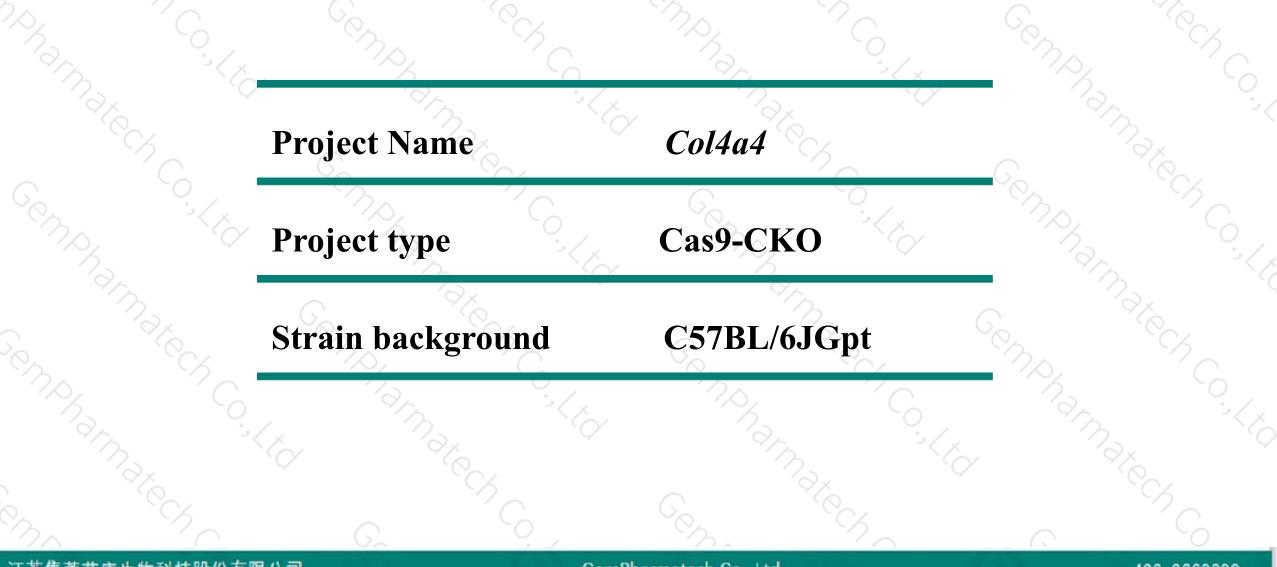


Col4a4 Cas9-CKO Strategy

Designer: Reviewer: Design Date: Huimin Su Ruirui Zhang 2019/8/30

Project Overview





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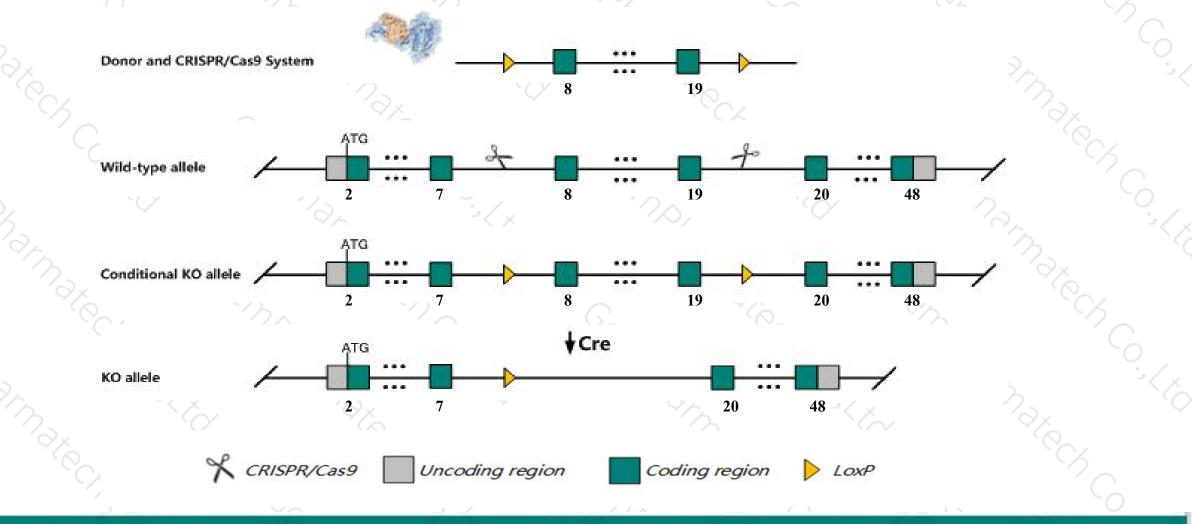
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Conditional Knockout strategy



400-9660890

This model will use CRISPR/Cas9 technology to edit the Col4a4 gene. The schematic diagram is as follows:



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The Col4a4 gene has 2 transcripts. According to the structure of Col4a4 gene, exon8-exon19 of Col4a4-201 (ENSMUST00000087050.6) transcript is recommended as the knockout region. The region contains 715bp coding sequence. Knock out the region will result in disruption of protein function.

In this project we use CRISPR/Cas9 technology to modify *Col4a4* 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.



- According to the existing MGI data, Mice homozygous for an ENU-induced mutation develop an early nephritic syndrome associated with uremia, proteinuria, hematuria, leukocyturia, and focal segmental glomerulosclerosis, and die prematurely of kidney failure. Some homozygotes exhibit moderatesensorineural hearing loss.
- The Col4a4 gene is located on the Chr1. 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)



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Col4a4 collagen, type IV, alpha 4 [Mus musculus (house mouse)]

Gene ID: 12829, updated on 12-Aug-2019

- Summary

Official Symbol	Col4a4 provided by MGI
Official Full Name	collagen, type IV, alpha 4 provided by MGI
Primary source	MGI:MGI:104687
See related	Ensembl:ENSMUSG0000067158
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	[a]4(IV); E130010M05Rik
Expression	Biased expression in kidney adult (RPKM 12.0), lung adult (RPKM 10.4) and 6 other tissues See more
Orthologs	human all

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Transcript information (Ensembl)



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The gene has 2 transcripts, all transcripts are shown below:

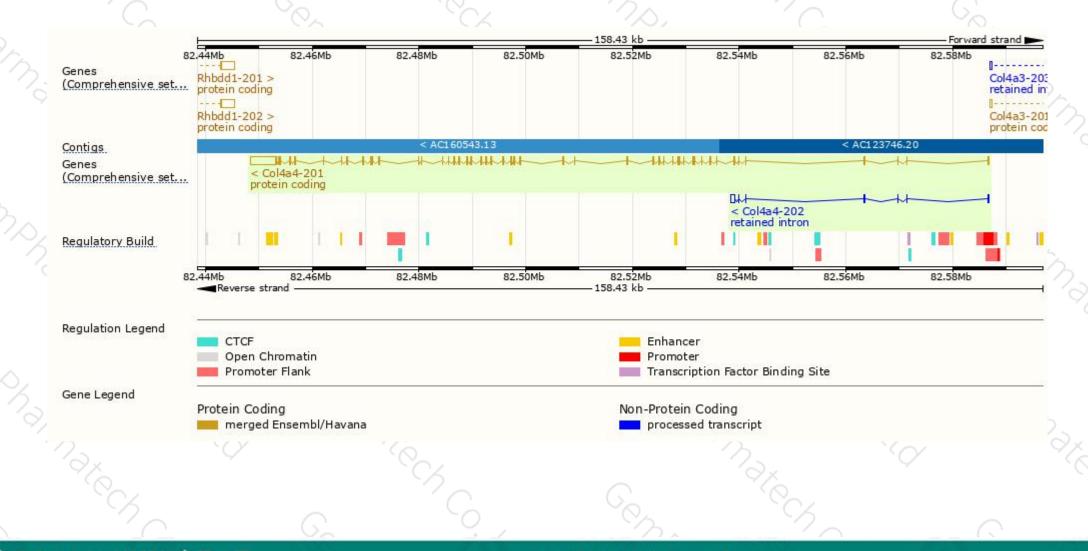
Name 🍦	Transcript ID 💧	bp 🖕	Protein 🛔	Biotype 💧	CCDS 🍦	UniProt 🖕	F	lags	*
Col4a4-201	ENSMUST0000087050.6	10108	<u>1682aa</u>	Protein coding	<u>CCDS35630</u> &	<u>Q9QZR9</u> ₽	TSL:1 GENCO	DE basic	APPRIS P1
Col4a4-202	ENSMUST00000135271.1	1263	No protein	Retained intron	_	10		TSL:5	

The strategy is based on the design of Col4a4-201 transcript, The transcription is shown below

Reverse strand	$\gamma_{\mathcal{O}_{\mathcal{O}_{\mathcal{O}_{\mathcal{O}_{\mathcal{O}_{\mathcal{O}}}}}}$	138.43 kb —	G _R	°C2	
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Genomic location distribution





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Protein domain



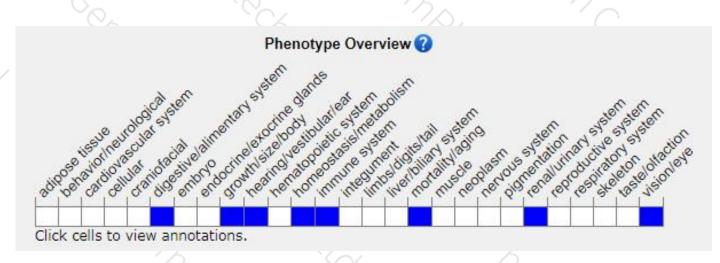
ENSMUSP00000084 MobiDB lite Low complexity (Seg)							
Superfamily SMART						C-type lectin fold	5
	llagen triple helix repeat					Collagen IV, non-collage Collagen IV, non-collage Collagen IV, non-collage	
PANTHER	PTHR24023:SF714		23 				
Gene3D	PTHR24023				*	Collagen IV, non-collage	er
All Seque	nce variants (dbSNP and	all other sources)					-1
Variant Legend	frame insertion	d all other sources)		sense variant			X
Variant Legend		d all other sources)	mis:				XC
Variant Legend	frame insertion lice region variant		miss syne	sense variant onymous variant		ne sta non n ear an a	XC

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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 an ENU-induced mutation develop an early nephritic syndrome associated with uremia, proteinuria, hematuria, leukocyturia, and focal segmental glomerulosclerosis, and die prematurely of kidney failure. Some homozygotes exhibit moderates ensorine ural hearing loss.

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



