

Aldoa Cas9-KO Strategy

Designer: Xiangli Bian

Reviewer: Yao Yu

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Overview

Target Gene Name

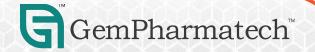
• Aldoa

Project Type

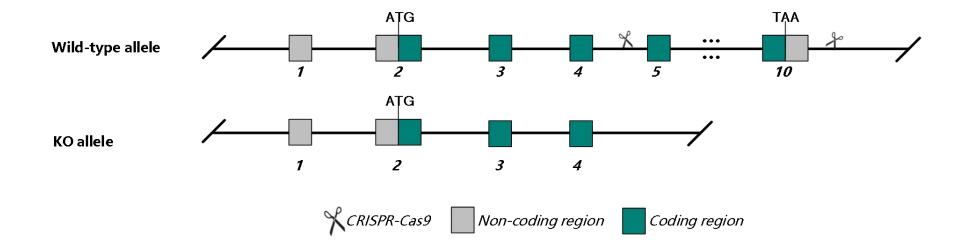
• Cas9-KO

Genetic Background

• C57BL/6JGpt



Strain Strategy



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



Technical Information

- The *Aldoa* gene has 12 transcripts. According to the structure of *Aldoa* gene, exon 5-10 of *Aldoa*-202 (ENSMUST00000087566.11) is recommended as the knockout region. The region contains 771 bp of coding sequence. Knocking out the region will result in disruption of gene function.
- In this project we use CRISPR-Cas9 technology to modify *Aldoa* gene. The brief process is as follows: gRNAs were transcribed in vitro. Cas9 and gRNAs 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 ontarget 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.

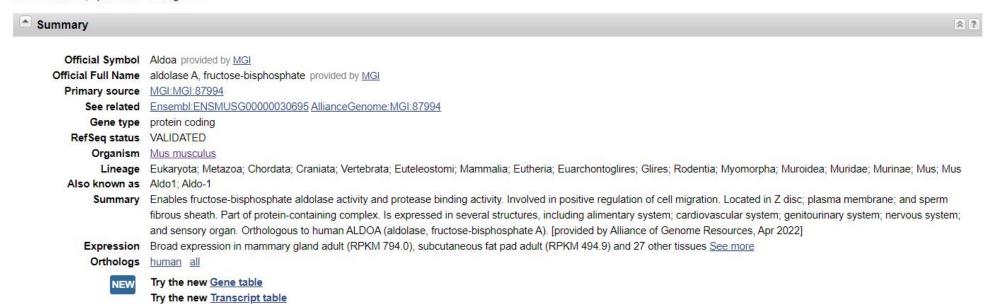


Gene Information

Aldoa aldolase A, fructose-bisphosphate [Mus musculus (house mouse)]

L Download Datasets

Gene ID: 11674, updated on 18-Aug-2023



Genomic context

☆ ?

Location: 7 F3; 7 69.25 cM

See Aldoa in Genome Data Viewer

Exon count: 12

https://www.ncbi.nlm.nih.gov/gene/11674

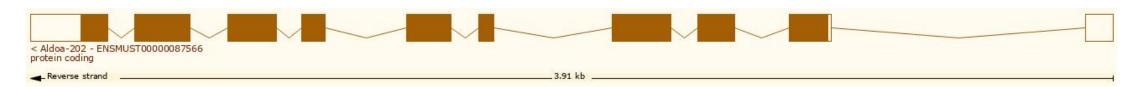


Transcript Information

The gene has 12 transcripts, all transcripts are shown below:

Transcript ID	Name 🎂	bp 🎂	Protein ▼	Biotype	CCDS	UniProt Match	Flags
ENSMUST00000087566.11	Aldoa-202	1549	<u>418aa</u>	Protein coding	CCDS52399 ₺	<u>A6ZI44</u> ₽	Ensembl Canonical GENCODE basic TSL:5
ENSMUST00000032934.12	Aldoa-201	1511	<u>364aa</u>	Protein coding	CCDS21845 ₺	P05064@Q5FWB7@	GENCODE basic APPRIS P1 TSL:1
ENSMUST00000106348.8	Aldoa-203	1460	<u>364aa</u>	Protein coding	CCDS21845 &	P05064@Q5FWB7@	GENCODE basic APPRIS P1 TSL:1
ENSMUST00000151137.8	Aldoa-210	776	<u>236aa</u>	Protein coding		D3YWI1 ₺	TSL:2 CDS 3' incomplete
ENSMUST00000205336.2	Aldoa-211	796	209aa	Protein coding		A0A0U1RPT5₽	TSL:5 CDS 3' incomplete
ENSMUST00000133514.8	Aldoa-204	626	<u>193aa</u>	Protein coding		D3YV98 ₽	TSL:3 CDS 3' incomplete
ENSMUST00000205890.2	Aldoa-212	762	<u>190aa</u>	Protein coding		A0A0U1RPN8₺	TSL:5 CDS 3' incomplete
ENSMUST00000141355.4	Aldoa-207	879	180aa	Protein coding		D3Z510@	TSL:5 CDS 3' incomplete
ENSMUST00000135707.8	Aldoa-205	945	No protein	Protein coding CDS not defined		-	TSL:5
ENSMUST00000139657.8	Aldoa-206	1352	No protein	Retained intron		(a)	TSL:2
ENSMUST00000147748.2	Aldoa-209	895	No protein	Retained intron		100	TSL:5
ENSMUST00000143509.8	Aldoa-208	891	No protein	Retained intron		-	TSL:2

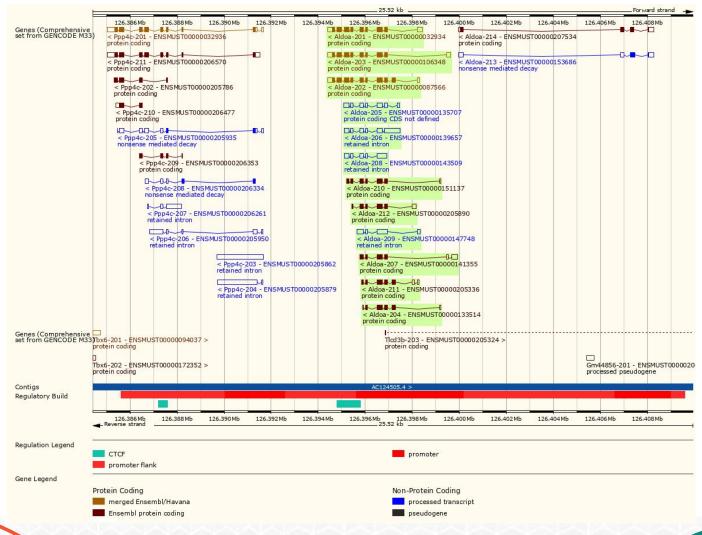
The strategy is based on the design of *Aldoa*-202 transcript, the transcription is shown below:





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

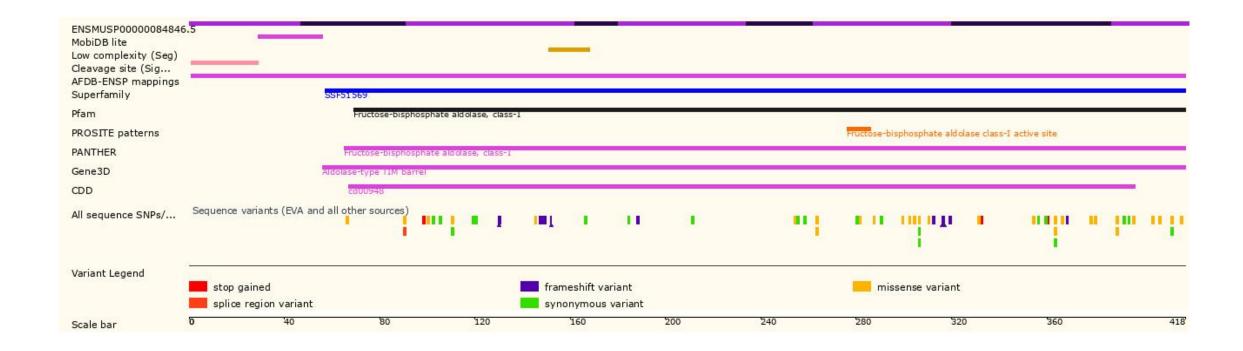
Genomic Information





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

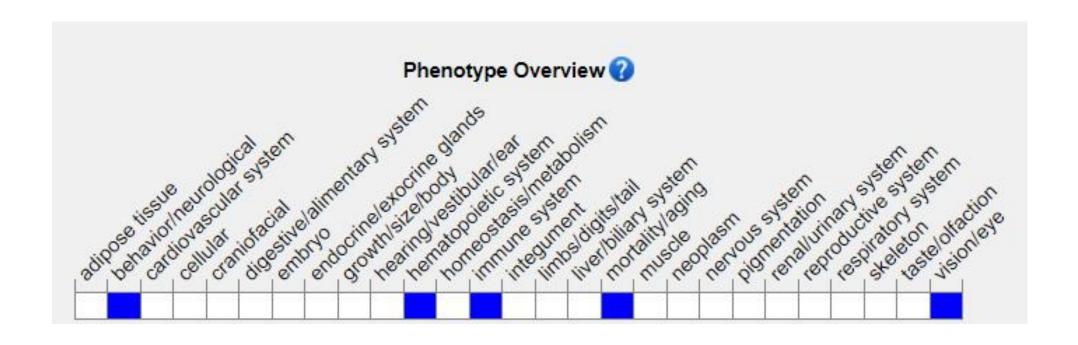
Protein Information

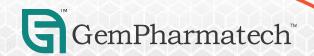




Source: https://www.ensembl.org

Mouse Phenotype Information (MGI)





Important Information

- The knockout region is about 2.5 kb away from the 5' of the *Ppp4c* gene, which may affect the regulation of this gene.
- The knockout region is about 0.5 kb away from the 5' of the *Tlcd3b* gene, which may affect the regulation of this gene.
- This stratergy may not affect *Aldoa*-213 and *Aldoa*-214 transcript.
- A part of amino acid sequence (162 aa) will still remain at the N-terminal of *Aldoa* gene
- *Aldoa* is located on Chr 7. 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 risks of the mutation on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

