

Arid4b Cas9-CKO Strategy

Designer: Huimin Su

Reviewer: Ruiuri Zhang

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Project Overview



Project Name

Arid4b

Project type

Cas9-CKO

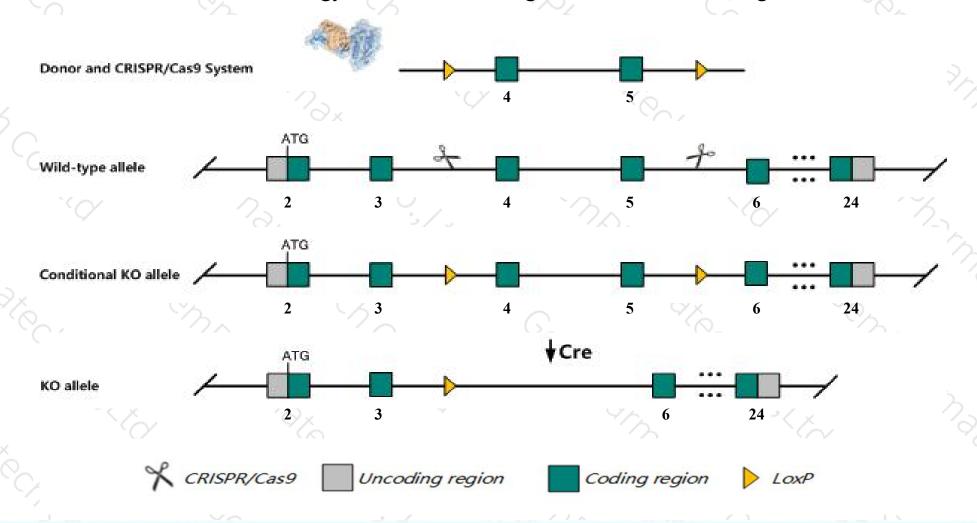
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Arid4b* gene. The schematic diagram is as follows:



Technical routes



- The *Arid4b* gene has 12 transcripts. According to the structure of *Arid4b* gene, exon4-exon5 of *Arid4b*-203(ENSMUST00000110534.7) transcript is recommended as the knockout region. The region contains 157bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Arid4b* 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 homozygous for a null allele die pre-implantation.
- > The *Arid4b* 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)



Arid4b AT rich interactive domain 4B (RBP1-like) [Mus musculus (house mouse)]

Gene ID: 94246, updated on 26-Jun-2020

Summary

☆ ?

Official Symbol Arid4b provided by MGI

Official Full Name AT rich interactive domain 4B (RBP1-like) provided by MGI

Primary source MGI:MGI:2137512

See related Ensembl: ENSMUSG00000039219

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 BCAA; BRCAA1; Rbp1I1; SAP180; RBBP1L1

Expression Ubiquitous expression in CNS E11.5 (RPKM 10.1), CNS E14 (RPKM 7.3) and 26 other tissues See more

Orthologs human all

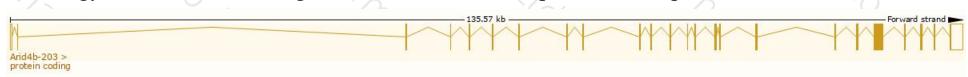
Transcript information (Ensembl)



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

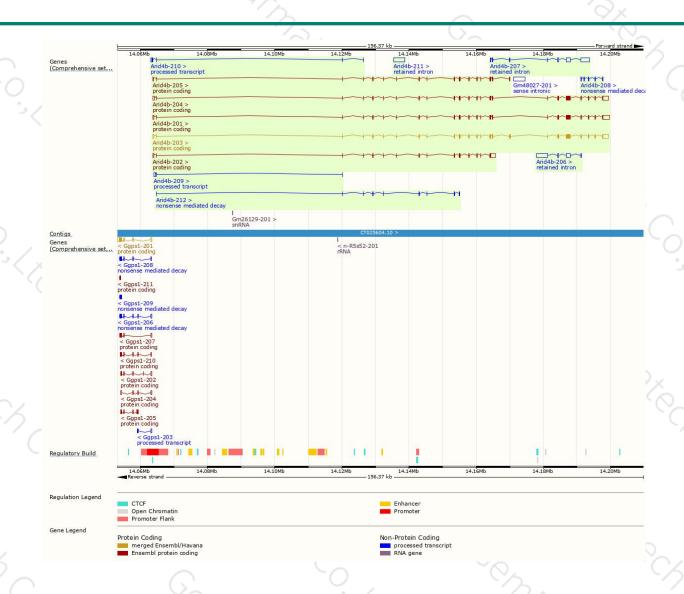
Name	Transcript ID	bp 👙	Protein A	Biotype	CCDS	UniProt #	Flags
Arid4b-208	ENSMUST00000151151.1	745	<u>31aa</u>	Nonsense mediated decay	-	A0A1Y7VKA9₽	CDS 5' incomplete TSL:3
Arid4b-212	ENSMUST00000222928.1	815	<u>128aa</u>	Nonsense mediated decay	*	A0A1Y7VIX3&	TSL:5
Arid4b-202	ENSMUST00000110533.1	3020	<u>473aa</u>	Protein coding	-	Z4YL44@	TSL:1 GENCODE basic
Arid4b-205	ENSMUST00000129488.7	1802	<u>540aa</u>	Protein coding	일	Z4YMH1₽	CDS 3' incomplete TSL:1
Arid4b-201	ENSMUST00000039538.14	5844	<u>1227aa</u>	Protein coding	CCDS36600@	A2CG63 ₺	TSL:5 GENCODE basic
Arid4b-204	ENSMUST00000110536.7	5783	<u>1227aa</u>	Protein coding	CCDS36600@	A2CG63 &	TSL:5 GENCODE basic
Arid4b-203	ENSMUST00000110534.7	5864	<u>1314aa</u>	Protein coding	CCDS36599 ₽	A2CG63 ₺	TSL:1 GENCODE basic APPRIS P1
Arid4b-209	ENSMUST00000151182.1	678	No protein	Processed transcript	-	9.5	TSL:3
Arid4b-210	ENSMUST00000155553.7	571	No protein	Processed transcript	-	(*)	TSL:3
Arid4b-206	ENSMUST00000148941.1	4828	No protein	Retained intron	-	-	TSL:2
Arid4b-207	ENSMUST00000149579.2	4571	No protein	Retained intron	-		TSL:5
Arid4b-211	ENSMUST00000222346.1	3353	No protein	Retained intron	일	<u> 92</u> 59	TSL:NA
	7/3/		777				L.V

The strategy is based on the design of Arid4b-203 transcript, the transcription is shown below:



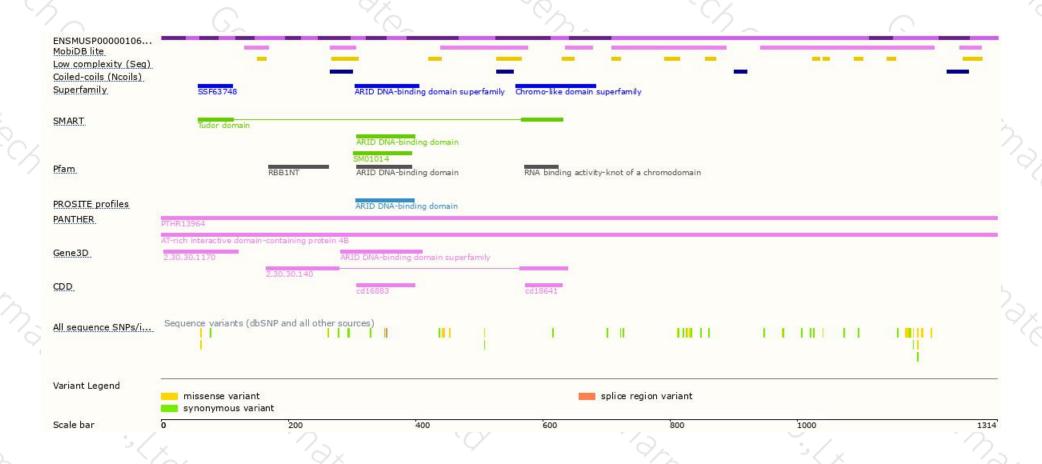
Genomic location distribution





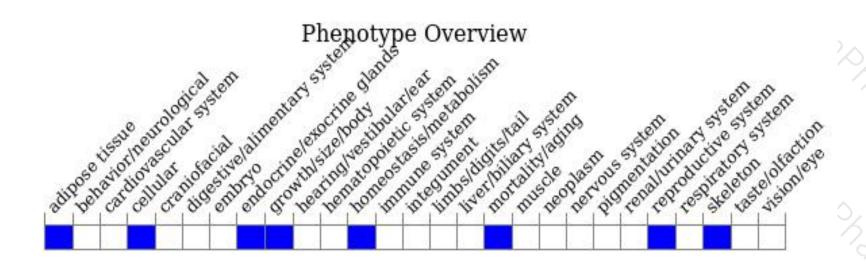
Protein domain





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, mice homozygous for a null allele die pre-implantation.



If you have any questions, you are welcome to inquire. Tel: 400-9660890





