

Avil Cas9-KO Strategy

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Project Overview



Project Name
Avil

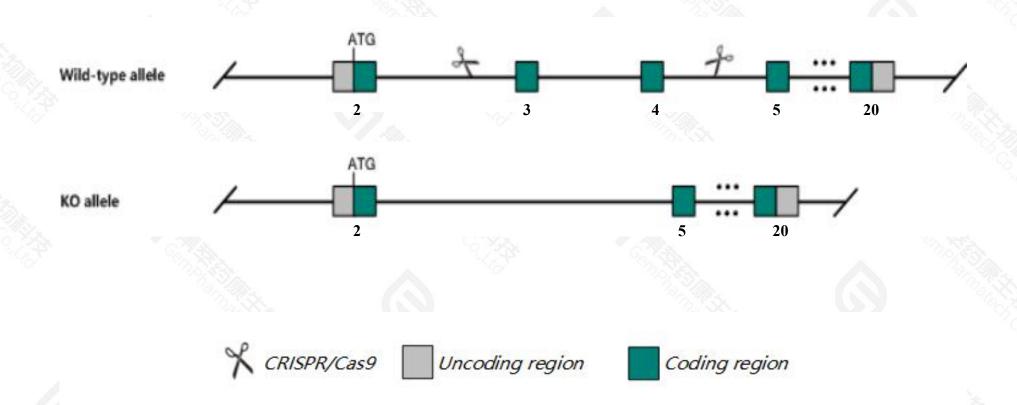
Project type
Cas9-KO

Strain background
C57BL/6JGpt

Knockout strategy



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



Technical routes



- The *Avil* gene has 4 transcripts. According to the structure of *Avil* gene, exon3-exon4 of *Avil-* 201(ENSMUST00000026500.12) transcript is recommended as the knockout region. The region contains 272bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Avil* gene. The brief process is as follows: CRISPR/Cas9 system 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.

Notice



- > According to the existing MGI data, homozygotes null mice show partial embryonic lethality before E10.5, but surviving mice are fertile and exhibit no abnormal behavior into adult. The regenerative axon growth and remodeling of sensory nerves are abnormal in homozygous null mice.
- > The Avil gene is located on the Chr10. 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)



Avil advillin [Mus musculus (house mouse)]

Gene ID: 11567, updated on 26-Jan-2021

Summary

☆ ?

Official Symbol Avil provided by MGI
Official Full Name advillin provided by MGI

Primary source MGI:MGI:1333798

See related Ensembl:ENSMUSG00000025432

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 Advil, DOC, DOC6

Expression Biased expression in stomach adult (RPKM 6.0), small intestine adult (RPKM 1.4) and 5 other tissuesSee more

Orthologs <u>human</u> all

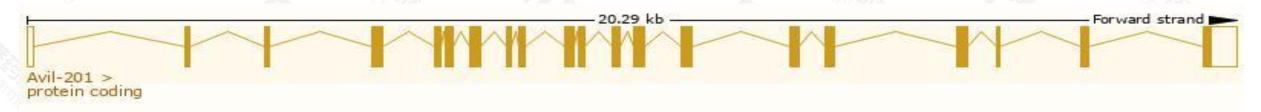
Transcript information (Ensembl)



The gene has 4 transcripts, all transcripts are shown below:

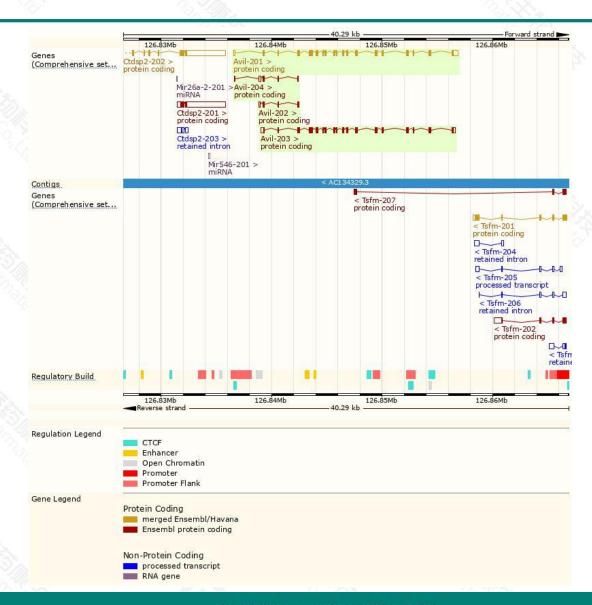
10. 100							
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Avil-201	ENSMUST00000026500.12	3033	819aa	Protein coding	CCDS24221		TSL:1, GENCODE basic, APPRIS P1,
Avil-203	ENSMUST00000129173.2	2925	<u>819aa</u>	Protein coding	CCDS24221		TSL:5 , GENCODE basic , APPRIS P1 ,
Avil-204	ENSMUST00000142698.8	419	<u>54aa</u>	Protein coding	72		CDS 3' incomplete , TSL:5 ,
Avil-202	ENSMUST00000126816.8	350	<u>78aa</u>	Protein coding	-		CDS 3' incomplete , TSL:5 ,

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



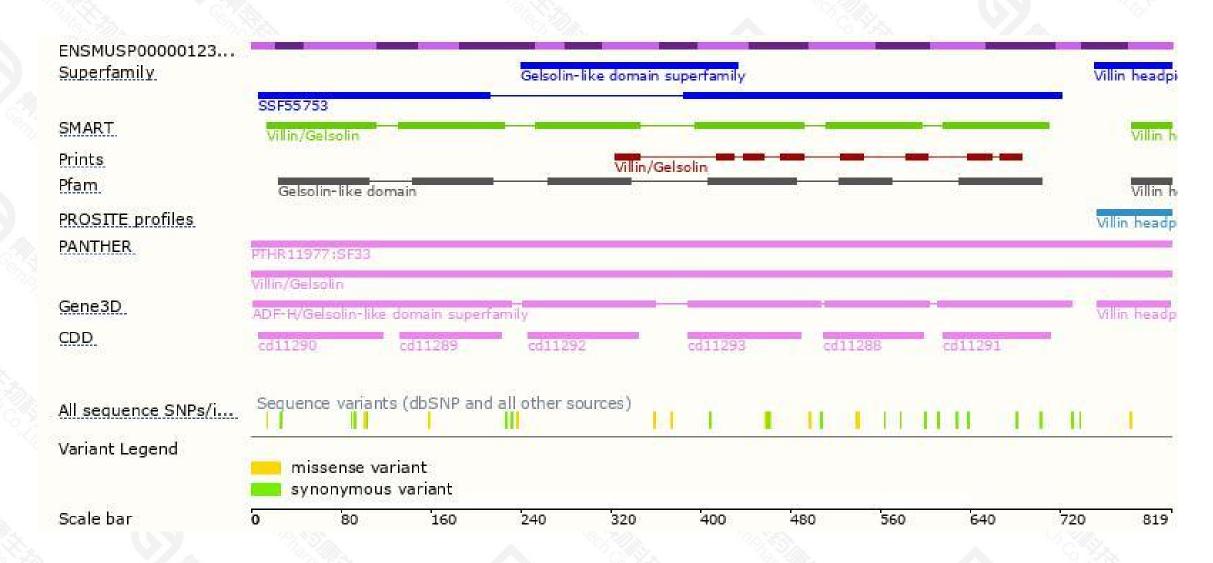
Genomic location distribution





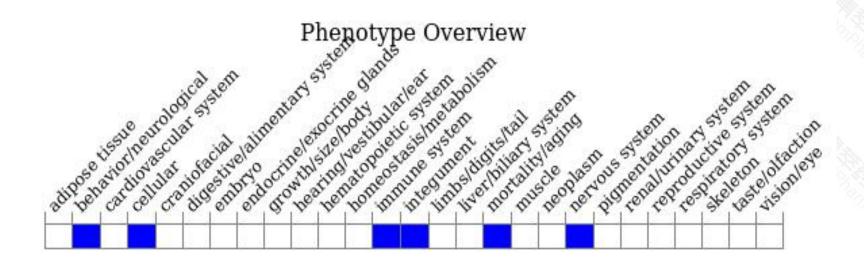
Protein domain





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 null mice show partial embryonic lethality before E10.5, but surviving mice are fertile and exhibit no abnormal behavior into adult. The regenerative axon growth and remodeling of sensory nerves are abnormal in homozygous null mice.



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

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