

Amhr2 Cas9-CKO Strategy

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Overview

Target Gene Name

• *Amhr2*

Project Type

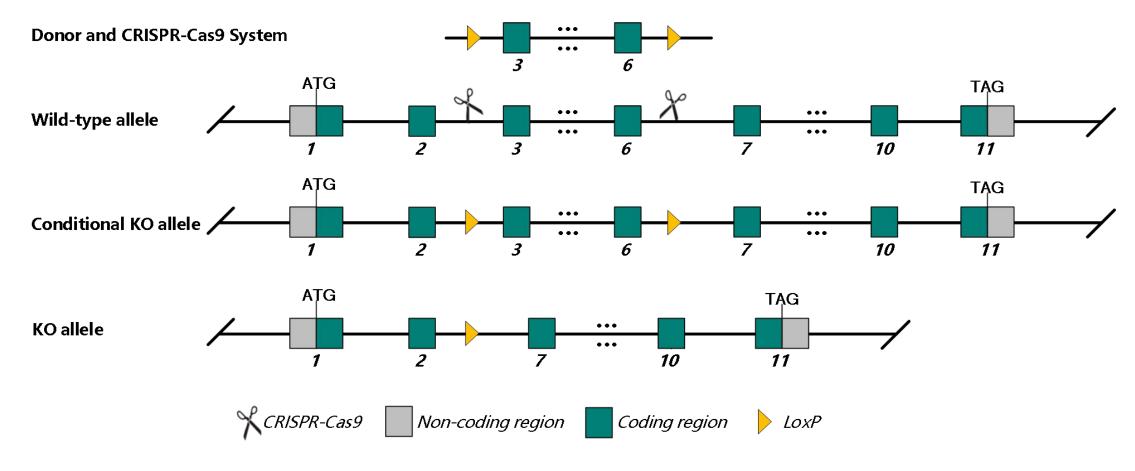
• Cas9-CKO

Genetic Background

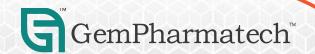
• C57BL/6JGpt



Strain Strategy



Schematic representation of CRISPR-Cas9 engineering used to edit the Amhr2 gene.



Technical Information

- The *Amhr2* gene has 5 transcripts. According to the structure of *Amhr2* gene, exon 3-6 of *Amhr2*-201 (ENSMUST00000023809.11) is recommended as the knockout region. The region contains 608 bp of coding sequence. Knockout the region will result in disruption of gene function.
- In this project we use CRISPR-Cas9 technology to modify *Amhr2* 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 on-target amplicon sequencing. A stable F1-generation mouse strain was obtained by mating positive F0-generation mice with C57BL/6JGpt mice and confirmation of the desired mutant allele was carried out by PCR and on-target amplicon sequencing.
- 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.

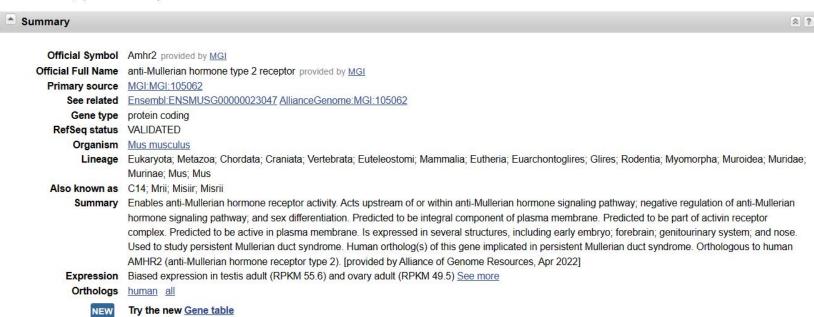


Gene Information

Amhr2 anti-Mullerian hormone type 2 receptor [Mus musculus (house mouse)]

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Gene ID: 110542, updated on 31-May-2023



Genomic context

Location: 15 F3; 15 57.58 cM

Try the new Transcript table

See Amhr2 in Genome Data Viewer

Exon count: 12

https://www.ncbi.nlm.nih.gov/gene/110542

☆ ?

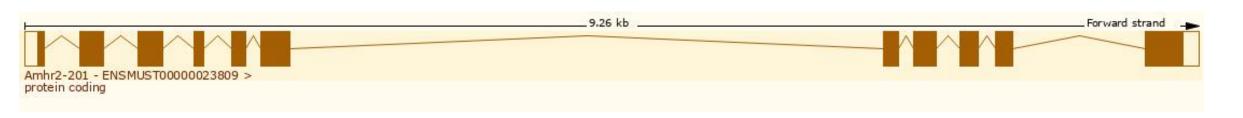


Transcript Information

The gene has 5 transcripts, all transcripts are shown below:

Show/hide columns (1 hidden)							Filter
Transcript ID	Name A	bp 🛊	Protein	Biotype	CCDS	UniProt Match	Flags
ENSMUST00000023809.11	Amhr2-201	1938	<u>568aa</u>	Protein coding	CCDS27882 ₺	Q8K592₽	Ensembl Canonical GENCODE basic APPRIS P1 TSL:1
ENSMUST00000161376.2	Amhr2-202	364	No protein	Retained intron		6#8	TSL:2
ENSMUST00000162893.2	Amhr2-203	611	<u>190aa</u>	Protein coding		F6V9X7配	TSL:5 CDS 5' incomplete
ENSMUST00000229278.2	Amhr2-204	1735	<u>448aa</u>	Protein coding		A0A2R8W6H9 @	GENCODE basic
ENSMUST00000229566.2	Amhr2-205	209	<u>61aa</u>	Protein coding		A0A2R8VHE2₺	CDS 3' incomplete

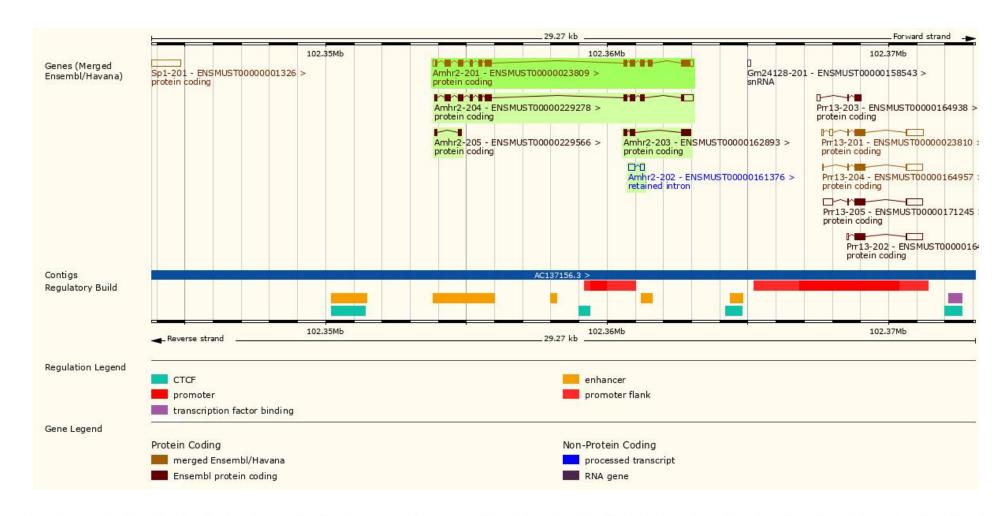
The strategy is based on the design of *Amhr2*-201 transcript, the transcription is shown below:



Source: http://asia.ensembl.org/

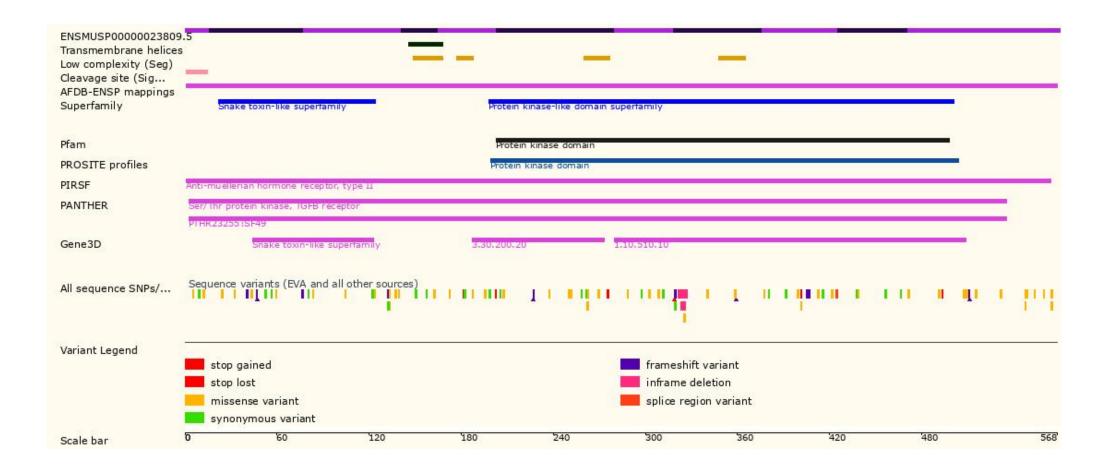


Genomic Information





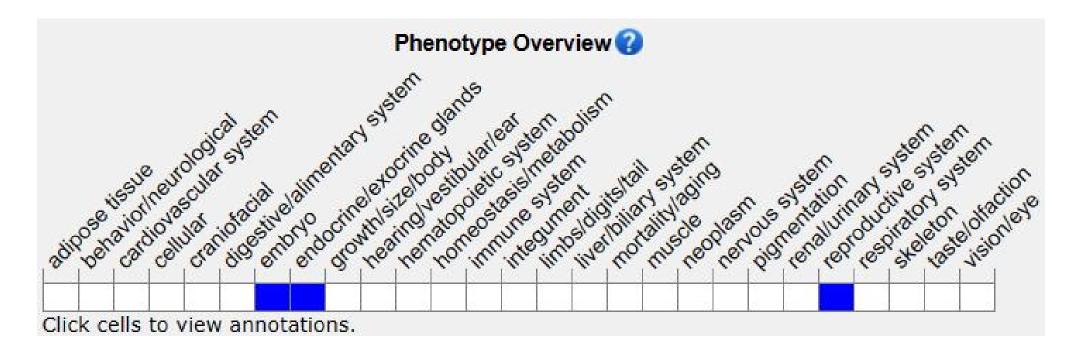
Protein Information





Source: : https://www.ensembl.org

Mouse Phenotype Information (MGI)



Homozygous null mutant males have a complete male reproductive tract, but also a uterus and oviducts. Functional sperm are produced, but most males are infertile because female reproductive organs block sperm transfer.



Source: https://www.informatics.jax.org

Important Information

- According to the existing MGI data, homozygous null mutant males have a complete male reproductive tract, but also a uterus and oviducts. Functional sperm are produced, but most males are infertile because female reproductive organs block sperm transfer.
- The intron 2-3 is 276 bp, the loxp insertion may affect the regulation of this gene.
- This stratergy may not affect Amhr2-202 and Amhr2-203 transcript.
- *Amhr2* is located on Chr 15. If the knockout mice are crossed with other mouse strains to obtain double homozygous mutant offspring, please avoid the situation that the second gene is 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 the existing technology level.

