

# ***Fads6* Cas9-KO Strategy**

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# Project Overview

**Project Name**

*Fads6*

**Project type**

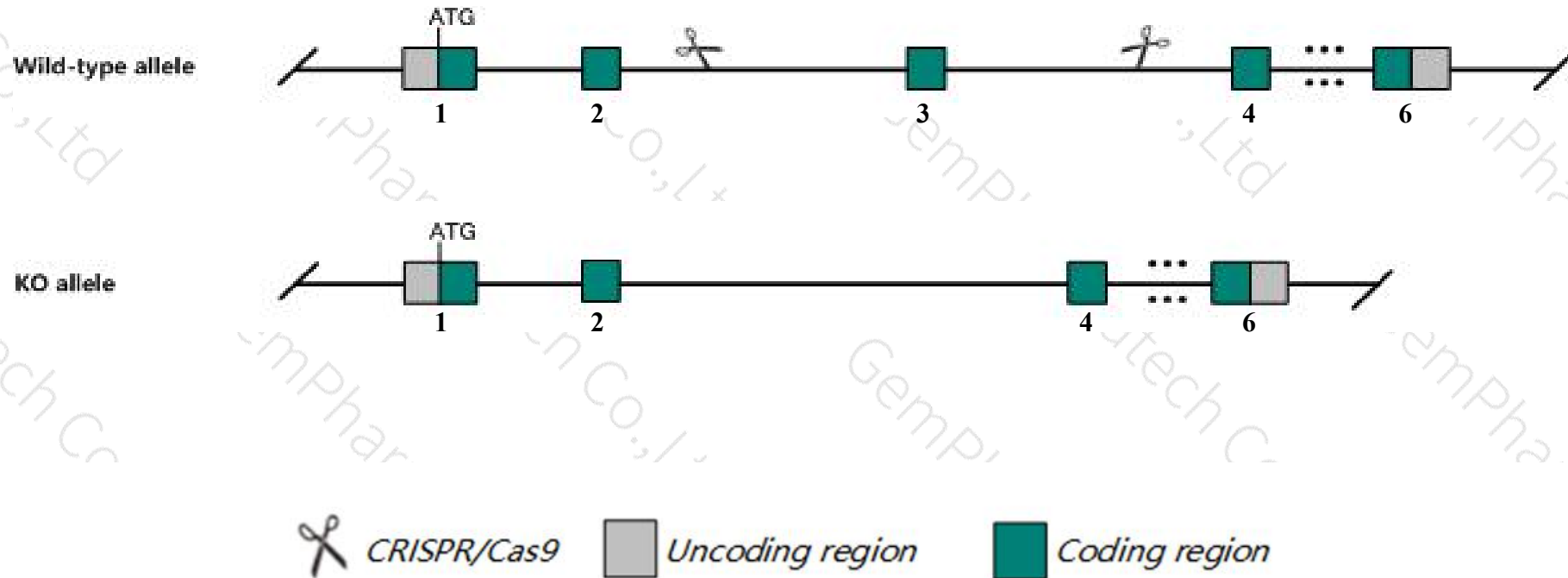
**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Fads6* gene. The schematic diagram is as follows:



- The *Fads6* gene has 2 transcripts. According to the structure of *Fads6* gene, exon3 of *Fads6-201* (ENSMUST00000056153.7) transcript is recommended as the knockout region. The region contains 181bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Fads6* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Fads6* gene is located on the Chr11. 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)

## Fads6 fatty acid desaturase domain family, member 6 [ *Mus musculus* (house mouse) ]

Gene ID: 328035, updated on 7-Dec-2019

### Summary

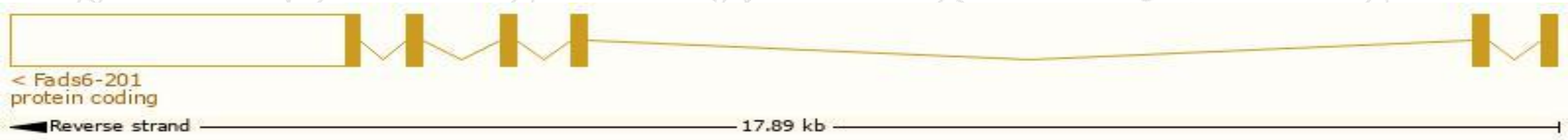
Official Symbol	Fads6 provided by <a href="#">MGI</a>
Official Full Name	fatty acid desaturase domain family, member 6 provided by <a href="#">MGI</a>
Primary source	<a href="#">MGI:MGI:3039592</a>
See related	<a href="#">Ensembl:ENSMUSG00000044788</a>
Gene type	protein coding
RefSeq status	VALIDATED
Organism	<a href="#">Mus musculus</a>
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	BC044804
Expression	Biased expression in adrenal adult (RPKM 34.0), liver adult (RPKM 20.1) and 11 other tissues <a href="#">See more</a>
Orthologs	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

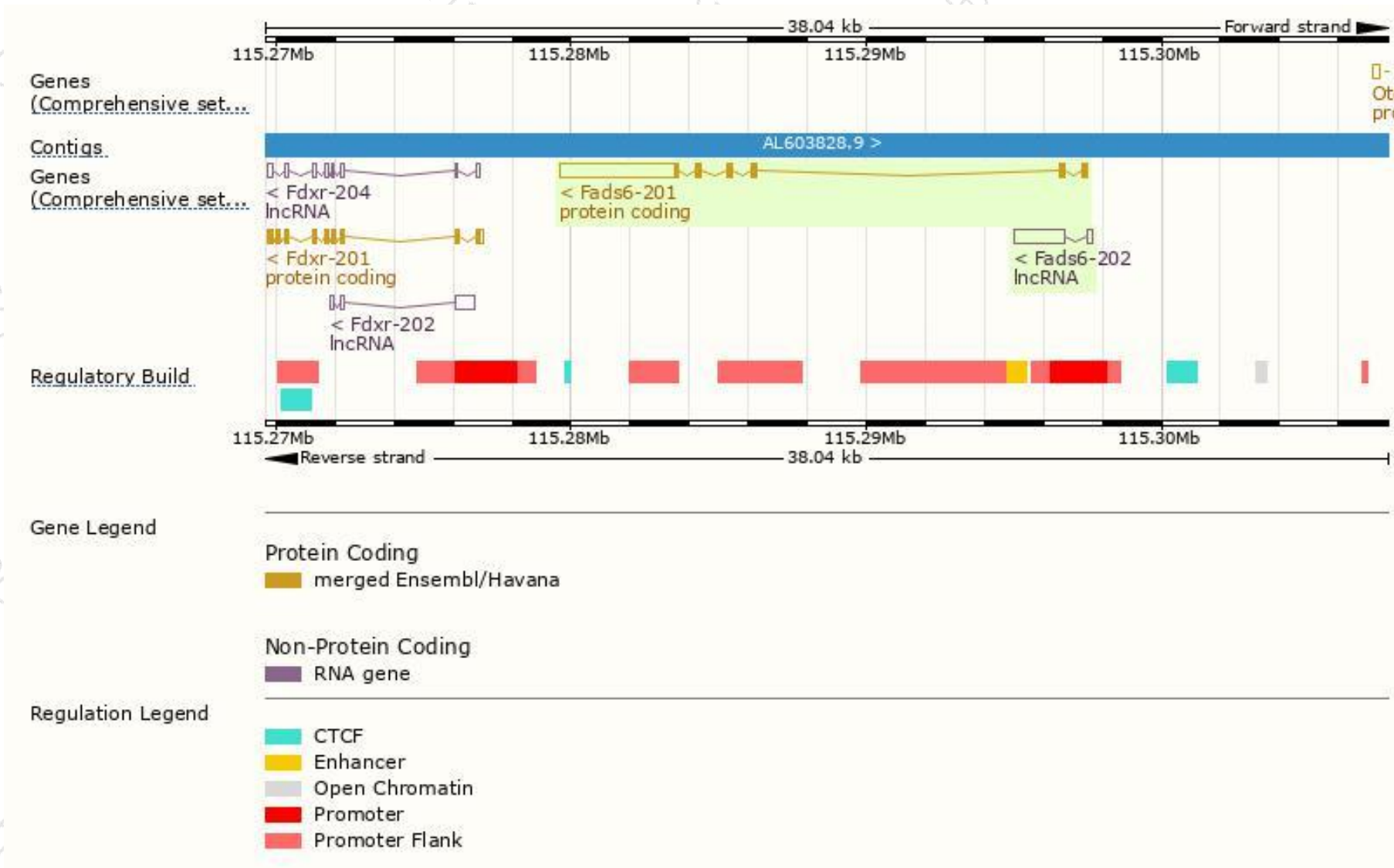
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Fads6-201	<a href="#">ENSMUST00000056153.7</a>	4951	<a href="#">342aa</a>	Protein coding	<a href="#">CCDS25626</a>	<a href="#">Q80UG1</a>	TSL:1 GENCODE basic APPRIS P1
Fads6-202	<a href="#">ENSMUST00000141006.1</a>	1890	No protein	lncRNA	-	-	TSL:1

The strategy is based on the design of *Fads6-201* transcript,The transcription is shown below

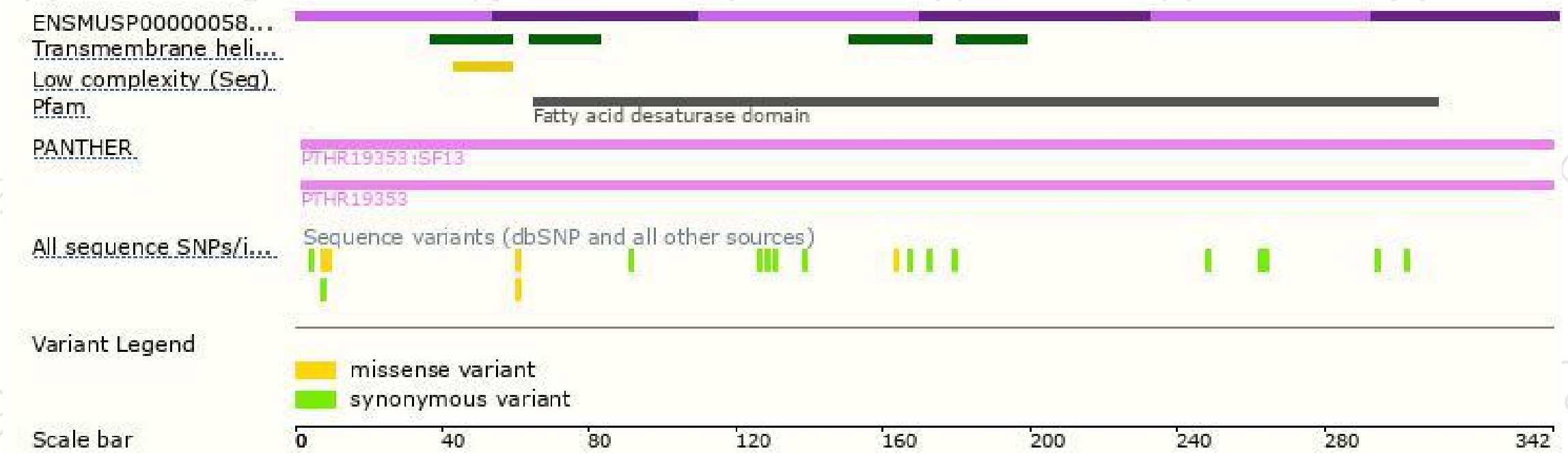


# Genomic location distribution





# Protein domain



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

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