

Xpr1 Cas9-KO Strategy

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Project Overview

Project Name

Xpr1

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Xpr1* gene. The schematic diagram is as follows:



- The *Xpr1* gene has 7 transcripts. According to the structure of *Xpr1* gene, exon2 of *Xpr1-201* (ENSMUST00000027741.11) transcript is recommended as the knockout region. The region contains 52bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Xpr1* gene. The brief process is as follows: CRISPR/Cas9 system v

- According to the existing MGI data, Homozygotes and heterozygotes for a variant from some wild Mus stocks, including *M. spretus*, support replication of xenotropic murine leukemia viruses and mink cell focus-forming murine leukemia viruses that are not replicated in most laboratory strains.
- The *Xpr1* gene is located on the Chr1. 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)

Xpr1 xenotropic and polytropic retrovirus receptor 1 [Mus musculus (house mouse)]

Gene ID: 19775, updated on 19-Mar-2019

Summary



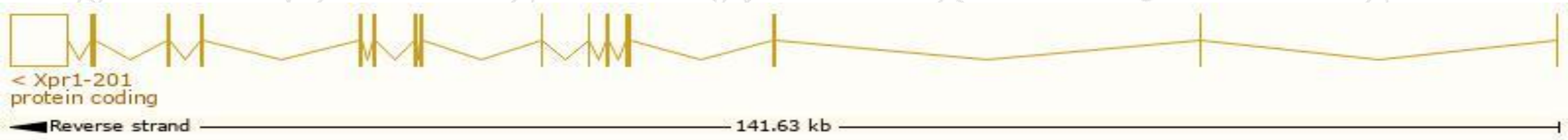
Official Symbol	Xpr1 provided by MGI
Official Full Name	xenotropic and polytropic retrovirus receptor 1 provided by MGI
Primary source	MGI:MGI:97932
See related	Ensembl:ENSMUSG00000026469
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Rmc-1, Rmc1, Sxv, Syg1, XR
Expression	Broad expression in whole brain E14.5 (RPKM 29.3), CNS E18 (RPKM 27.0) and 26 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

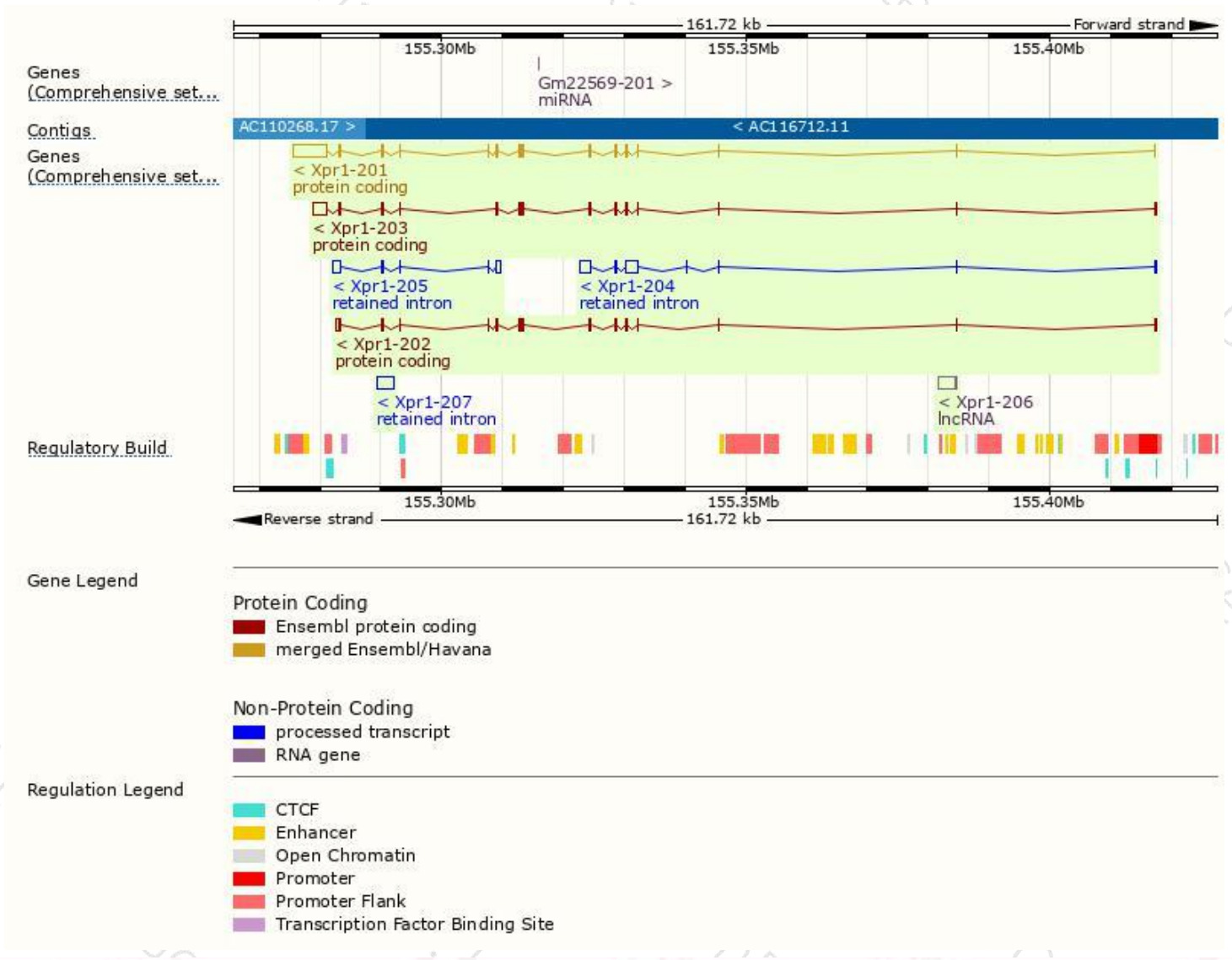
The gene has 7 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Xpr1-201	ENSMUST00000027741.11	7492	695aa	Protein coding	CCDS15384	Q9Z0U0	TSL:1 GENCODE basic APPRIS P1
Xpr1-203	ENSMUST00000111775.7	4094	630aa	Protein coding	-	D3Z2B4	TSL:5 GENCODE basic
Xpr1-202	ENSMUST00000111774.1	2753	679aa	Protein coding	-	Q9Z0U0	TSL:1 GENCODE basic
Xpr1-204	ENSMUST00000129271.7	4246	No protein	Retained intron	-	-	TSL:1
Xpr1-207	ENSMUST00000192359.1	2738	No protein	Retained intron	-	-	TSL:NA
Xpr1-205	ENSMUST00000139925.1	2443	No protein	Retained intron	-	-	TSL:1
Xpr1-206	ENSMUST00000143529.2	2881	No protein	lncRNA	-	-	TSL:2

The strategy is based on the design of *Xpr1-201* transcript,The transcription is shown below



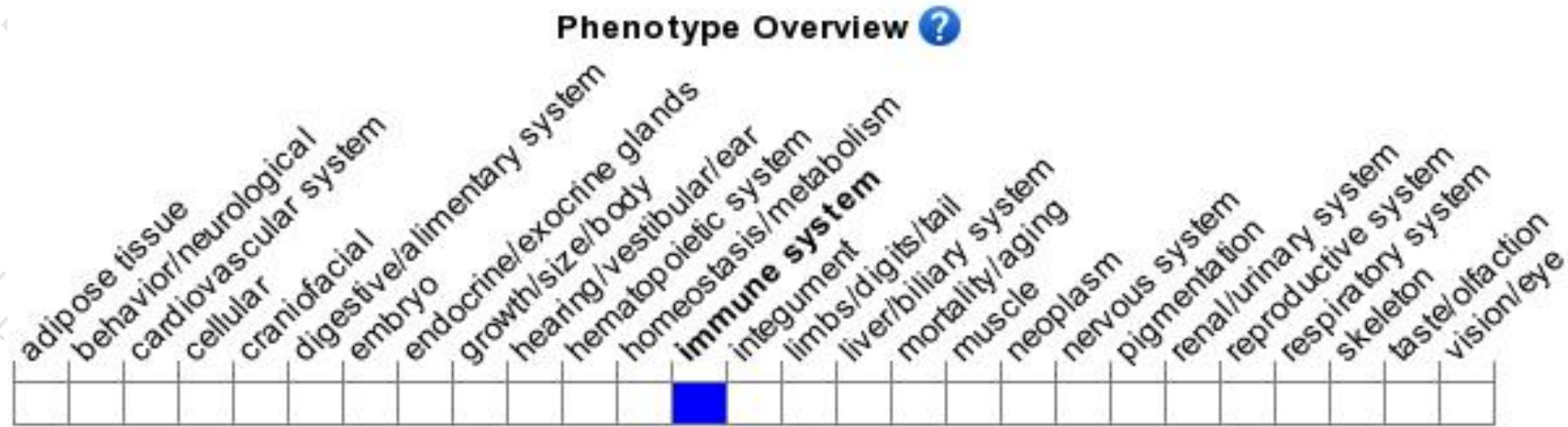
Genomic location distribution



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/>).

According to the existing MGI data, Homozygotes and heterozygotes for a variant from some wild Mus stocks, including *M. spretus*, support replication of xenotropic murine leukemia viruses and mink cell focus-forming murine leukemia viruses that are not replicated in most laboratory strains.

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

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