

Abhd6 Cas9-CKO Strategy

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Project Overview



Project Name

Abhd6

Project type

Cas9-CKO

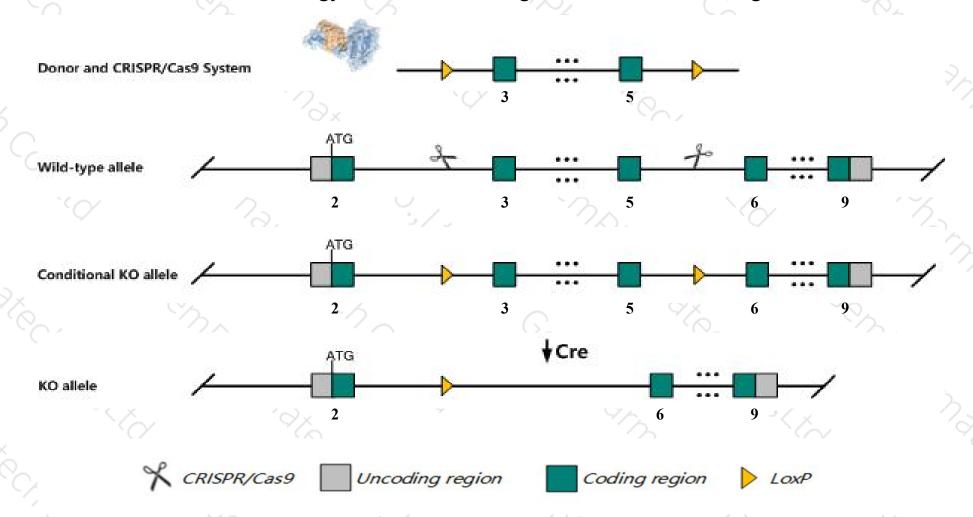
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Abhd6* gene. The schematic diagram is as follows:



Technical routes



- The *Abhd6* gene has 3 transcripts. According to the structure of *Abhd6* gene, exon3-exon5 of *Abhd6-202* (ENSMUST00000166497.8) transcript is recommended as the knockout region. The region contains 404bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Abhd6* 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, homozygous null mice show increased glucose-stimulated insulin secretion from islets which exhibit elevated monoacylglycerol content in response to glucose.
- ➤ The *Abhd6* gene is located on the Chr14. 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)



Abhd6 abhydrolase domain containing 6 [Mus musculus (house mouse)]

Gene ID: 66082, updated on 13-Mar-2020

Summary

☆ ?

Official Symbol Abhd6 provided by MGI

Official Full Name abhydrolase domain containing 6 provided by MGI

Primary source MGI:MGI:1913332

See related Ensembl: ENSMUSG00000025277

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 0610041D24Rik, AA673485, AV065425

Expression Ubiquitous expression in adrenal adult (RPKM 11.3), large intestine adult (RPKM 9.3) and 28 other tissuesSee more

Orthologs human all

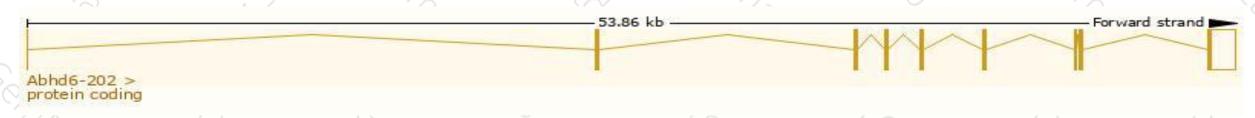
Transcript information (Ensembl)



The gene has 3 transcripts, all transcripts are shown below:

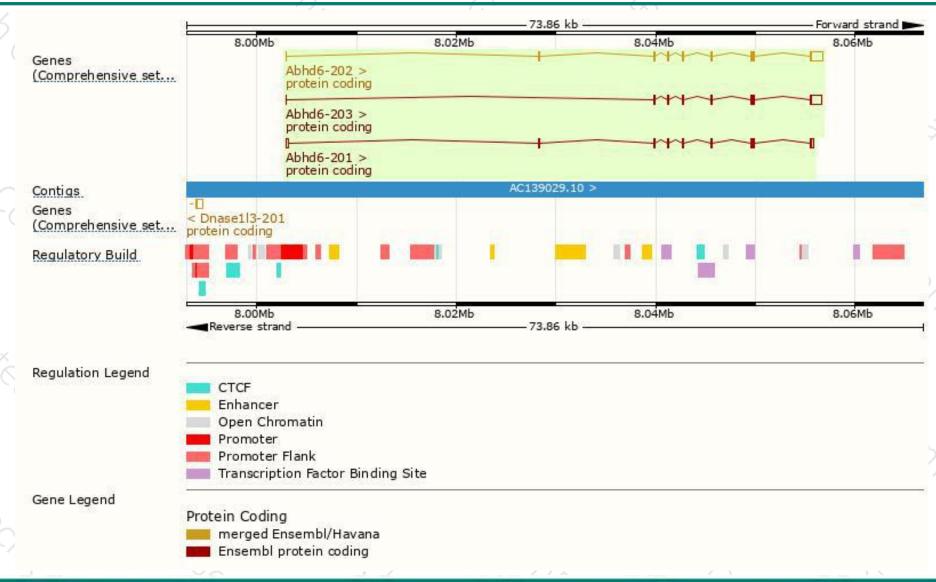
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Abhd6-202		2226		Protein coding			TSL:1 GENCODE basic APPRIS is a system to annotate alternatively spliced transcripts based on a range of computational methods to identify the most functionally important transcript(s) of a gene. APPRIS P
					CCD320000	30 Sept. 2000 - 10 Sept. 200	Septimental control of the septiment of
Abhd6-201	ENSMUST00000026313.3	1401	336aa	Protein coding	CCDS26808	Q8R2Y0	TSL:1 GENCODE basic APPRIS is a system to annotate alternatively spliced transcripts based on a range of computational methods to identify the most functionally important transcript(s) of a gene. APPRIS P
Abhd6-203	ENSMUST00000225234.1	1958	289aa	Protein coding	0	Q8R2Y0 Q9D375	GENCODE basic

The strategy is based on the design of *Abhd6-202* transcript, the transcription is shown below



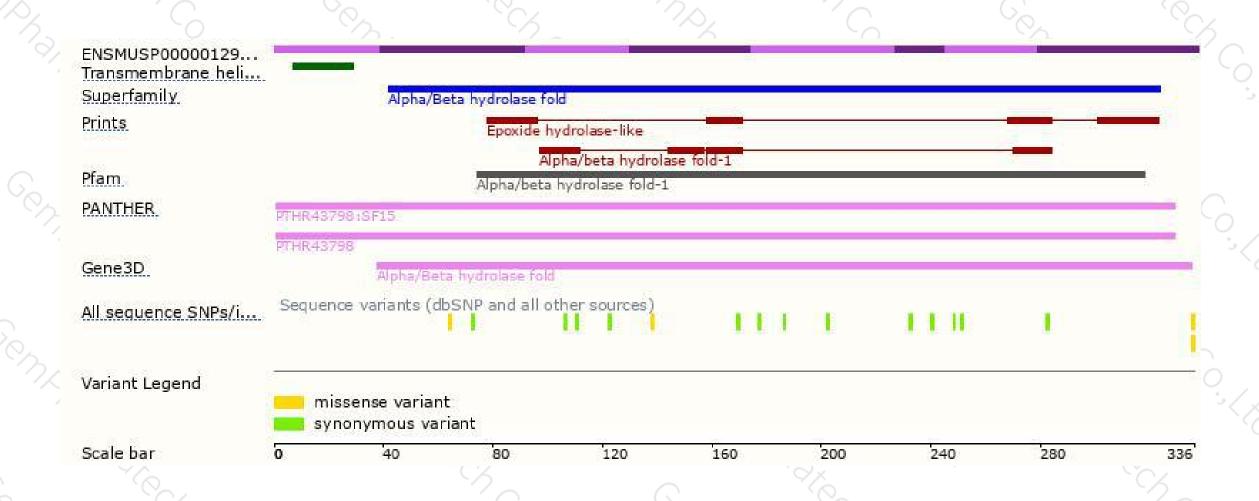
Genomic location distribution





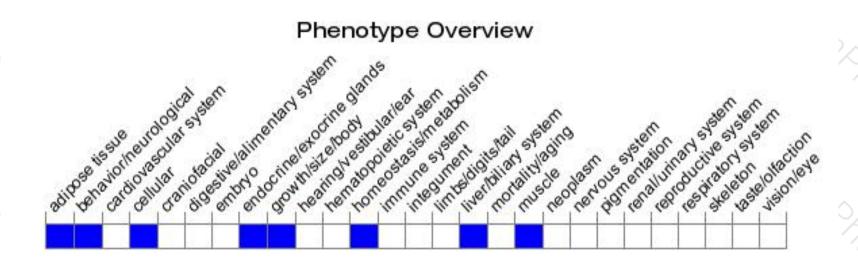
Protein domain





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, homozygous null mice show increased glucose-stimulated insulin secretion from islets which exhibit elevated monoacylglycerol content in response to glucose.



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





