

# Col2a1 Cas9-CKO Strategy

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## **Project Overview**



**Project Name** 

Col2a1

**Project type** 

Cas9-CKO

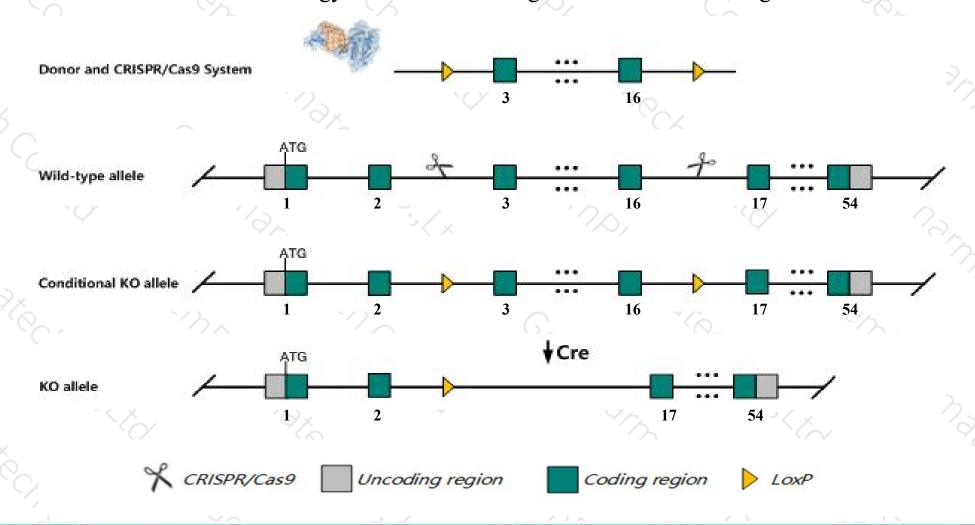
Strain background

C57BL/6JGpt

## Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Col2a1 gene. The schematic diagram is as follows:



### Technical routes



- The *Col2a1* gene has 10 transcripts. According to the structure of *Col2a1* gene, exon3-exon16 of *Col2a1-201* (ENSMUST00000023123.14) transcript is recommended as the knockout region. The region contains 734bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Col2a1* 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, Mutations in this locus affect cartilage development. Homozygotes die perinatally with anomalies such as shortened limbs without epiphiseal growth plates, cleft palate and persistence of notochord. Heterozygotes are dwarfed with reduced cartilage matrix.
- The Col2a1 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

### Gene information (NCBI)



#### Col2a1 collagen, type II, alpha 1 [Mus musculus (house mouse)]

Gene ID: 12824, updated on 9-Apr-2019

#### Summary

☆ ?

Official Symbol Col2a1 provided by MGI

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

Primary source MGI:MGI:88452

See related Ensembl: ENSMUSG00000022483

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 Col2, Col2a, Col2a-1, Del1, Dmm, Lpk, M100413, Rgsc413, Rgsc856

Summary This gene encodes the alpha-1 subunit of the fibril-forming type II collagen, the major component of cartilage and the vitreous humor of the

eye. The encoded preproprotein forms homotrimeric, triple helical procollagen that undergoes proteolytic processing during fibirl formation.

Mice harboring certain mutations in this gene exhibit severe chondrodysplasia characterized by short limbs and trunch, craniofacial

deformities and cleft palate. A complete lack of the encoded protein in mice results in postnatal lethality. Alternative splicing results in multiple

transcript variants encoding different isoforms that may undergo similar proteolytic processing. [provided by RefSeq, Dec 2015]

Expression Biased expression in limb E14.5 (RPKM 504.7), CNS E14 (RPKM 67.6) and 1 other tissueSee more

Orthologs <u>human</u> all

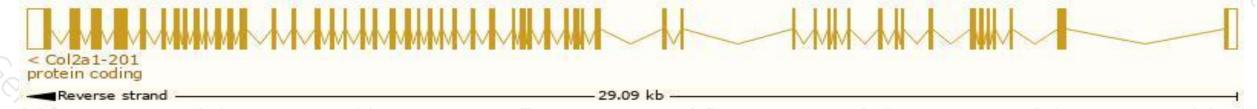
## Transcript information (Ensembl)



The gene has 10 transcripts, all transcripts are shown below:

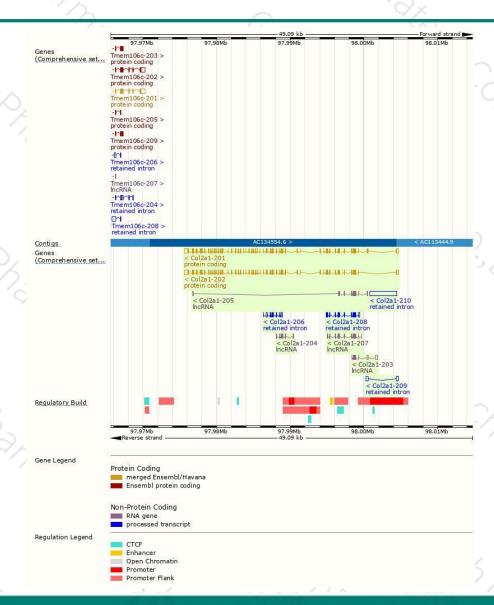
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Col2a1-201	ENSMUST00000023123.14	5104	1487aa	Protein coding	CCDS37189	P28481	TSL:1 GENCODE basic APPRIS P3
Col2a1-202	ENSMUST00000088355.11	4862	<u>1419aa</u>	Protein coding	CCDS49716	P28481	TSL:1 GENCODE basic APPRIS ALT1
Col2a1-210	ENSMUST00000230148.1	3666	No protein	Retained intron	94	0.20	
Col2a1-208	ENSMUST00000139246.7	809	No protein	Retained intron	82	3523	TSL:5
Col2a1-206	ENSMUST00000131910.7	654	No protein	Retained intron	85	1.7	TSL:5
Col2a1-209	ENSMUST00000140064.1	613	No protein	Retained intron	89		TSL:2
Col2a1-203	ENSMUST00000127879.1	715	No protein	IncRNA	<u> </u>	12	TSL:3
Col2a1-207	ENSMUST00000133488.7	626	No protein	IncRNA	ė.	525	TSL:5
Col2a1-205	ENSMUST00000131560.2	575	No protein	IncRNA	85	-	TSL:3
Col2a1-204	ENSMUST00000128547.1	491	No protein	IncRNA	i -		TSL:3
			74.0			7	

The strategy is based on the design of Col2a1-201 transcript, The transcription is shown below



### Genomic location distribution





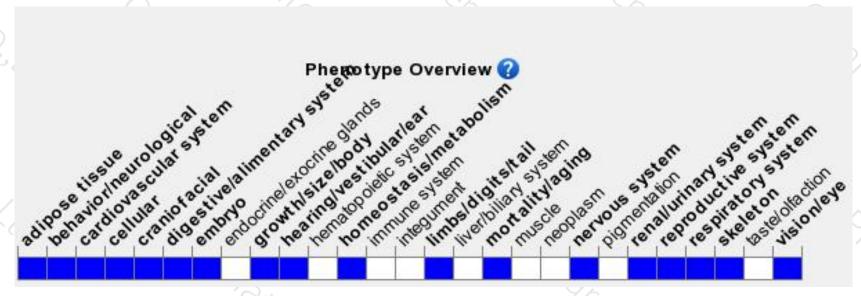
## Protein domain





## 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, Mutations in this locus affect cartilage development. Homozygotes die perinatally with anomalies such as shortened limbs without epiphiseal growth plates, cleft palate and persistence of notochord. Heterozygotes are dwarfed with reduced cartilage matrix.



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





