



Atad2b Cas9-CKO Strategy

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Project Overview

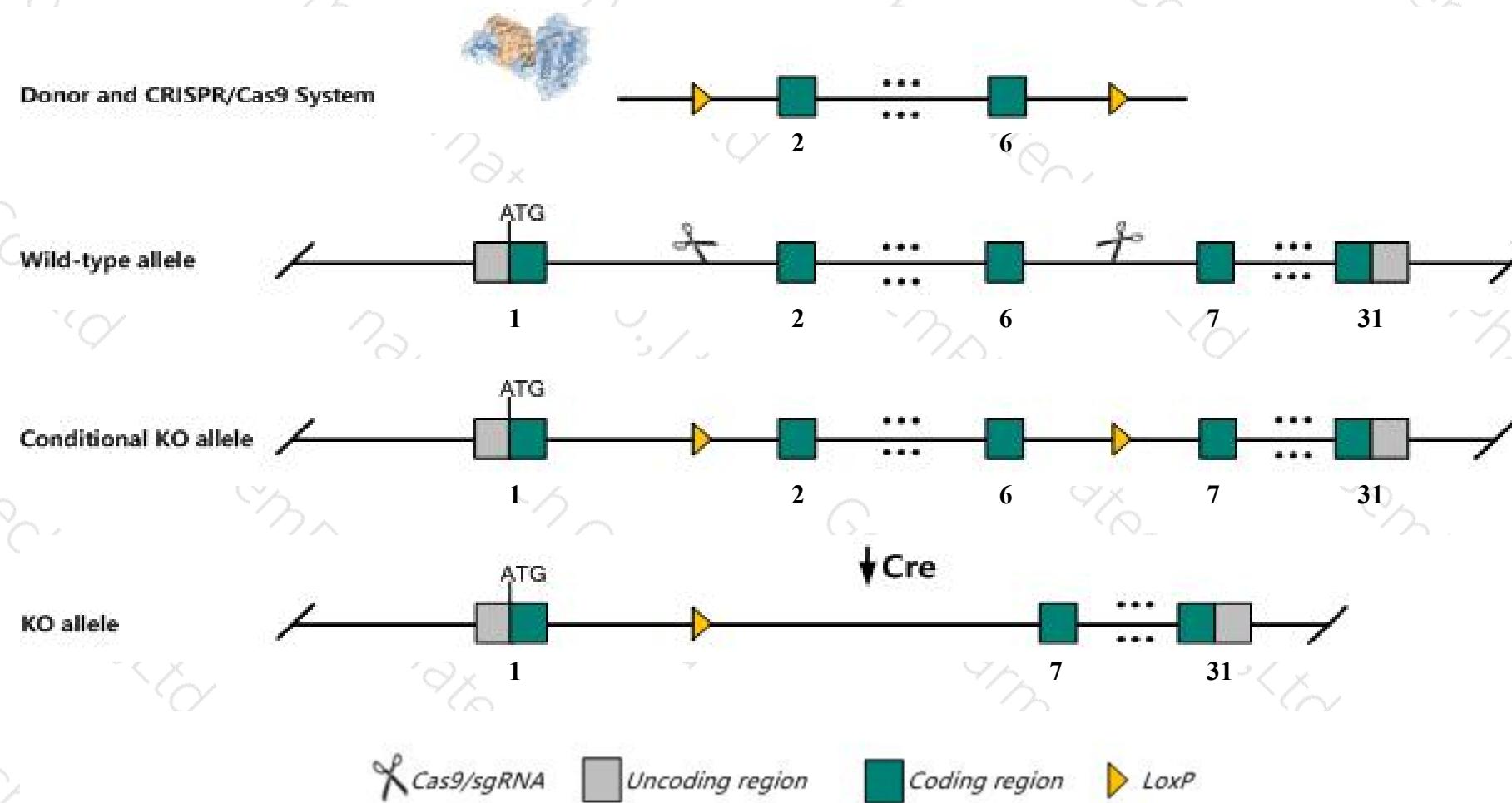
Project Name***Atad2b***

Project type**Cas9-CKO**

Strain background**C57BL/6JGpt**

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Atad2b* gene. The schematic diagram is as follows:



Technical routes

- The *Atad2b* gene has 5 transcripts. According to the structure of *Atad2b* gene, exon2-exon6 of *Atad2b*-201(ENSMUST00000045664.6) transcript is recommended as the knockout region. The region contains 568bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Atad2b* 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.



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Notice

- According to the existing MGI data, mice homozygous for a transgenic gene disruption exhibit reduced body size and fertility in female mice.
- Transcript *Atad2b*-203&205 may not be affected.
- The *Atad2b* gene is located on the Chr12. 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)

Atad2b ATPase family, AAA domain containing 2B [Mus musculus (house mouse)]

Gene ID: 320817, updated on 13-Mar-2020

Summary



Official Symbol Atad2b provided by [MGI](#)

Official Full Name ATPase family, AAA domain containing 2B provided by [MGI](#)

Primary source [MGI:MGI:2444798](#)

See related [Ensembl:ENSMUSG00000052812](#)

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 1110014E10Rik, BC032887, C79189, D530031C13Rik

Expression Ubiquitous expression in CNS E11.5 (RPKM 5.7), CNS E14 (RPKM 4.6) and 28 other tissues [See more](#)

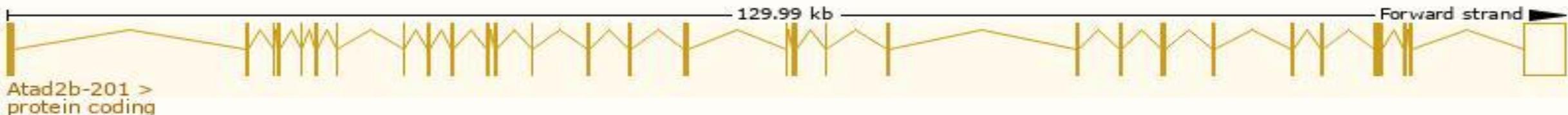
Orthologs [human](#) [all](#)

Transcript information (Ensembl)

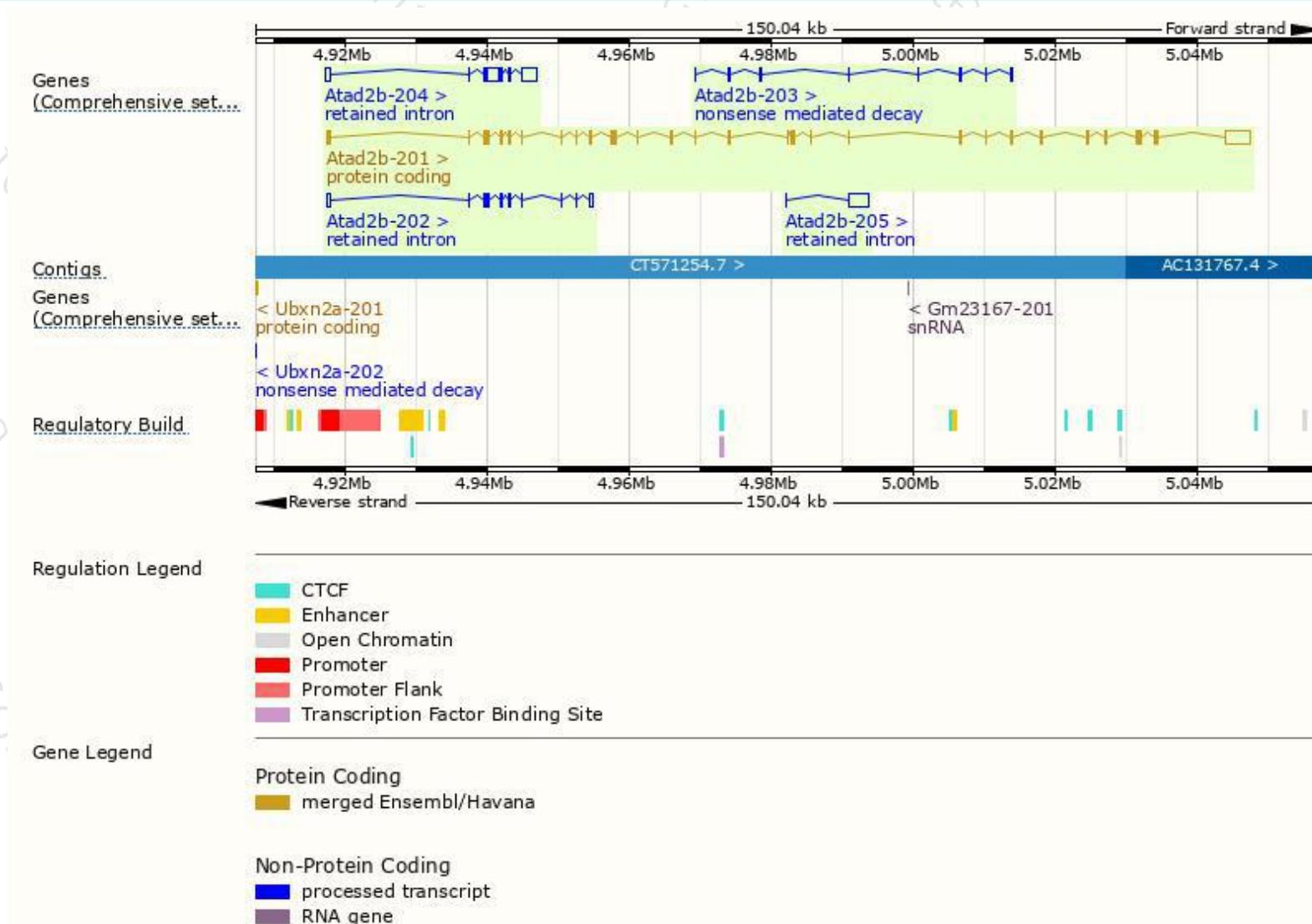
The gene has 5 transcripts, all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Atad2b-201	ENSMUST00000045664.6	8035	1460aa	Protein coding	-	E9Q166	TSL:5 GENCODE basic APPRIS P1
Atad2b-203	ENSMUST00000218859.1	1343	259aa	Nonsense mediated decay	-	A0A1W2P7J4	CDS 5' incomplete TSL:1
Atad2b-204	ENSMUST00000219187.1	4829	No protein	Retained intron	-	-	TSL:1
Atad2b-205	ENSMUST00000220408.1	3034	No protein	Retained intron	-	-	TSL:1
Atad2b-202	ENSMUST00000218303.1	1824	No protein	Retained intron	-	-	TSL:1

The strategy is based on the design of *Atad2b-201* transcript, the transcription is shown below:



Genomic location distribution



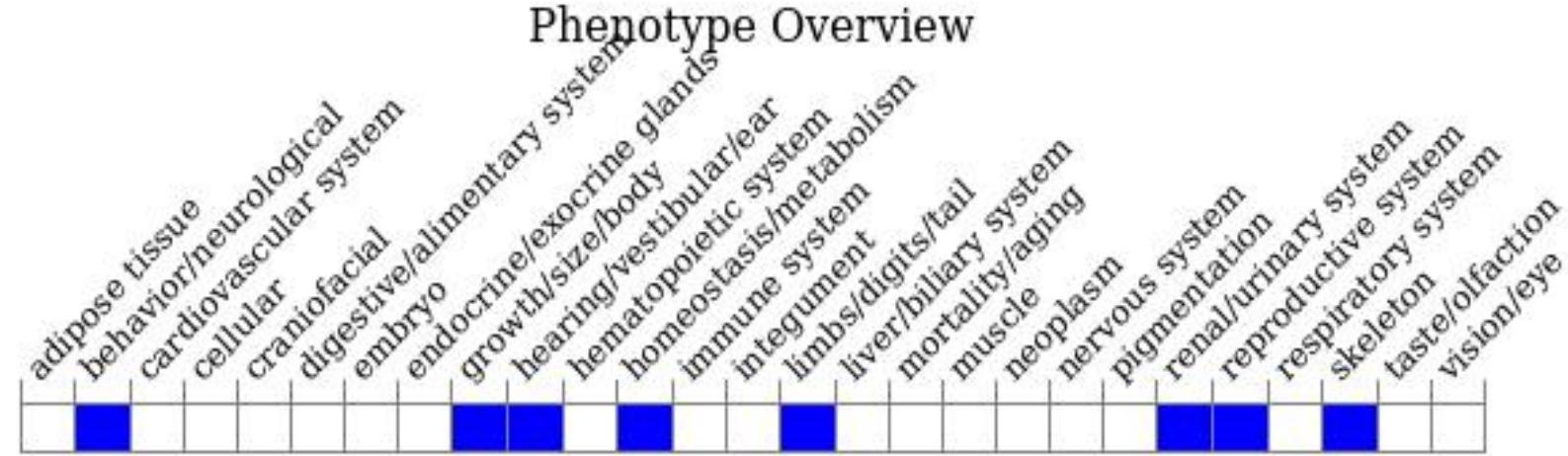
Protein domain





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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 transgenic gene disruption exhibit reduced body size and fertility in female mice.



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

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