

Fzd6 Cas9-KO Strategy

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Project Overview

Project Name

Fzd6

Project type

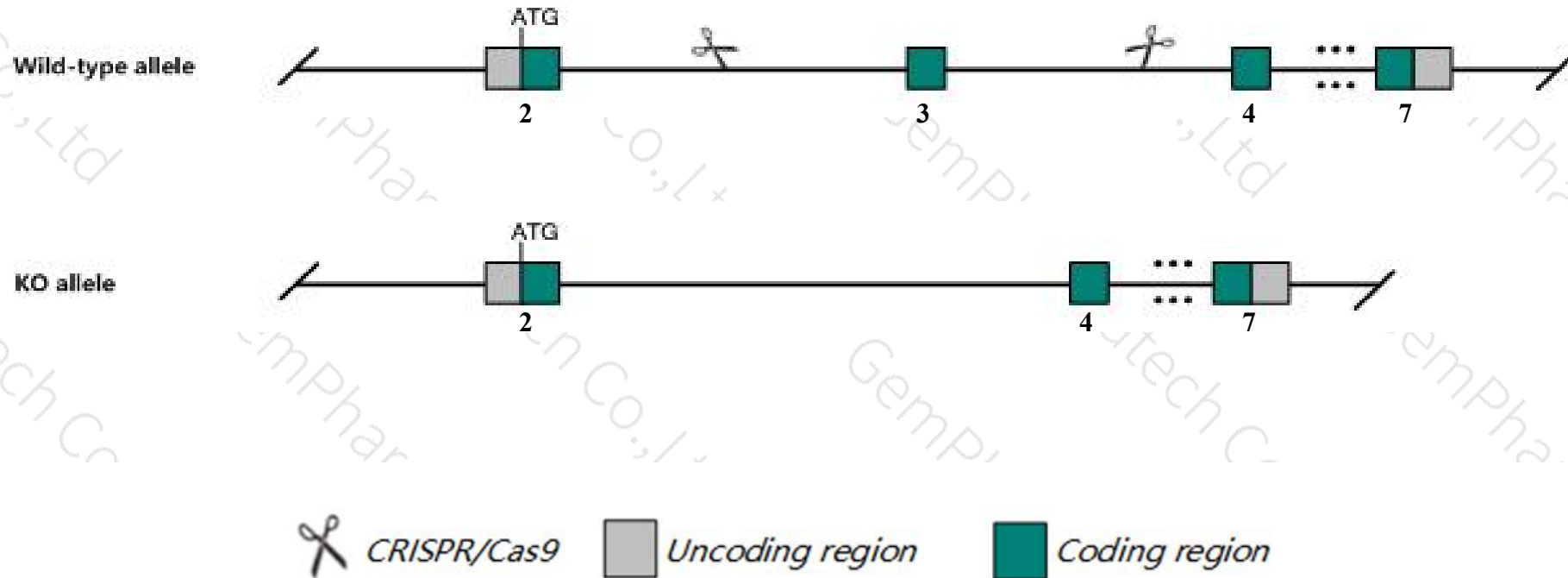
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Fzd6* gene. The schematic diagram is as follows:



- The *Fzd6* gene has 5 transcripts. According to the structure of *Fzd6* gene, exon3 of *Fzd6-202* (ENSMUST00000179165.8) transcript is recommended as the knockout region. The region contains 197bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Fzd6* gene. The brief process is as follows: CRISPR/Cas9 system v

- According to the existing MGI data, Homozygous mice for one mutation display abnormal hair follicle orientation. Another mutation of this gene does not appear to result in a phenotype.
- The *Fzd6* gene is located on the Chr15. 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)

Fzd6 frizzled class receptor 6 [Mus musculus (house mouse)]

Gene ID: 14368, updated on 19-Feb-2019

Summary



Official Symbol	Fzd6 provided by MGI
Official Full Name	frizzled class receptor 6 provided by MGI
Primary source	MGI:MGI:108474
See related	Ensembl:ENSMUSG00000022297
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	Fz6
Expression	Ubiquitous expression in bladder adult (RPKM 8.3), limb E14.5 (RPKM 5.7) and 26 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

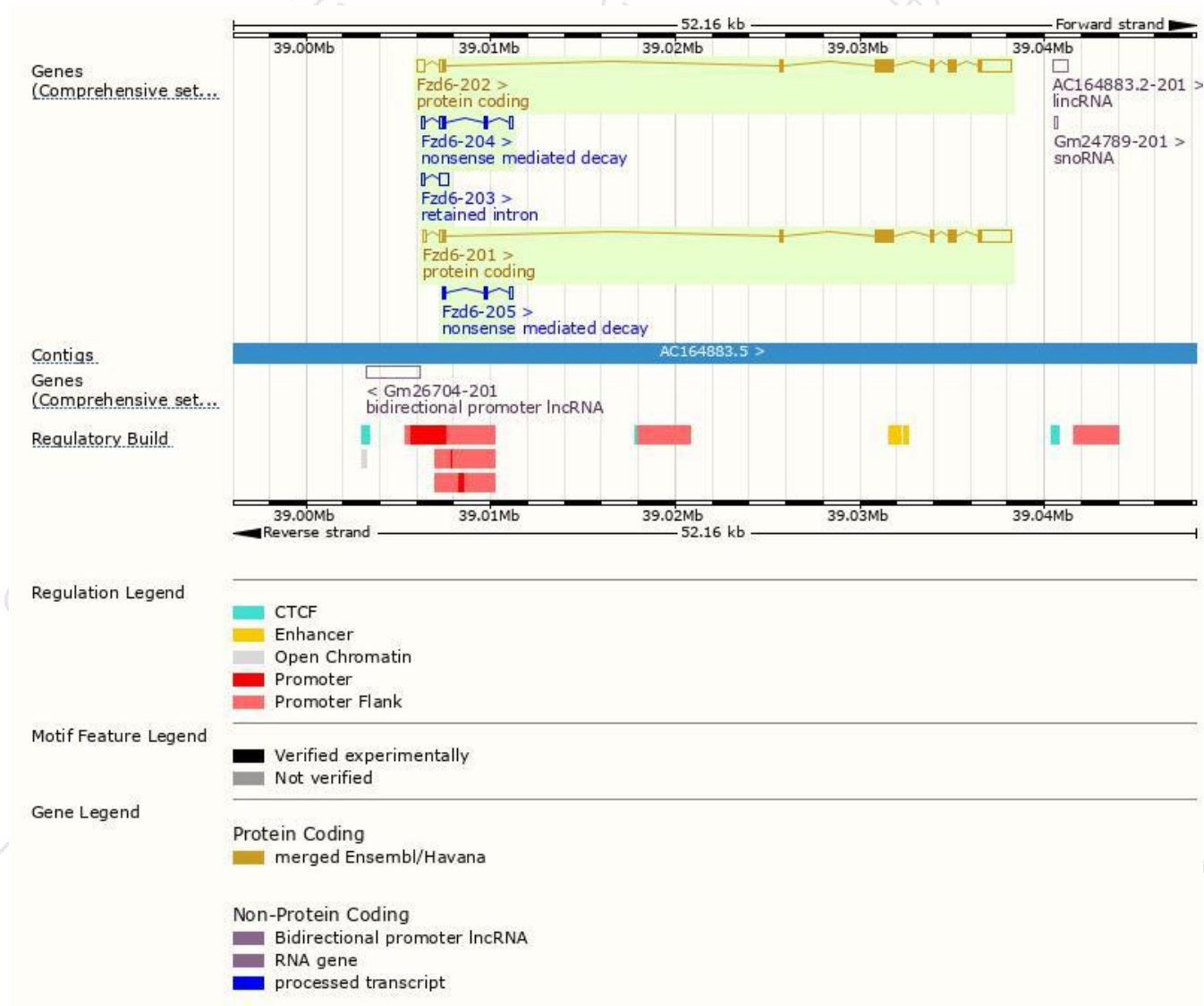
The gene has 5 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Fzd6-202	ENSMUST00000179165.8	4325	709aa	Protein coding	CCDS27441	Q542J1 Q61089	TSL:1 GENCODE basic APPRIS P1
Fzd6-201	ENSMUST00000022906.7	4081	709aa	Protein coding	CCDS27441	Q542J1 Q61089	TSL:1 GENCODE basic APPRIS P1
Fzd6-204	ENSMUST00000228491.1	750	68aa	Nonsense mediated decay	-	A0A2I3BQF8	
Fzd6-205	ENSMUST00000228922.1	458	80aa	Nonsense mediated decay	-	A0A2I3BR53	
Fzd6-203	ENSMUST00000227606.1	647	No protein	Retained intron	-	-	

The strategy is based on the design of *Fzd6-202* transcript,The transcription is shown below



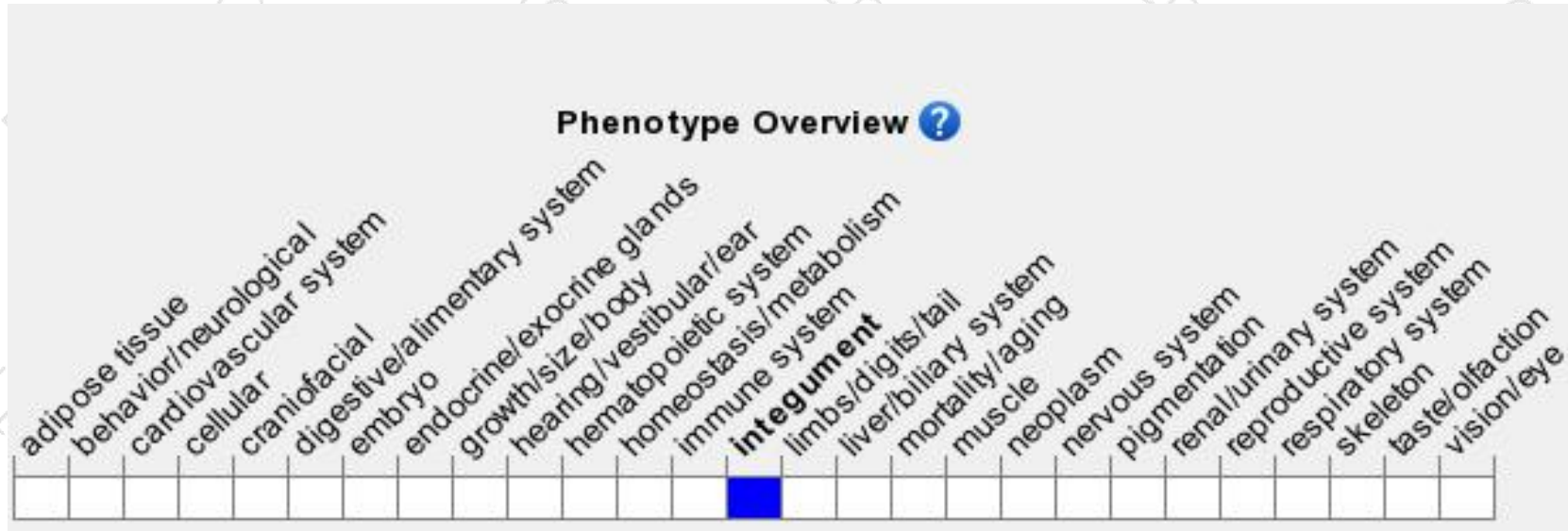
Genomic location distribution



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 mice for one mutation display abnormal hair follicle orientation. Another mutation of this gene does not appear to result in a phenotype.

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

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