

Cers4 Cas9-CKO Strategy

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Project Overview



Project Name Cers4

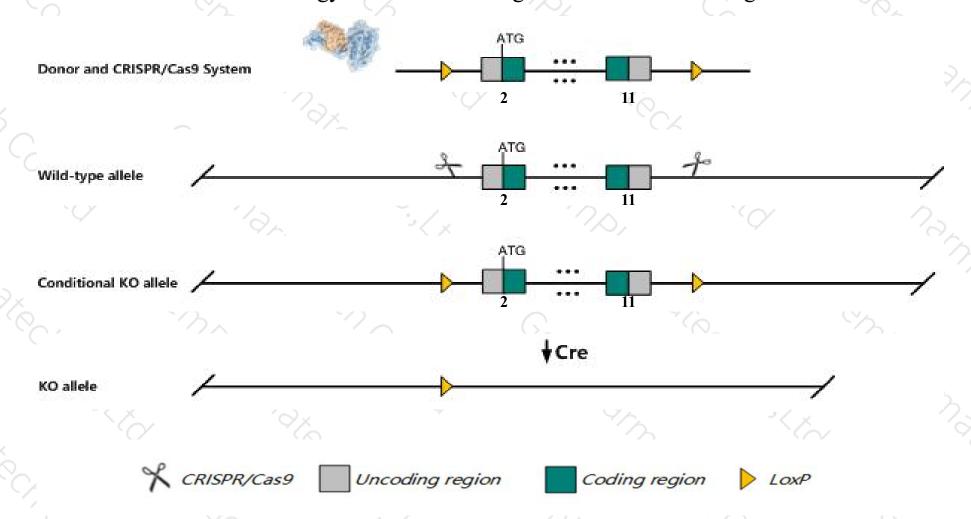
Project type Cas9-CKO

Strain background C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Cers4 gene. The schematic diagram is as follows:



Technical routes



- The *Cers4* gene has 9 transcripts. According to the structure of *Cers4* gene, exon2-exon11 of *Cers4-201* (ENSMUST00000008350.15) transcript is recommended as the knockout region. The region contains all of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Cers4* gene. The brief process is as follows:CRISPR/Cas9 system and Donor were microinjected into the fertilized eggs of C57BL/6JGpt mice. Fertilized eggs were transplanted to obtain positive F0 mice which were confirmed by PCR and sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6JGpt mice.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

Notice



- > According to the existing MGI data, Mice homozygous for a knock-out allele exhibit altered lipid composition of the sebum and hair follicle dystrophy that results in a progressive form of alopecia.
- > The *Cers4* gene is located on the Chr8. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)



Cers4 ceramide synthase 4 [Mus musculus (house mouse)]

Gene ID: 67260, updated on 31-Jan-2019

Summary

☆ ?

Official Symbol Cers4 provided by MGI

Official Full Name ceramide synthase 4 provided by MGI

Primary source MGI:MGI:1914510

See related Ensembl:ENSMUSG00000008206

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 2900019C14Rik, Lass4, Trh1

Expression Ubiquitous expression in genital fat pad adult (RPKM 11.0), lung adult (RPKM 8.8) and 23 other tissues See more

Orthologs <u>human all</u>

Transcript information (Ensembl)



The gene has 9 transcripts, all transcripts are shown below:

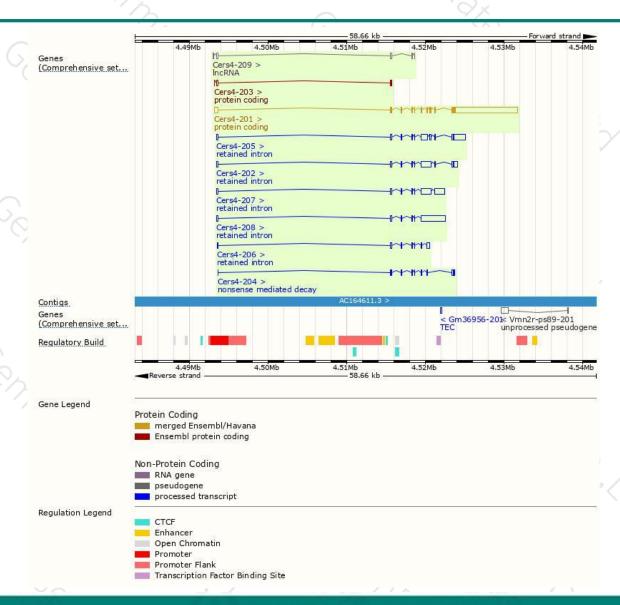
7 No.				/)		
Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
ENSMUST00000008350.15	9475	<u>393aa</u>	Protein coding	CCDS22086	Q9D6J1	TSL:1 GENCODE basic APPRIS P1
ENSMUST00000176042.1	387	<u>56aa</u>	Protein coding	-	H3BKZ9	CDS 3' incomplete TSL:3
ENSMUST00000176130.9	874	<u>211aa</u>	Nonsense mediated decay	828	H3BL50	TSL:5
ENSMUST00000176932.7	3674	No protein	Retained intron	3528	20	TSL:2
ENSMUST00000176267.8	3369	No protein	Retained intron	(27)	=	TSL:2
ENSMUST00000176837.7	3116	No protein	Retained intron	-	* .	TSL:1
ENSMUST00000175781.4	2539	No protein	Retained intron	825	=	TSL:1
ENSMUST00000176705.1	908	No protein	Retained intron	1528	20	TSL:2
ENSMUST00000177010.9	561	No protein	IncRNA	-	· ·	TSL:5
	ENSMUST00000176042.1 ENSMUST00000176042.1 ENSMUST00000176130.9 ENSMUST00000176932.7 ENSMUST00000176267.8 ENSMUST00000176837.7 ENSMUST00000175781.4 ENSMUST00000176705.1	ENSMUST00000176042.1 387 ENSMUST00000176042.1 387 ENSMUST00000176130.9 874 ENSMUST00000176932.7 3674 ENSMUST00000176267.8 3369 ENSMUST00000176837.7 3116 ENSMUST00000175781.4 2539 ENSMUST00000176705.1 908	ENSMUST00000008350.15 9475 393aa ENSMUST00000176042.1 387 56aa ENSMUST00000176130.9 874 211aa ENSMUST00000176932.7 3674 No protein ENSMUST00000176267.8 3369 No protein ENSMUST00000176837.7 3116 No protein ENSMUST00000175781.4 2539 No protein ENSMUST00000176705.1 908 No protein	ENSMUST00000008350.15 9475 393aa Protein coding ENSMUST00000176042.1 387 56aa Protein coding ENSMUST00000176130.9 874 211aa Nonsense mediated decay ENSMUST00000176932.7 3674 No protein Retained intron ENSMUST00000176267.8 3369 No protein Retained intron ENSMUST00000176837.7 3116 No protein Retained intron ENSMUST00000175781.4 2539 No protein Retained intron ENSMUST00000176705.1 908 No protein Retained intron	ENSMUST00000008350.15 9475 393aa Protein coding CCDS22086 ENSMUST00000176042.1 387 56aa Protein coding - ENSMUST00000176130.9 874 211aa Nonsense mediated decay - ENSMUST00000176932.7 3674 No protein Retained intron - ENSMUST00000176267.8 3369 No protein Retained intron - ENSMUST00000176837.7 3116 No protein Retained intron - ENSMUST00000175781.4 2539 No protein Retained intron - ENSMUST00000176705.1 908 No protein Retained intron -	ENSMUST00000008350.15 9475 393aa Protein coding CCDS22086 Q9D6J1 ENSMUST00000176042.1 387 56aa Protein coding - H3BKZ9 ENSMUST00000176130.9 874 211aa Nonsense mediated decay - H3BL50 ENSMUST00000176932.7 3674 No protein Retained intron - - ENSMUST00000176267.8 3369 No protein Retained intron - - ENSMUST00000176837.7 3116 No protein Retained intron - - ENSMUST00000175781.4 2539 No protein Retained intron - - ENSMUST00000176705.1 908 No protein Retained intron - -

The strategy is based on the design of Cers4-201 transcript, The transcription is shown below



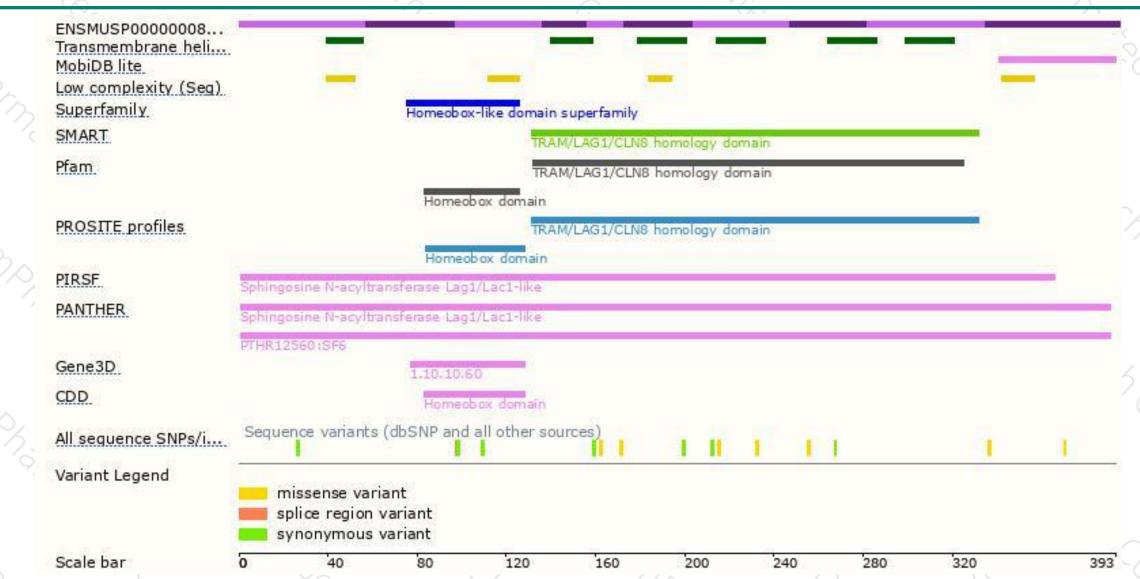
Genomic location distribution





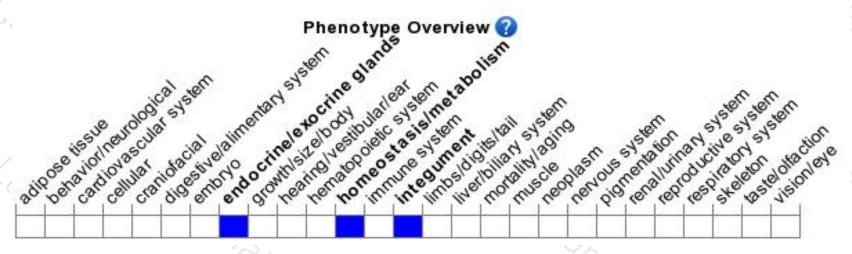
Protein domain





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, Mice homozygous for a knock-out allele exhibit altered lipid composition of the sebum and hair follicle dystrophy that results in a progressive form of alopecia.



If you have any questions, you are welcome to inquire. Tel: 400-9660890





