# *Amfr* Cas9-CKO Strategy

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Design Date: 2019-11-20
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# Project Overview

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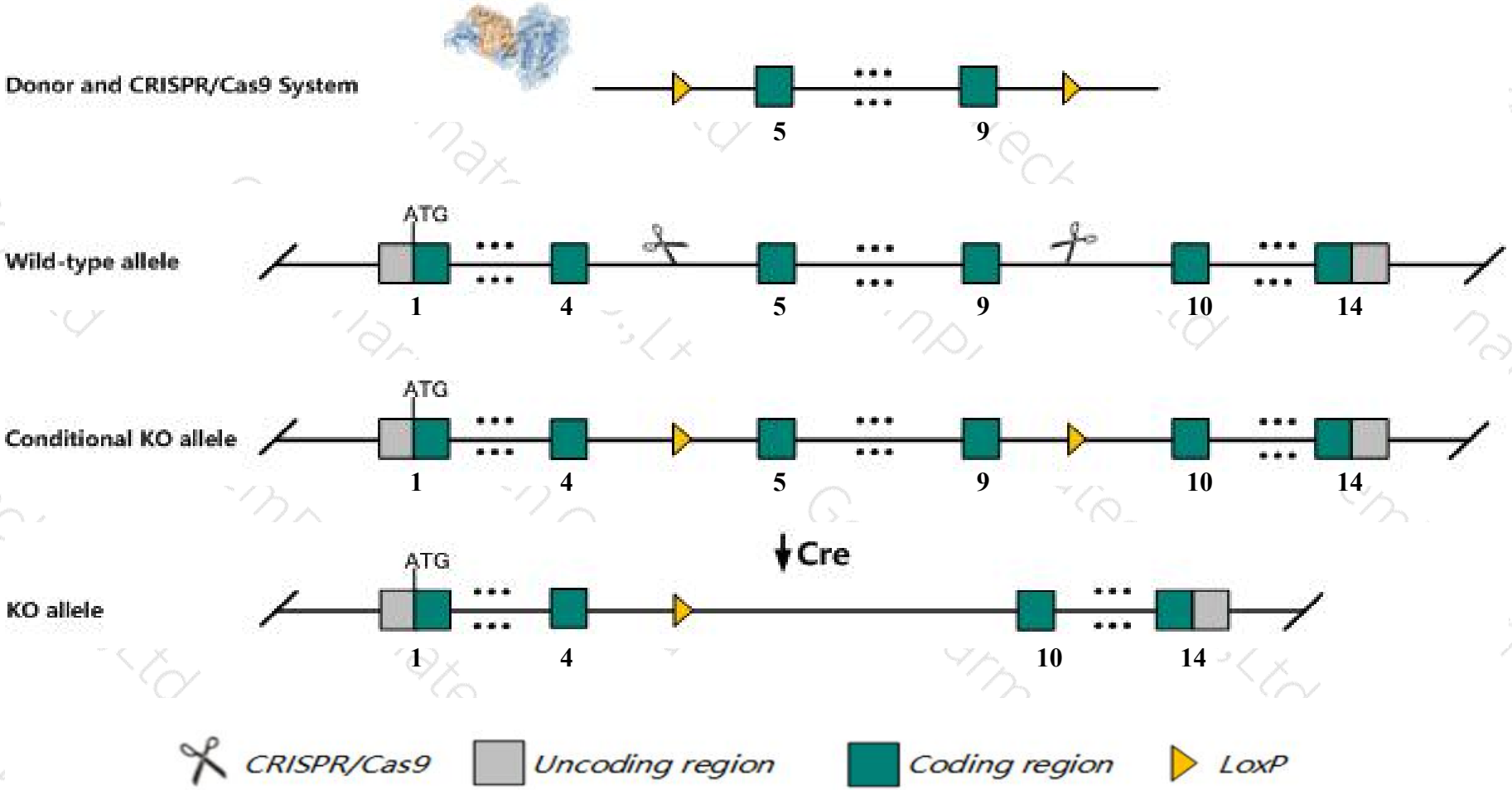
| **Project Name** | ***Amfr*** |
| **Project type** | **Cas9-CKO** |
| **Strain background** | **C57BL/6JGpt** |

# Conditional Knockout strategy

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



# Technical routes

- The *Amfr* gene has 4 transcripts. According to the structure of *Amfr* gene, exon5-exon9 of *Amfr-201* (ENSMUST00000053766.13) transcript is recommended as the knockout region.The region contains 610bp coding sequence. Knock out the region will result in disruption of protein function.

- In this project we use CRISPR/Cas9 technology to modify *Amfr* 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 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 for a gene-trapped null allele are obese and develop liver steatosis and/or hepatic inflammation resembling nonalcoholic steatohepatitis. Some mice develop liver tumors. Mice homozygous for another knock-out allele exhibit normal HMGCR turnover in mouse embryonic fibroblasts.

- The *Amfr* gene is located on the Chr8. 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）

集萃药康 GemPharmatech

## Amfr autocrine motility factor receptor [ *Mus musculus* (house mouse) ]

Gene ID: 23802, updated on 10-Oct-2019

### Summary

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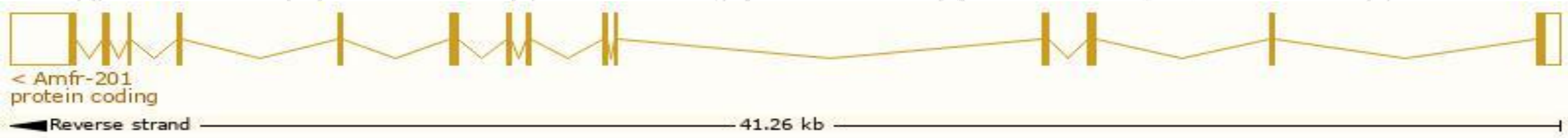
|---|---|
| **Official Symbol** | Amfr provided by MGI |
| **Official Full Name** | autocrine motility factor receptor provided by MGI |
| **Primary source** | MGI:MGI:1345634 |
| **See related** | Ensembl:ENSMUSG00000031751 |
| **Gene type** | protein coding |
| **RefSeq status** | VALIDATED |
| **Organism** | Mus musculus |
| **Lineage** | Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus |
| **Also known as** | gp78 |
| **Expression** | Ubiquitous expression in adrenal adult (RPKM 95.7), liver adult (RPKM 63.4) and 28 other tissues See more |
| **Orthologs** | human all |

# Transcript information（Ensembl）

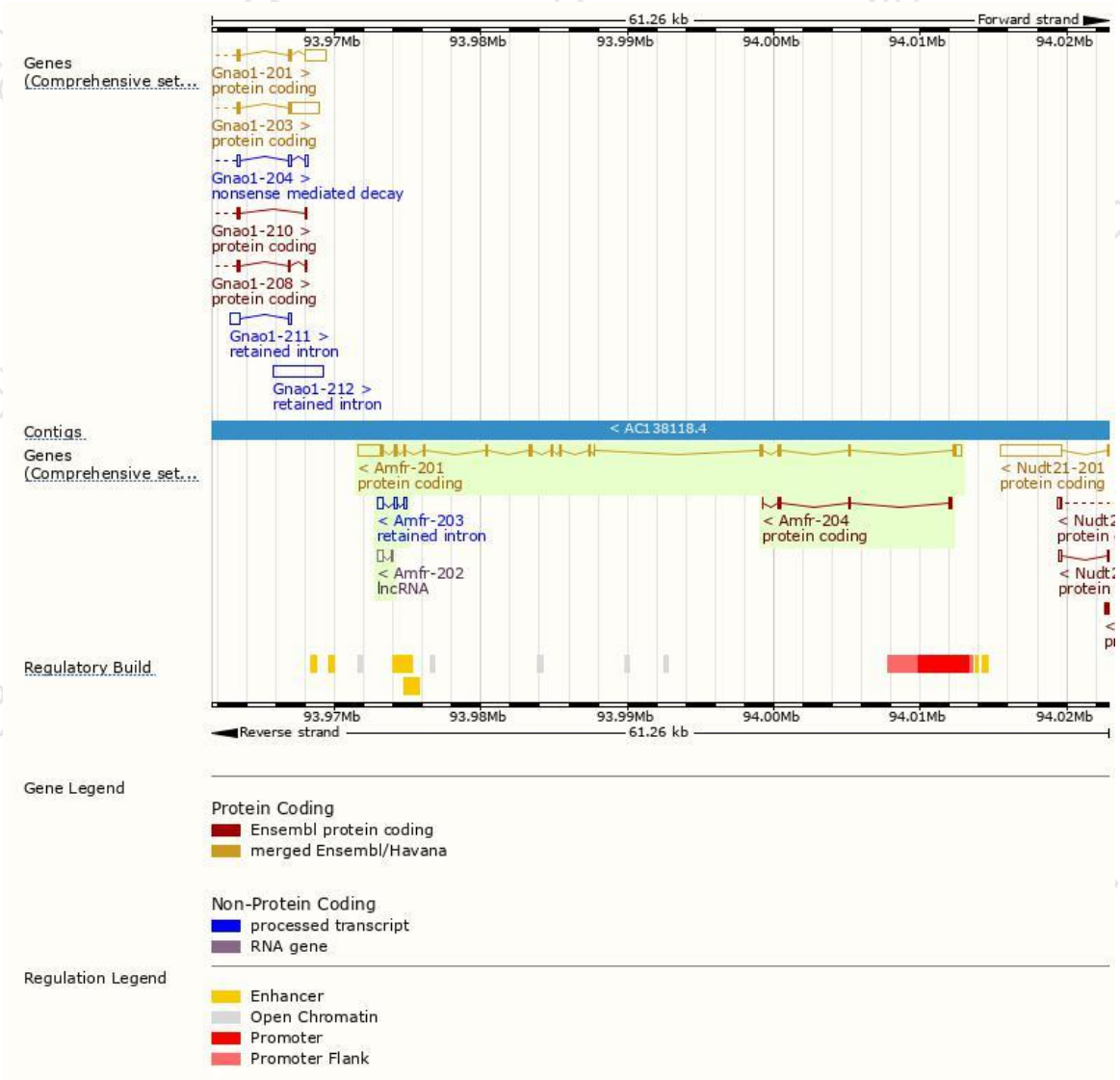
The gene has 4 transcripts,all transcripts are shown below:

| Name | Transcript ID | bp | Protein | Biotype | CCDS | UniProt | Flags |
|---|---|---|---|---|---|---|---|
| Amfr-201 | ENSMUST00000053766.13 | 3880 | 639aa | Protein coding | CCDS22533 | Q3TCI2 Q9R049 | TSL:1 GENCODE basic APPRIS P1 |
| Amfr-204 | ENSMUST00000143265.1 | 414 | 105aa | Protein coding | - | H3BJC0 | CDS 3' incomplete TSL:5 |
| Amfr-203 | ENSMUST00000139702.1 | 779 | No protein | Retained intron | - | - | TSL:2 |
| Amfr-202 | ENSMUST00000137475.1 | 392 | No protein | lncRNA | - | - | TSL:5 |

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

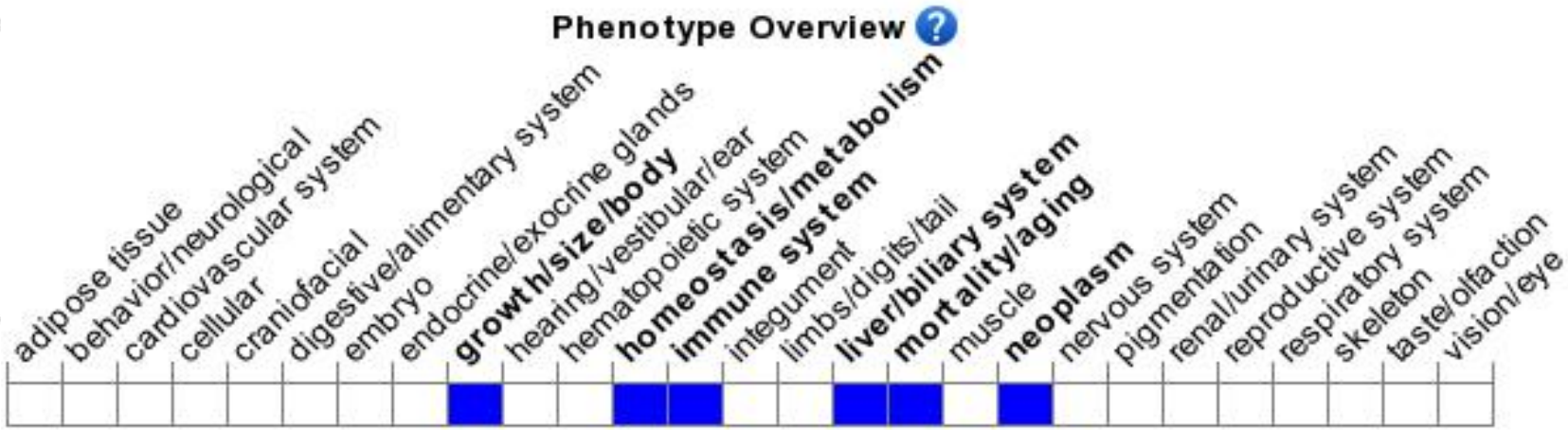


# Genomic location distribution



# 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/).*

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

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