Otof-L1010P Mouse Model Strategy -CRISPR/Cas9 technology

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Reviewer: Xueting Zhang

Design Date: 2021-1-12

Project Overview



Project Name

Otof-L1010P

Project type

Cas9-ki(PM)

Strain background

C57BL/6JGpt

Technical Description



- The mouse *Otof* gene has 5 transcripts. The human *OTOF* gene has 7 transcripts.
- According to the structure of *Otof* gene and requirements of customer, the 1011th amino acid(Leu) of human *OTOF* gene corresponds to the 1010th amino acid(Leu) of mouse *Otof* gene after comparing homology of mouse *Otof* gene and human *Otof* gene. This project produced *Otof*-L1010P point mutation on exon25 of the transcript of *Otof*-201(ENSMUST00000074171.9, NM_031875.2). The 1010th amino acids will be mutated from L(Leu) to P(Pro), and the corresponding codon will be mutated to CCG by the CTG.
- The mouse *Otof*-201 transcript contains 47 exons. The translation initiation site ATG is located at exon1, and the translation termination site TGA is located at exon46, encoding 1997aa.
- In this project, *Otof* 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.



A comparison of the aa homology of human and mouse Otof gene

	976	980	990	1000 10	10	1020 1030	1040	1050	1060
Human OTOF protein			VFFINQSQC	CTEVLNETLCPTWDQML	VFDNLEI	YGEAHELRDDPPIIVIE	IYDQDSMGKAD	FMGRTFAKE	LVKMADEAYC
Mouse Otof protein	975	DSSGLSDPFAR	VFFINQSQC	TEVLNETLCPTWDQML	VFDNLEI	YGEAHELRDDPPIIVIE	IYDQDSMGKAD)FMGRTFAKP	LVKMADEAYC
1									>
				muta	nt sit	e			
Consensus	076	DSSCISDDFAD	/FFINOSOC	TEVI NETI COTWOOMI	VEDNIEI	YGEAHELRDDPPIIVIE	TVDODSMCVAT	NEMCDTEN VD	UWMA DEA VC
Consensus	370	1	VEE THOSE	, LEVENETECPT WDQFIL	VEDNLEI	J	TIDQUSHGRAL	PERIORI PARE	LVRIADEATC
	To: 0.1			consensus position	s: 97.2%	identity positions: 94.8	% aln: 981		
			T.	N O		* / *,			

The 1011th amino acid(Leu) of human *Otof* gene corresponds to the 1010th amino acid(Leu) of mouse *Otof* gene after comparing homology of mouse *Otof* gene and human *OTOF* gene.

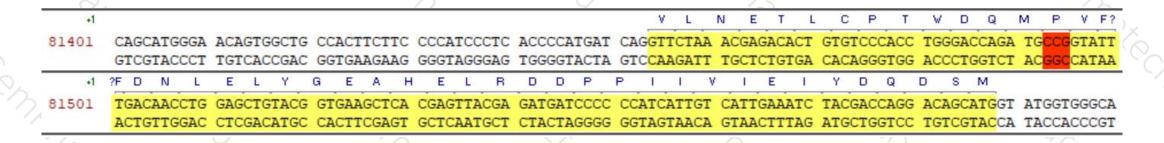
Mutation Site



Before mutation

-1																	٧	L	N	Е	Т	L	С	Р	Т	W	D	Q	М	L	V F	?
81401	CAGCATO	GGA	ACA	GTG	GCTG	CC	ACT	TCTT	C	CCCA	TCC	CTC	ACC	CCA	TGAT	CAG	GTT	CTA	A A	GAG	ACZ	CT	GTG	TCC	CACC	TG	GGA	CAGA	TG	CTG	GTAT	Т
02102	GTCGTAC																															
+1	?F D N	L	E	L	Υ	G	Е	Α	Н	Е	L	R	D	D	Р	Р	I	ı	٧	1	E	1	Y	D	Q	D	S	М				
81501	TGACAAC	CTG	GAG	CTG	TACG	GT	GAA	GCTC	A	CGAG	TTA	CGA	GAT	GAT	cccc	CCA	TCA	TTG	r cz	ATTO	SAAZ	ATC	TAC	GAC	CAGG	AC	AGC	ATGGI	AT	GGT	GGGC	A
	ACTGTTG	GAC	CTC	GAC	ATGC	CA	CTT	CGAG	T	GCTC	AAT	GCT	CTA	CTA	GGGG	GGT	AGT	AAC	A G	CAAC	TTT	rag	ATG	CTG	GTCC	TG	TCG	racca	TA	CCA	CCCG	Γ

After mutation

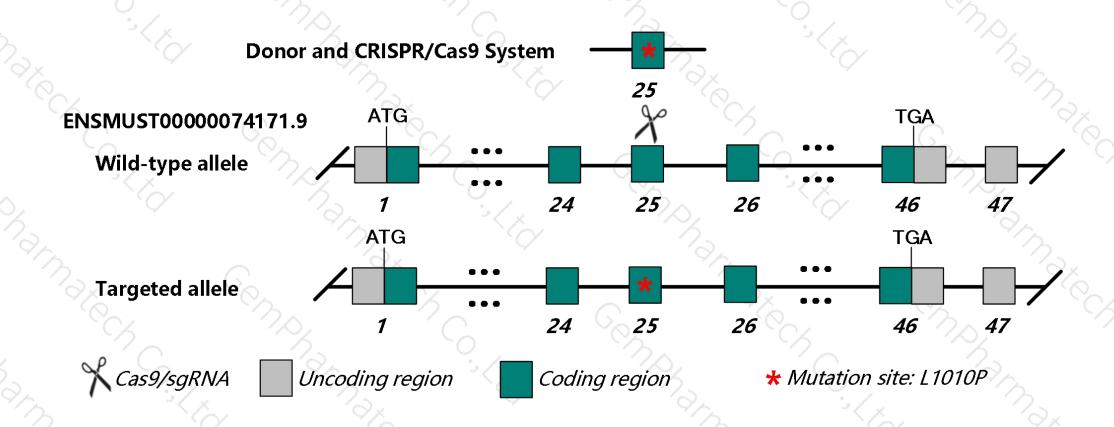


The yellow region is exon25 of Otof-201, the red region represents the mutation site.

Strategy



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



Notice



- According to the data of MGI, Homozygous mutants have no detectable auditory brainstem response at any frequency tested. Otoacoustic transmission distortion products are detected. Direct electrical stimulation of cochlear ganglia elicits brainstem responses. On depolarization, inner hair cells release almost no neurotransmitter.
- > One or Two synonymous mutations of amino acids will be intronduced on exon25 of *Otof*.
- Mouse *Otof* gene is located on Chr5. Please take the loci in consideration when breeding this mutation mice with other gene modified strains, if the other gene is also on Chr5, 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)



Otof otoferlin [Mus musculus (house mouse)]

Gene ID: 83762, updated on 17-Dec-2020

Summary

☆ ?

Official Symbol Otof provided by MGI
Official Full Name otoferlin provided by MGI
Primary source MGI:MGI:1891247

See related Ensembl: ENSMUSG00000062372

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

Expression Biased expression in CNS E18 (RPKM 4.0), cortex adult (RPKM 3.4) and 9 other tissues See more

Orthologs <u>human</u> all



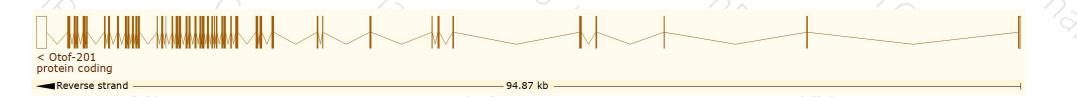
Transcript information (Ensembl)



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

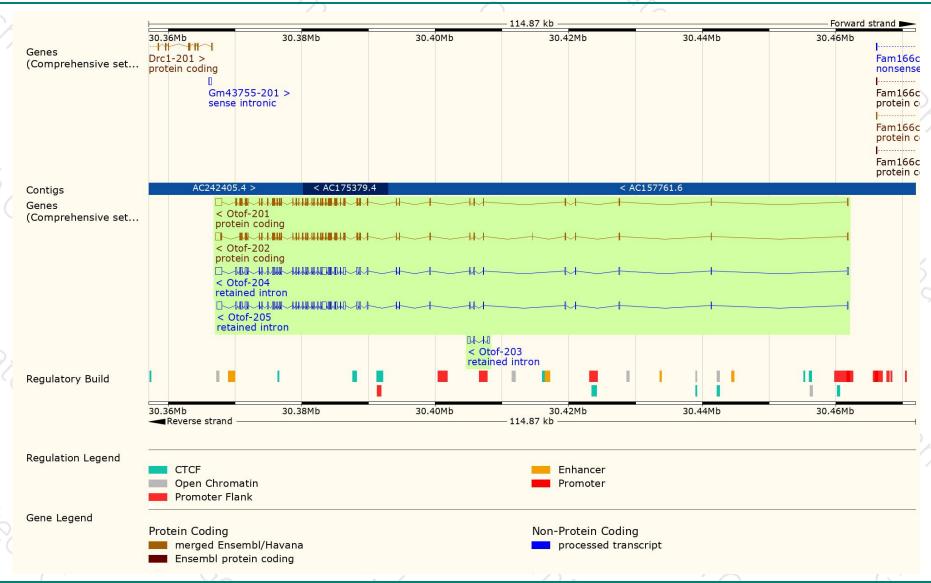
					<u> </u>								
Name A	Transcript ID 👙	bp 👙	Protein	Biotype 👙	CCDS 🍦	UniProt Match	Flags						
Otof-201	ENSMUST00000074171.9	7129	1997aa	Protein coding	CCDS19157&	E9PYR6@	TSL:1 GENCODE basic APPRIS P1						
Otof-202	ENSMUST00000114747.8	6881	<u>1992aa</u>	Protein coding	CCDS51452&	A0A0R4J1K2&	TSL:1 GENCODE basic						
Otof-203	ENSMUST00000128177.1	686	No protein	Retained intron		-	TSL:2						
Otof-204	ENSMUST00000144125.7	7419	No protein	Retained intron	-	-	TSL:2						
Otof-205	ENSMUST00000150734.7	7063	No protein	Retained intron	2	-	TSL:1						

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



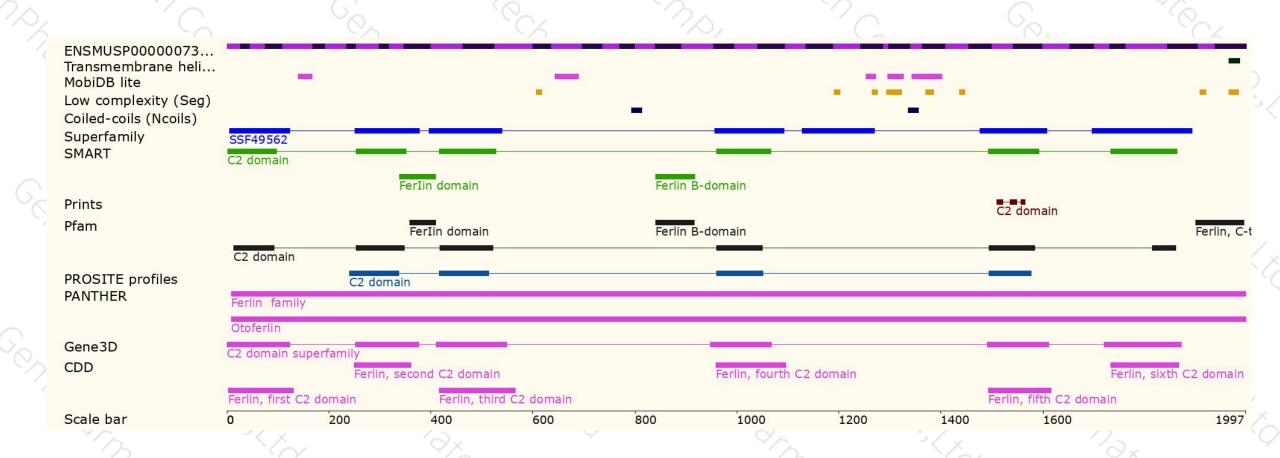
Genomic location distribution





Protein domain



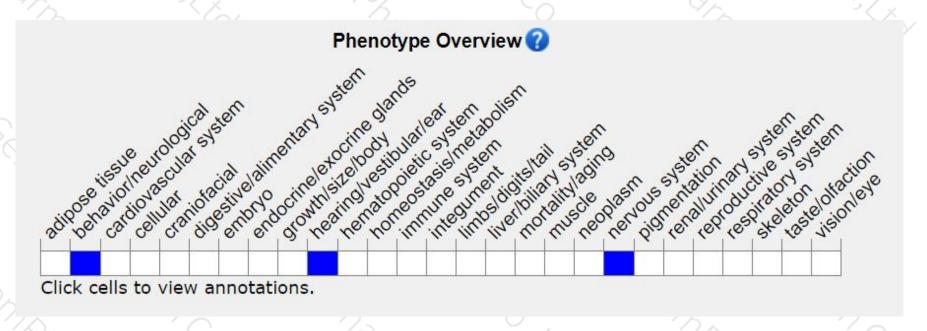


Mouse phenotype description(MGI)



URL link is as follows:

http://www.informatics.jax.org/marker/MGI:1891247



Homozygous mutants have no detectable auditory brainstem response at any frequency tested. Otoacoustic transmission distortion products are detected. Direct electrical stimulation of cochlear ganglia elicits brainstem responses. On depolarization, inner hair cells release almost no neurotransmitter.

If you have any questions, please feel free to contact us. Tel: 025-5864 1534





