

Zdhhc19-C142S cas9-ki(PM) Mouse Model Strategy

-CRISPR/Cas9 technology

Designer: Xueting Zhang

Reviewer: Yanhua Shen

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Project Overview

Project Name

Zdhhc19-C142S

Project type

cas9-ki(PM)

Strain background

C57BL/6JGpt

Technical Description

- The mouse *Zdhhc19* gene has 5 transcripts.
- According to the structure of *Zdhhc19* gene and requirements of customer, this project produced *Zdhhc19*-C142S point mutation on exon5 of the transcript of *Zdhhc19*-201(ENSMUST00000064192.8, NM_199309.2). The 142th amino acids will be mutated from C to S, and the corresponding codon will be mutated to AGC by the TGC.
- The mouse *Zdhhc19*-201 transcript contains 9 exons. The translation initiation site ATG is located at exon2, and the translation termination site TAG is located at exon9, encoding 347aa.
- In this project, *Zdhhc19* gene will be modified by CRISPR/Cas9 technology. The brief process is as follows: In vitro, sgRNA and donor vectors were constructed. Cas9, sgRNA and donor were injected into the fertilized eggs of C57BL/6JGpt mice for homologous recombination, and obtained positive F0 mice identified by PCR and sequencing analysis. The stable inheritable positive F1 mice model was obtained by mating F0 mice with C57BL/6JGpt mice.

Targeted Mutation Site

Before mutation

+3		D	F	D	H	H	C	K	W	V	N	N	C	I	G	H	R	N	F	R	L	F	M	L	L	V	L	S	L	C	L
12701	CCTGTAACAG	GACTTCGACC	ACCAT	TGCAA	GTGGGTCAAT	AACTGCATCG	GTCACCGCAA	CTTCCGCCTC	TTCATGCTGC	TGGTCCTGTC	ACTCTGTCTC	GGACATTGTC	CTGAAGCTGG	TGGTAACGTT	CACCCAGTTA	TTGACGTAGC	CAGTGGCGTT	GAAGGCGGAG	AAGTACGACG	ACCAGGACAG	TGAGACAGAG										
+3		Y	S	G	A	L	L	V	T	C	L	T	F	L	F	R	T	R	H	L	P	F	S	L	D	K	G	M			
12801	TACTCGGGAG	CCCTGCTGGT	CACCTGCCTG	ACATTCTAT	TTGCGACAAG	GCATCTGCC	TTCTCCCTGG	ACAAGGGGAT	GGCGTATCTA	CTTATGCGCG	ATGAGCCCTC	GGGACGACCA	GTGGACGGAC	TGTAAGGATA	AAGCGTGTTC	CGTAGACGGG	AAGAGGGACC	TGTTCCCCTA	CCGCATAGAT	GAATACGCGC											

After mutation

+3		D	F	D	H	H	S	K	W	V	N	N	C	I	G	H	R	N	F	R	L	F	M	L	L	V	L	S	L	C	L
12701	CCTGTAACAG	GACTTCGACC	ACCATAGCAA	GTGGGTCAAT	AACTGCATCG	GTCACCGCAA	CTTCGCCTC	TTCATGCTGC	TGGTCCTGTC	ACTCTGTCTC	GGACATTGTC	CTGAAGCTGG	TGGTATCGTT	CACCCAGTTA	TTGACGTAGC	CAGTGGCGTT	GAAGGCGGAG	AAGTACGACG	ACCAGGACAG	TGAGACAGAG											
+3		Y	S	G	A	L	L	V	T	C	L	T	F	L	F	R	T	R	H	L	P	F	S	L	D	K	G	M			
12801	TACTCGGGAG	CCCTGCTGGT	CACCTGCCTG	ACATTCTAT	TTGCGACAAG	GCATCTGCCC	TTCTCCCTGG	ACAAGGGGAT	GGCGTATCTA	CTTATGCGCG	ATGAGCCCTC	GGGACGACCA	GTGGACGGAC	TGTAAGGATA	AAGCGTGTTC	CGTAGACGGG	AAGAGGGACC	TGTTCCCCTA	CCGCATAGAT	GAATACGCGC											

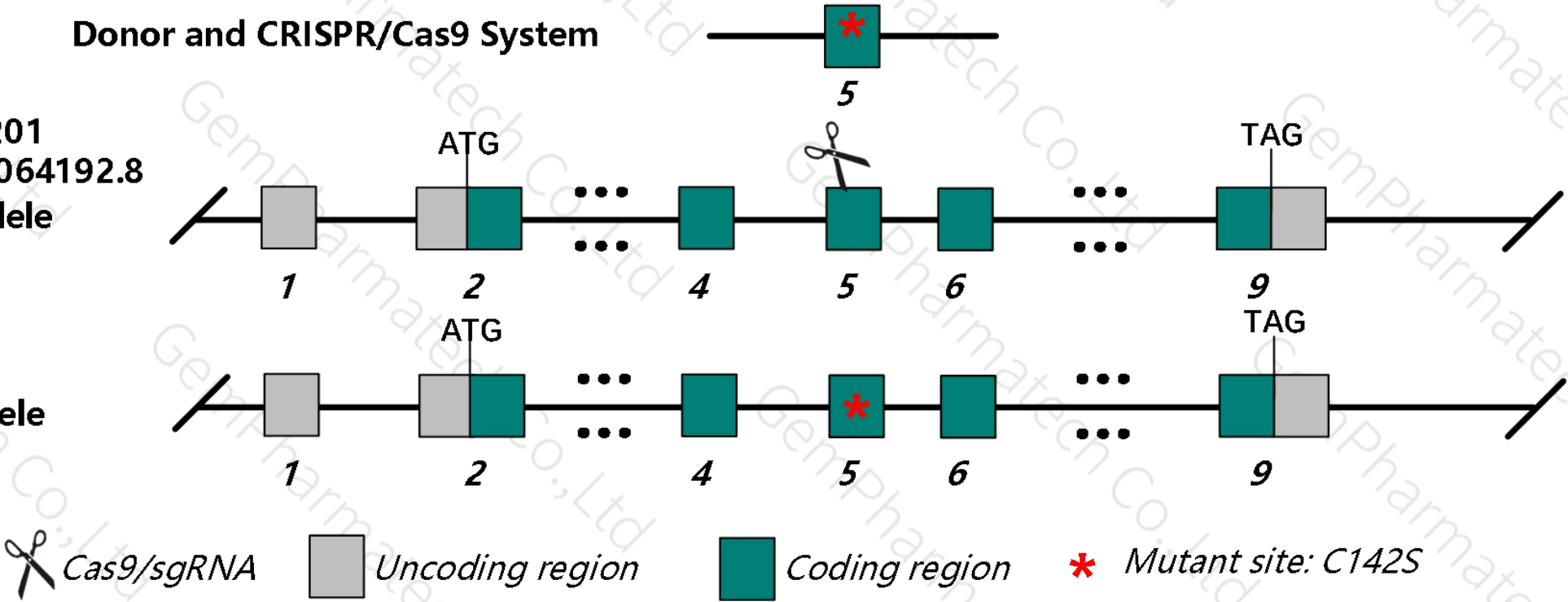
The blue region is exon5 of *Zdhhc19-201*, the yellow region represents the targeted mutation site.

This model uses CRISPR/Cas9 technology to edit the *Zdhhc19* gene and the schematic diagram is as follow:

Donor and CRISPR/Cas9 System

Zdhhc19-201
ENSMUST00000064192.8
Wild-type allele

Targeted allele



- One or Two synonymous mutations of amino acids will be introduced on exon5 of *Zdhhc19*.
- Mouse *Zdhhc19* gene is located on Chr16. Please take the loci in consideration when breeding this mutation mice with other gene modified strains, if the other gene is also on Chr16, it may be extremely hard to get double gene positive homozygotes.
- The scheme is designed according to the genetic information in the existing database. Due to the complex process of gene transcription and translation, it cannot be predicted completely at the present technology level.

Gene name and location (NCBI)

Zdhhc19 zinc finger, DHHC domain containing 19 [*Mus musculus* (house mouse)]

Download Datasets

Gene ID: 245308, updated on 17-Dec-2020

Summary

- Official Symbol

Zdhhc19 provided by MGI
- Official Full Name

zinc finger, DHHC domain containing 19 provided by MGI
- Primary source

MGI:MGI:2682948
- See related

Ensembl:ENSMUSG00000052363
- Gene type

protein coding
- RefSeq status

PROVISIONAL
- Organism

Mus musculus
- Lineage

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
- Also known as

Gm616; Gm1744
- Expression

Restricted expression toward testis adult (RPKM 76.6) See more
- Orthologs

human all
- NEW

Try the new Gene table
- Try the new Transcript table

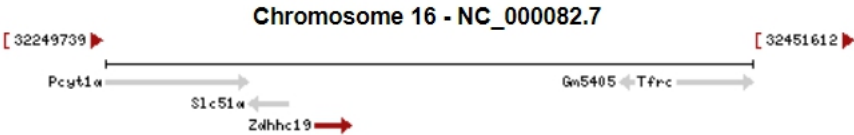
Genomic context

Location: 16; 16 B3

See Zdhhc19 in Genome Data Viewer

Exon count: 12

Annotation release	Status	Assembly	Chr	Location
109	current	GRCm39 (GCF_000001635.27)	16	NC_000082.7 (32315059..32326032)
108.20200622	previous assembly	GRCm38.p6 (GCF_000001635.26)	16	NC_000082.6 (32496241..32507214)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	16	NC_000082.5 (32496367..32507300)



Transcript information (Ensembl)

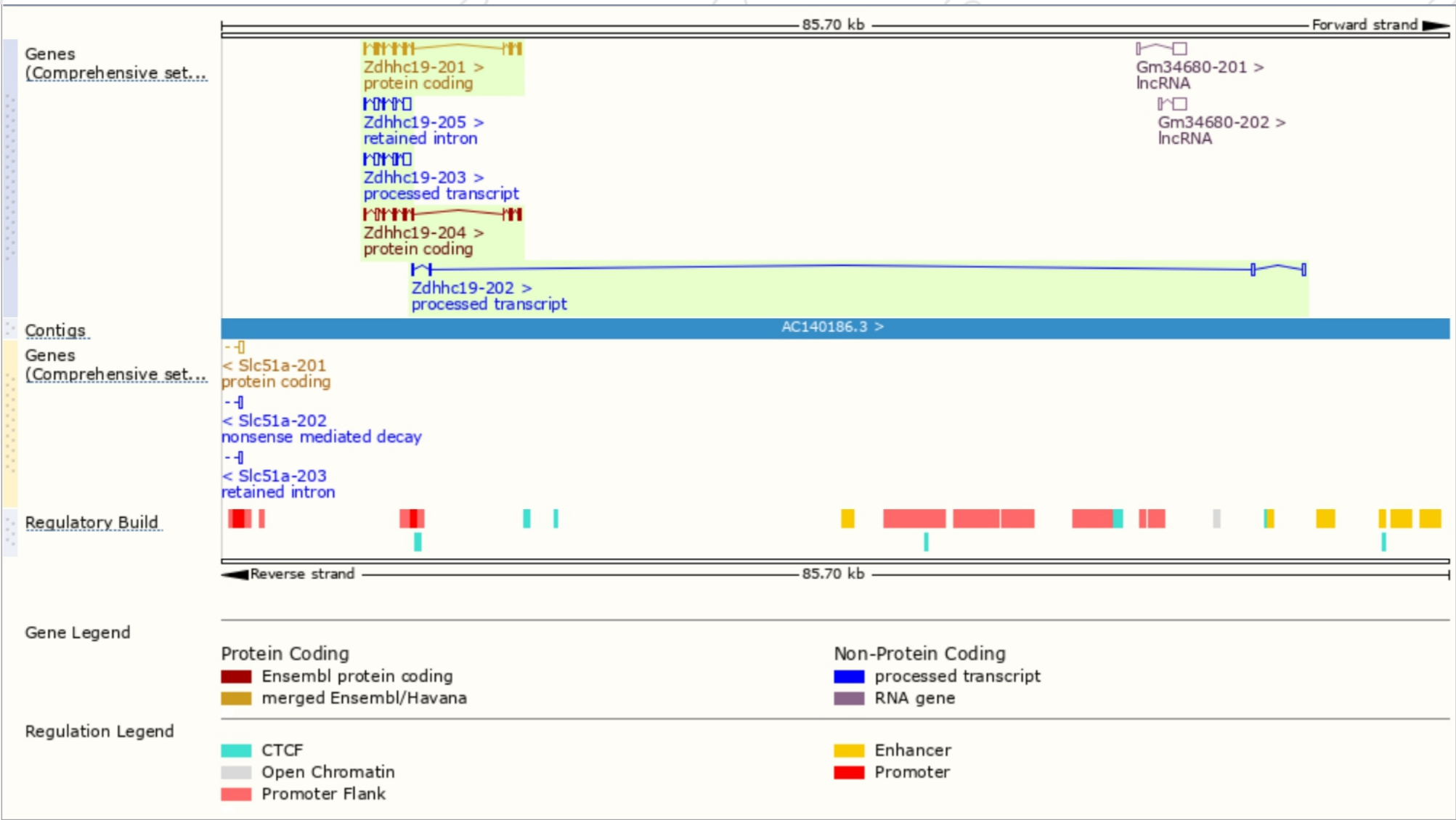
The gene has 5 transcripts, and all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt Match	Flags
Zdhhc19-201	ENSMUST00000064192.8	1323	347aa	Protein coding	CCDS28122	Q810M5	TSL:1 Gencode basic APPRIS P2
Zdhhc19-204	ENSMUST00000231510.2	1181	251aa	Protein coding	-	A0A338P7I3	Gencode basic APPRIS ALT2
Zdhhc19-203	ENSMUST00000231293.2	1098	No protein	Processed transcript	-	-	-
Zdhhc19-202	ENSMUST00000160832.2	631	No protein	Processed transcript	-	-	TSL:3
Zdhhc19-205	ENSMUST00000232434.2	1067	No protein	Retained intron	-	-	-

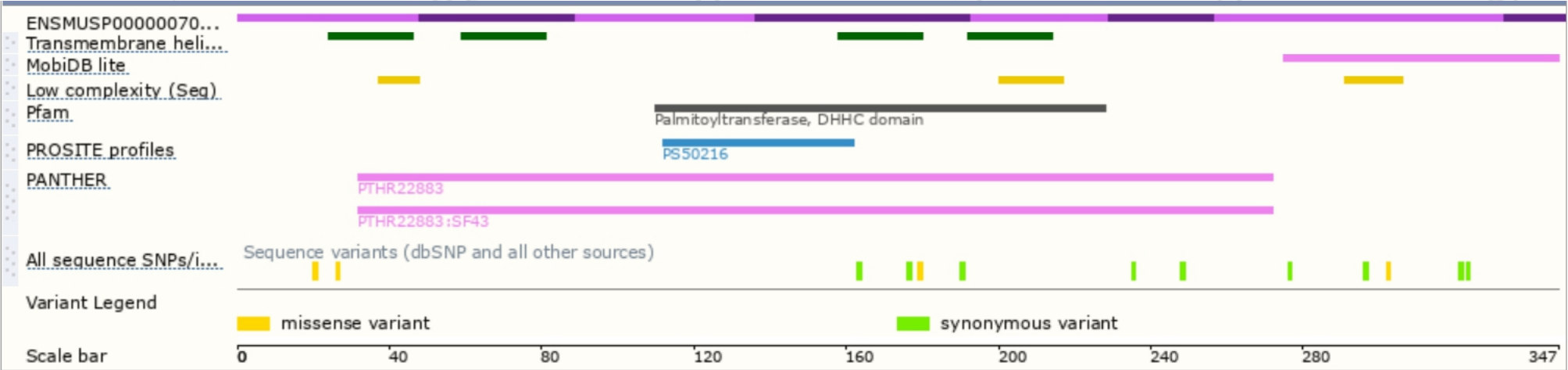
The strategy is based on the design of *Zdhhc19-201* transcript, the transcription is shown below:



Genomic location distribution



Protein domain



If you have any questions, please feel free to contact us.

Tel: 025-5864 1534



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