

Arsb Cas9-CKO Strategy

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Project Overview



Project Name Arsb

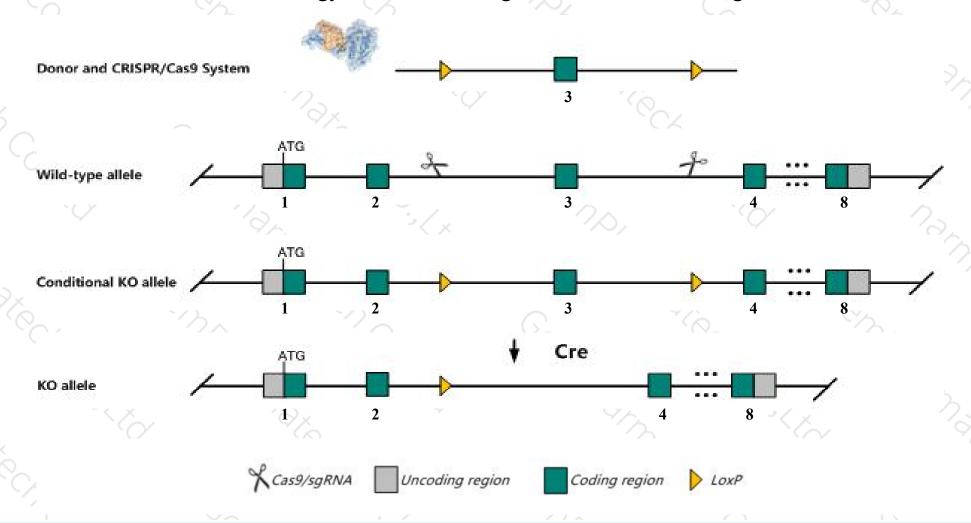
Project type Cas9-CKO

Strain background C57BL/6JGpt

Conditional Knockout strategy



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



Technical routes



- ➤ The Arsb gene has 2 transcripts. According to the structure of Arsb gene, exon3 of Arsb-201

 (ENSMUST00000091403.5) transcript is recommended as the knockout region. The region contains 191bp coding sequence.

 Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Arsb* gene. The brief process is as follows:sgRNA was transcribed in vitro, donor vector was constructed.Cas9, sgRNA 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 was 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, Homozygous mutation of this gene results in development of shortened limbs and snout and a broadened head after 4 weeks of age. Mutant animals have elevated concentrations of glucosaminoglycans in the urine and irregular cartilage structure.
- ➤ The KO region is close to *Mir5624* gene. Knockout the region may affect the function of *Mir5624* gene.
- The *Arsb* gene is located on the Chr13. 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)



Arsb arylsulfatase B [Mus musculus (house mouse)]

Gene ID: 11881, updated on 5-Nov-2019

Summary

2 1

Official Symbol Arsb provided by MGI

Official Full Name arylsulfatase B provided by MGI

Primary source MGI:MGI:88075

See related Ensembl: ENSMUSG00000042082

Gene type protein coding
RefSeq status VALIDATED
Organism <u>Mus musculus</u>

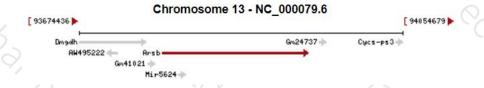
Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;

Muroidea; Muridae; Murinae; Mus; Mus

Also known as As1; As-1; As-1s; As-1s; As-1r; As1-r; As1-r; As1-r; As1-t; Asr-1; Asr-1; Al480648; 1110007C02Rik

Expression Ubiquitous expression in kidney adult (RPKM 45.3), ovary adult (RPKM 25.9) and 28 other tissues See more

Orthologs <u>human</u> all



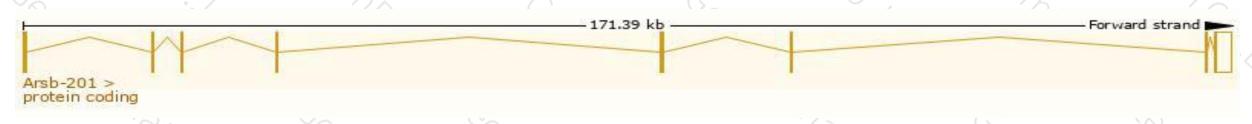
Transcript information (Ensembl)



The gene has 2 transcripts, all transcripts are shown below:

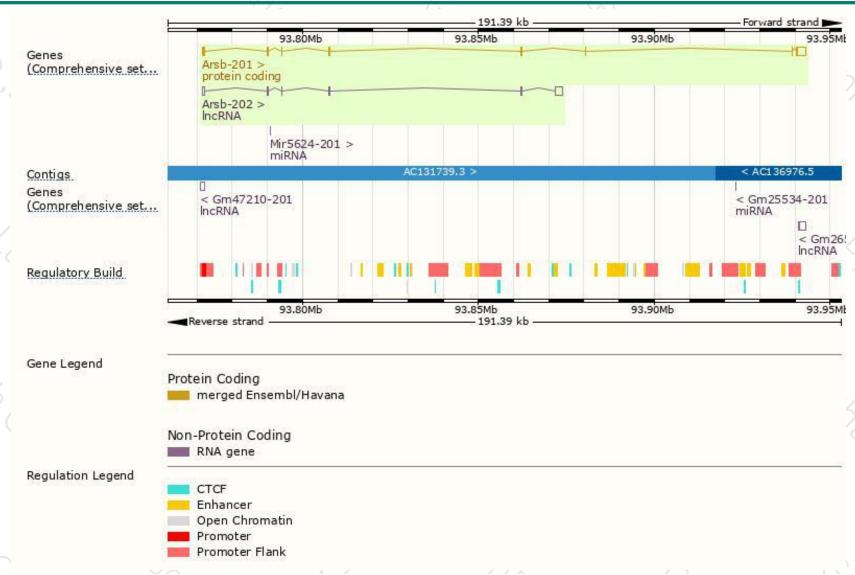
Name 👙	Transcript ID 🍦	bp 🌲	Protein 🍦	Translation ID 🍦	Biotype 🍦	CCDS 🍦	UniProt	Flags		
Arsb-201	ENSMUST00000091403.5	4038	<u>534aa</u>	ENSMUSP00000088964.4	Protein coding	CCDS36749₽	A0A0R4J138 &	TSL:1	GENCODE basic	APPRIS P1
Arsb-202	ENSMUST00000220652.1	3226	No protein	5 2	IncRNA	\$ T 8		TSL:1		

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



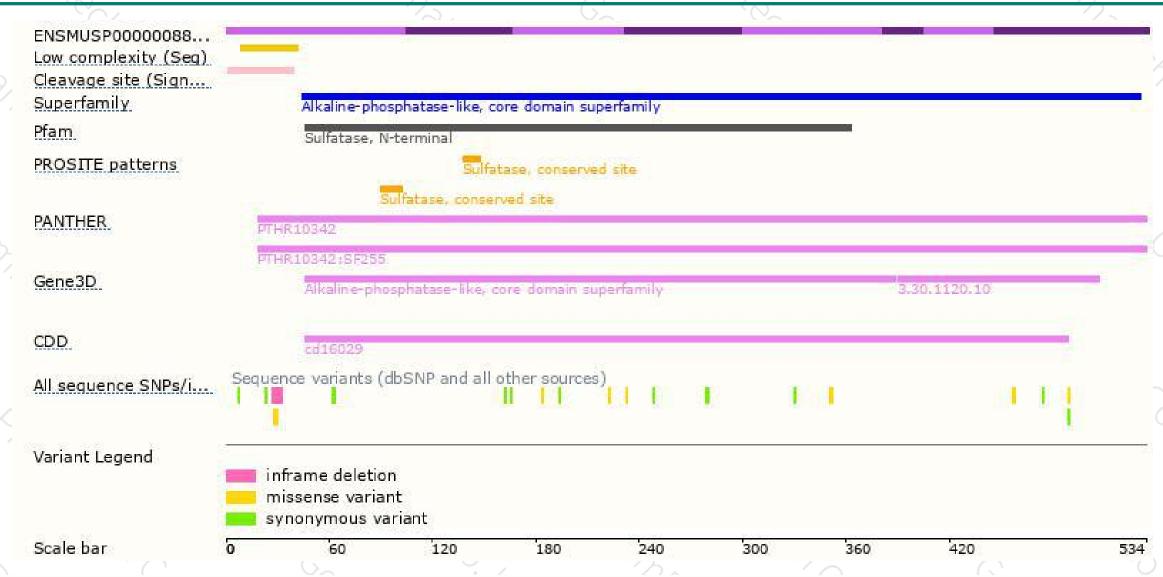
Genomic location distribution





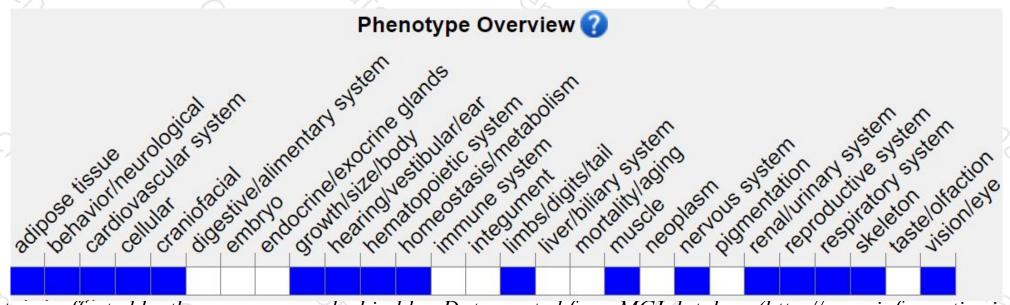
Protein domain





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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