

Dolar Day Co. Col4al Cas9-KO Strategy To hall alto color color

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Project Overview



Project Name

Col4a1

Project type

Cas9-KO

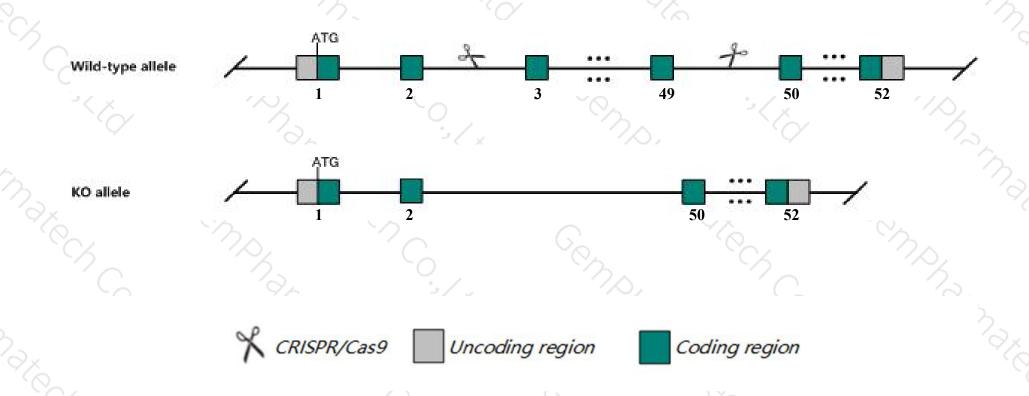
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Col4a1 gene. The schematic diagram is as follows:



Technical routes



- The *Col4a1* gene has 8 transcripts. According to the structure of *Col4a1* gene, exon3-exon49 of *Col4a1-201* (ENSMUST00000033898.9) transcript is recommended as the knockout region. The region contains 4496bp coding sequence Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Col4a1* gene. The brief process is as follows: CRISPR/Cas9 system

Notice



- ➤ According to the existing MGI data, Mice with ENU induced alleles have various eye and vision defects and may show bruising at birth. Mice carrying the G498V mutation have renal glomerular defects that resolve within the first weeks of life, but show retinal tortuosity, muscular dystrophy, brain hemorrhages, and renal cysts as adults.
- The *Col4a1* 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)



Col4a1 collagen, type IV, alpha 1 [Mus musculus (house mouse)]

Gene ID: 12826, updated on 27-Mar-2019

Summary

☆ ?

Official Symbol Col4a1 provided by MGI

Official Full Name collagen, type IV, alpha 1 provided by MGI

Primary source MGI:MGI:88454

See related Ensembl:ENSMUSG00000031502

Gene type protein coding
RefSeq status REVIEWED
Organism Mus musculus

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;

Muroidea; Muridae; Murinae; Mus; Mus

Also known as Bru, Col4a-1, Raw, Svc

Summary This gene encodes the alpha-1 subunit of the type IV collagens, an essential component of basement membranes. The encoded protein

forms a triple helical heterotrimer comprised of two alpha-1 and one alpha-2 subunits that assembles into a type IV collagen network. This gene is located adjacent to the gene encoding alpha-2 subunit. Mice lacking both the alpha-1 and alpha-2 subunits of collagen IV die in utero due to structural deficiencies in the basement membranes and certain mutations in this gene cause perinatal cerebral hemorrhage and

porencephaly. Alternative splicing of this gene results in multiple transcript variants. [provided by RefSeq, Nov 2015]

Expression Broad expression in subcutaneous fat pad adult (RPKM 160.4), lung adult (RPKM 107.4) and 21 other tissuesSee more

Orthologs human all

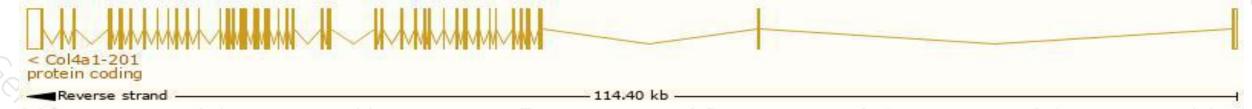
Transcript information (Ensembl)



The gene has 8 transcripts, all transcripts are shown below:

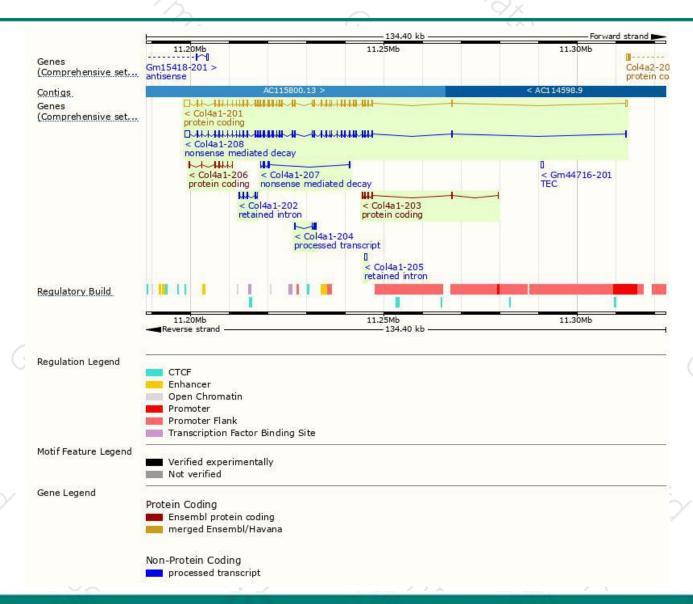
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Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
ENSMUST00000033898.9	6615	<u>1669aa</u>	Protein coding	CCDS40219	P02463	TSL:1 GENCODE basic APPRIS P1
ENSMUST00000209598.1	1170	328aa	Protein coding	8 5	A0A1B0GRC0	CDS 5' incomplete TSL:5
ENSMUST00000208095.1	545	<u>133aa</u>	Protein coding	84	A0A140LHU8	CDS 3' incomplete TSL:3
ENSMUST00000209735.1	6487	<u>1562aa</u>	Nonsense mediated decay	ė.	A0A1B0GSI7	TSL:2
ENSMUST00000209661.1	618	<u>126aa</u>	Nonsense mediated decay	2.5	A0A1B0GT69	CDS 5' incomplete TSL:5
ENSMUST00000208386.1	416	No protein	Processed transcript	8 7	-	TSL:3
ENSMUST00000130488.2	676	No protein	Retained intron	84	ų.	TSL:5
ENSMUST00000209000.1	604	No protein	Retained intron	62	-	TSL:NA
	ENSMUST00000033898.9 ENSMUST00000209598.1 ENSMUST00000208095.1 ENSMUST00000209735.1 ENSMUST00000209661.1 ENSMUST00000208386.1 ENSMUST00000130488.2	ENSMUST00000033898.9 6615 ENSMUST00000209598.1 1170 ENSMUST00000208095.1 545 ENSMUST00000209735.1 6487 ENSMUST00000209661.1 618 ENSMUST00000208386.1 416 ENSMUST00000130488.2 676	ENSMUST00000033898.9 6615 1669aa ENSMUST00000209598.1 1170 328aa ENSMUST00000208095.1 545 133aa ENSMUST00000209735.1 6487 1562aa ENSMUST00000209661.1 618 126aa ENSMUST00000208386.1 416 No protein ENSMUST00000130488.2 676 No protein	ENSMUST00000033898.9 6615 1669aa Protein coding ENSMUST00000209598.1 1170 328aa Protein coding ENSMUST00000208095.1 545 133aa Protein coding ENSMUST00000209735.1 6487 1562aa Nonsense mediated decay ENSMUST00000209661.1 618 126aa Nonsense mediated decay ENSMUST00000208386.1 416 No protein Processed transcript ENSMUST00000130488.2 676 No protein Retained intron	ENSMUST00000033898.9 6615 1669aa Protein coding CCDS40219 ENSMUST00000209598.1 1170 328aa Protein coding - ENSMUST00000208095.1 545 133aa Protein coding - ENSMUST00000209735.1 6487 1562aa Nonsense mediated decay - ENSMUST00000209661.1 618 126aa Nonsense mediated decay - ENSMUST00000208386.1 416 No protein Processed transcript - ENSMUST00000130488.2 676 No protein Retained intron -	ENSMUST00000033898.9 6615 1669aa Protein coding CCDS40219 P02463 ENSMUST00000209598.1 1170 328aa Protein coding - A0A1B0GRC0 ENSMUST00000208095.1 545 133aa Protein coding - A0A140LHU8 ENSMUST00000209735.1 6487 1562aa Nonsense mediated decay - A0A1B0GSI7 ENSMUST000000209661.1 618 126aa Nonsense mediated decay - A0A1B0GT69 ENSMUST000000208386.1 416 No protein Processed transcript - - ENSMUST000000130488.2 676 No protein Retained intron - -

The strategy is based on the design of Col4a1-201 transcript, The transcription is shown below



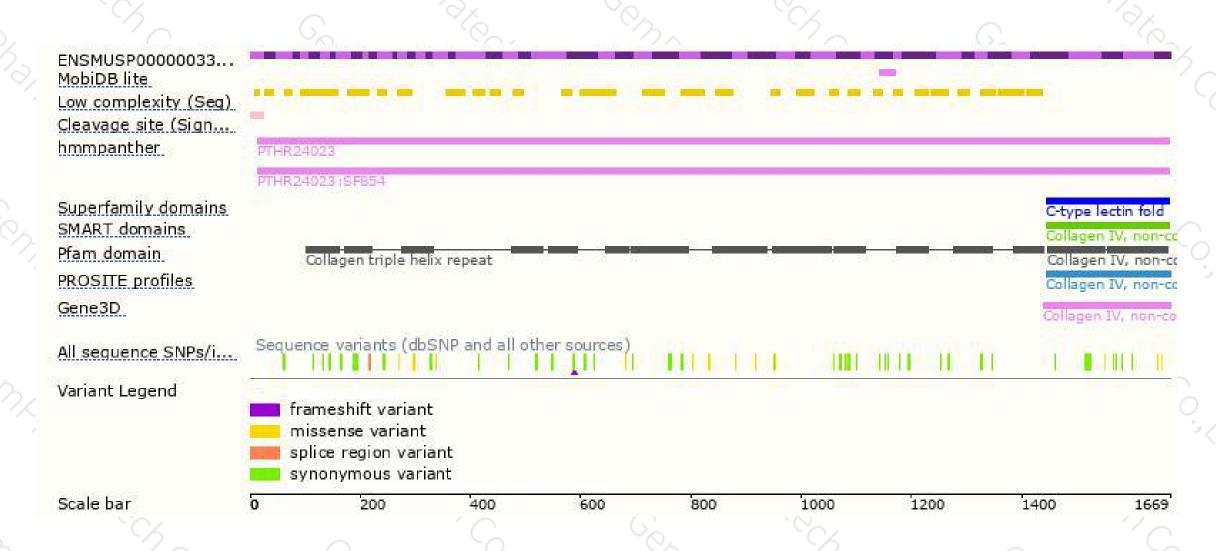
Genomic location distribution





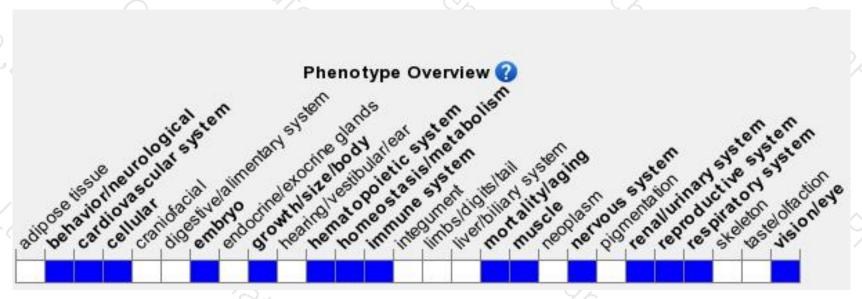
Protein domain





Mouse phenotype description(MGI)





Phenotypes affected by the gene are marked in blue.Data quoted from MGI database(http://www.informatics.jax.org/).

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





