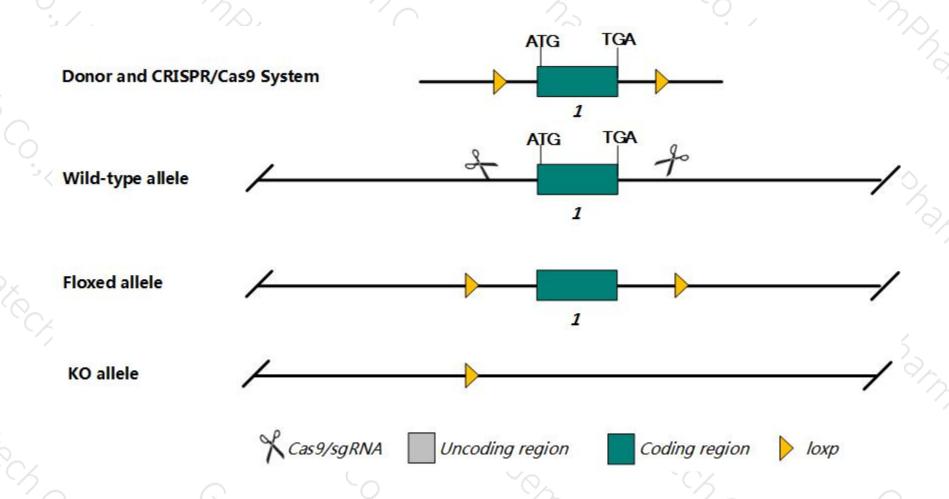
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Taar5* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Taar5* gene has 1 transcript. According to the structure of *Taar5* gene, exon1 of *Taar5-201*(ENSMUST00000092659.3) transcript is recommended as the knockout region. The region contains all coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Taar5* gene. The brief process is as follows:gRNA was transcribed in vitro, donor was constructed.Cas9, gRNA 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 knock-out allele exhibit abnormal olfactory system physiology.

- The *Taar5* 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)



Taar5 trace amine-associated receptor 5 [Mus musculus (house mouse)]

Gene ID: 215854, updated on 12-Aug-2019

Summary

Official Symbol Taar5 provided by MGI

Official Full Name trace amine-associated receptor 5 provided by MGI

Primary source MGI:MGI:2685073

See related Ensembl: ENSMUSG00000069706

RefSeq status PROVISIONAL
Organism Mus musculus

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

Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus

Also known as Gm227; taR-5 Orthologs <u>human</u> all

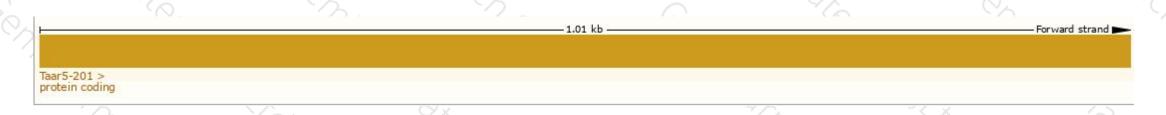
Transcript information (Ensembl)



The gene has 1 transcript, all transcripts are shown below:

Name 🍦	Transcript ID A	bp 🍦	Protein	Biotype	CCDS	UniProt	Flags
Taar5-201	ENSMUST00000092659.3	1014	<u>337aa</u>	Protein coding	CCDS23738₽	B2RTA0 ₽ Q5QD14 ₽	TSL:NA GENCODE basic APPRIS

The strategy is based on the design of *Taar5-201* transcript, The transcription is shown below



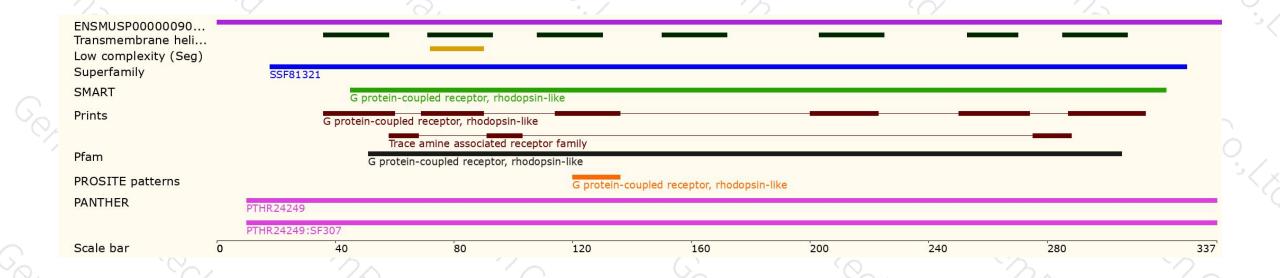
Genomic location distribution





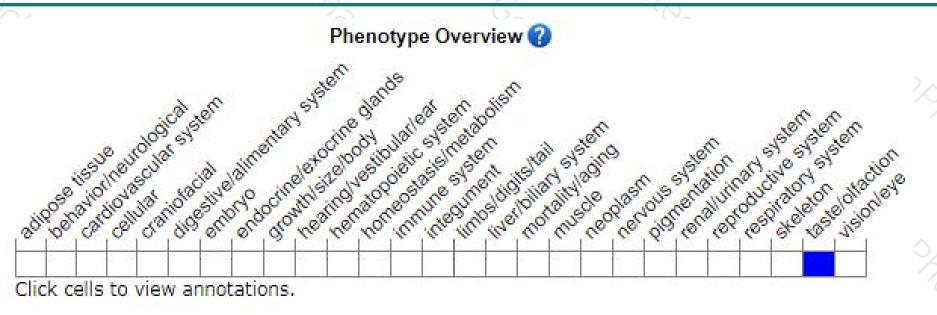
Protein domain





Mouse phenotype description(MGI)





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

Mice homozygous for a knock-out allele exhibit abnormal olfactory system physiology.



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





