

***Hacd2* Cas9-KO Strategy**

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Project Overview

Project Name

Hacd2

Project type

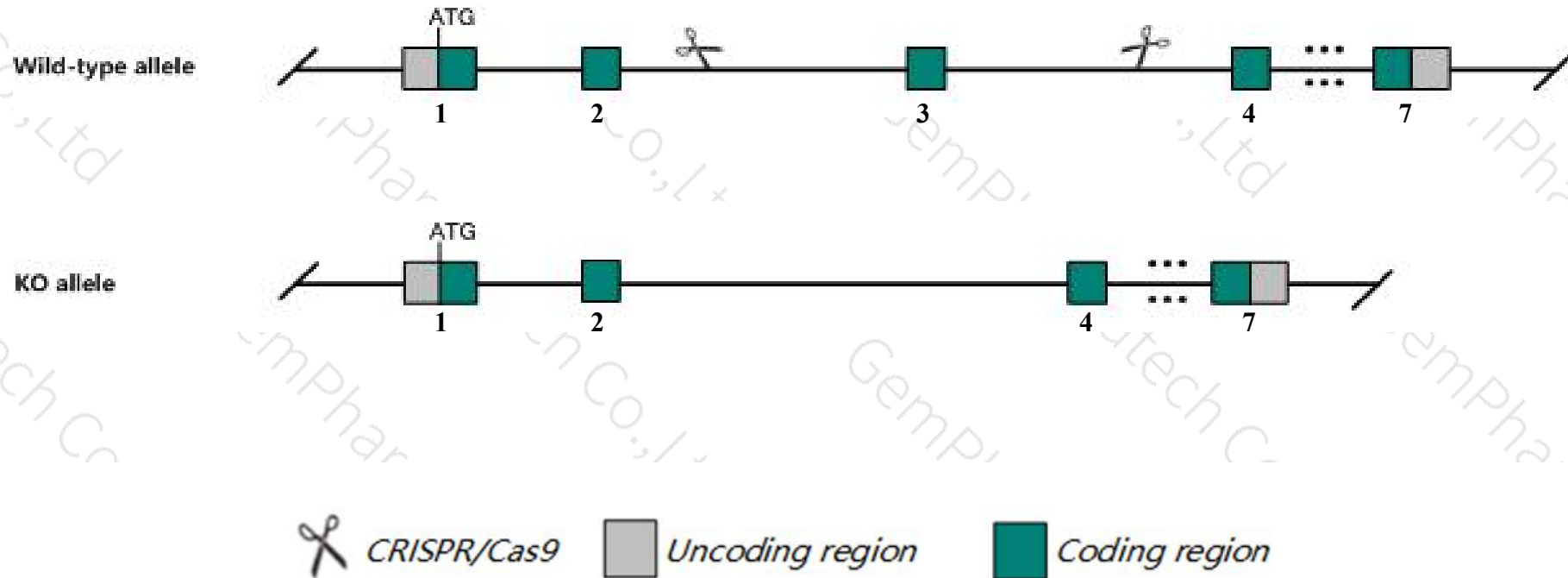
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Hacd2* gene. The schematic diagram is as follows:



- The *Hacd2* gene has 3 transcripts. According to the structure of *Hacd2* gene, exon3 of *Hacd2-201* (ENSMUST00000061156.9) transcript is recommended as the knockout region. The region contains 19bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Hacd2* gene. The brief process is as follows: CRISPR/Cas9 system

- The N-terminal of *Hacd2* gene will remain 91aa, it may remain the partial function of *Hacd2* gene.
- The knockout region is near to the N-terminal of *Gm21691* gene, this strategy may influence the regulatory function of the N-terminal of *Gm21691* gene.
- The *Hacd2* gene is located on the Chr16. 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)

Hacd2 3-hydroxyacyl-CoA dehydratase 2 [*Mus musculus* (house mouse)]

Gene ID: 70757, updated on 12-Aug-2019

Summary

Official Symbol Hacd2 provided by [MGI](#)
Official Full Name 3-hydroxyacyl-CoA dehydratase 2 provided by [MGI](#)
Primary source [MGI:MGI:1918007](#)
See related [Ensembl:ENSMUSG00000035376](#)
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 Hcad2; Ptplb; AI255777; AI481689; 6330408J20Rik
Expression Ubiquitous expression in genital fat pad adult (RPKM 24.6), subcutaneous fat pad adult (RPKM 18.9) and 28 other tissues [See more](#)
Orthologs [human](#) [all](#)

Genomic context

Location: 16; 16 B3 [See Hacd2 in Genome Data Viewer](#)
Exon count: 9

Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	16	NC_000082.6 (35022421..35109190)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	16	NC_000082.5 (35022507..35109261)

Transcript information (Ensembl)

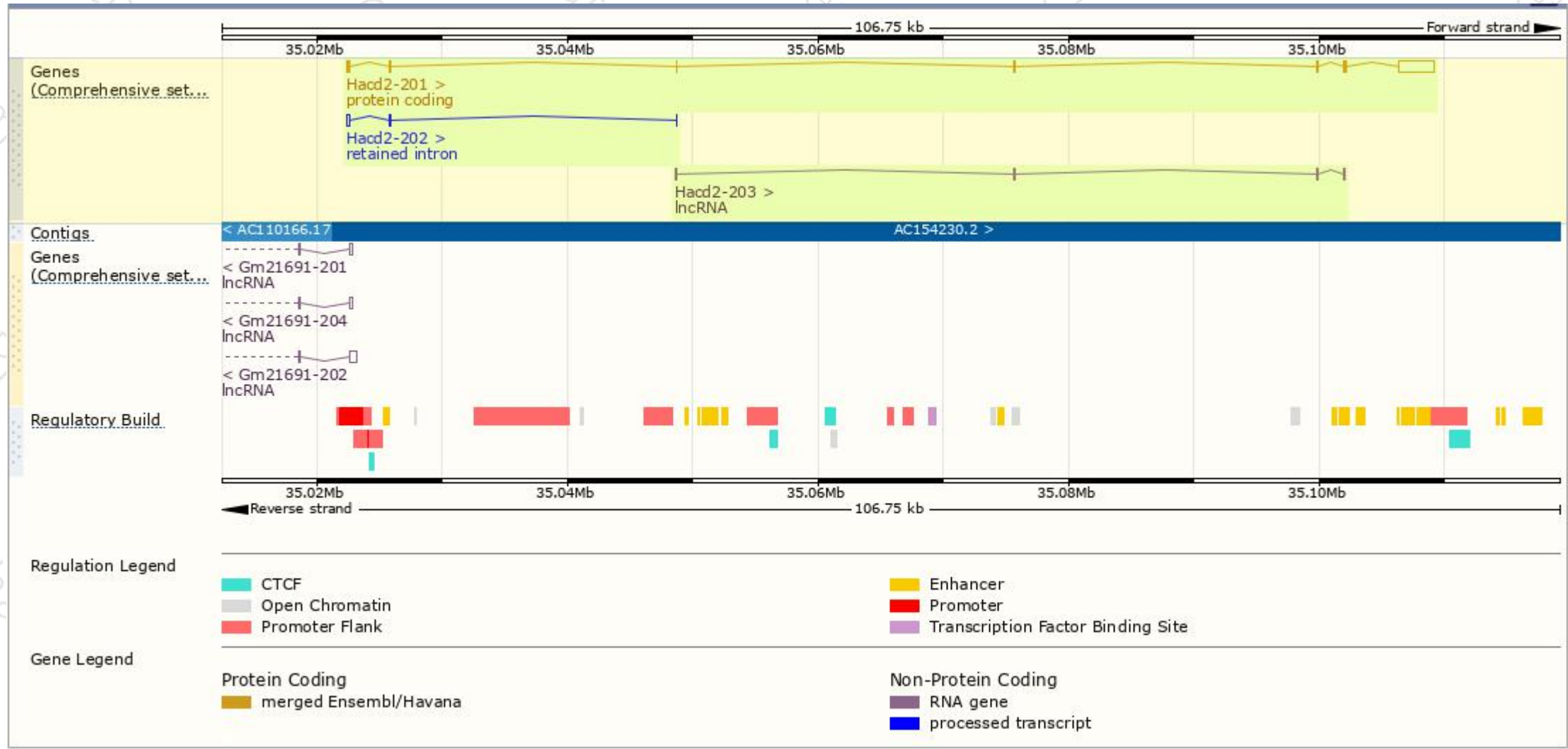
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Hacd2-201	ENSMUST00000061156.9	3598	254aa	Protein coding	CCDS37321	Q9D3B1	TSL:1 GENCODE basic APPRIS P1
Hacd2-203	ENSMUST00000231556.1	421	No protein	Processed transcript	-	-	
Hacd2-202	ENSMUST00000231516.1	389	No protein	Retained intron	-	-	

The strategy is based on the design of *Hacd2-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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