

***Zcchc14* Cas9-CKO Strategy**

Designer: Jia Yu

Reviewer: Xiaojing Li

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Project Overview

Project Name

Zcchc14

Project type

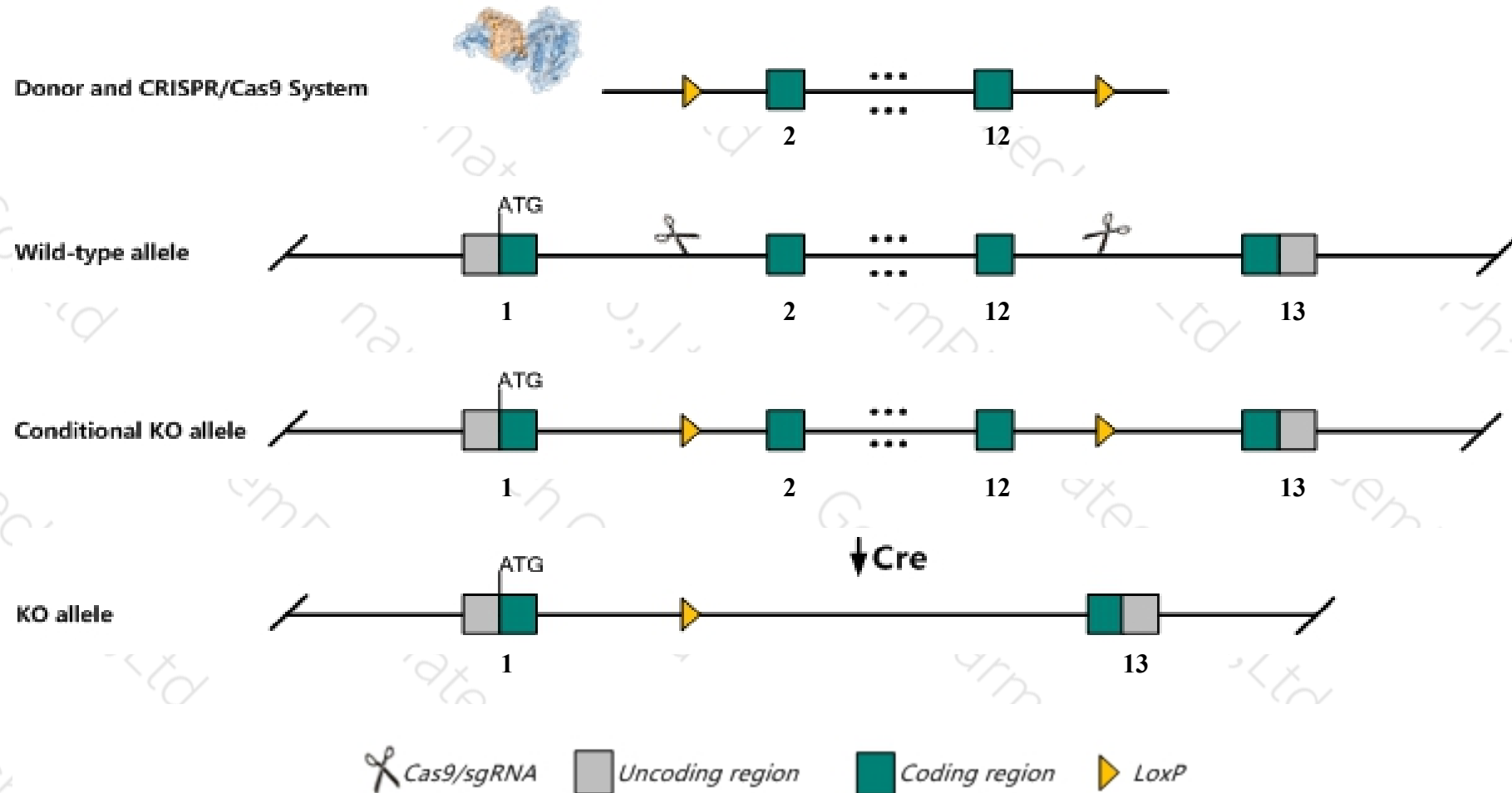
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Zcchc14* gene. The schematic diagram is as follows:



Technical routes

- The *Zcchc14* gene has 4 transcripts. According to the structure of *Zcchc14* gene, exon2-exon12 of *Zcchc14-201*(ENSMUST00000046386.4) transcript is recommended as the knockout region. The region contains 2638bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Zcchc14* gene. The brief process is as follows: sgRNA was transcribed in vitro, donor vector was constructed. Cas9, sgRNA 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 was 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.

- The *Zcchc14* 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)

Zcchc14 zinc finger, CCHC domain containing 14 [Mus musculus (house mouse)]

Gene ID: 142682, updated on 13-Mar-2020

Summary



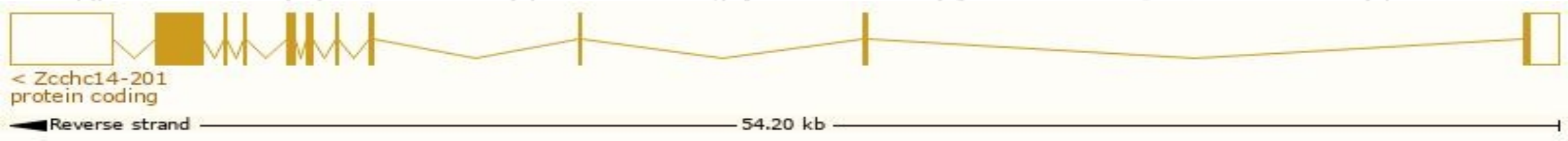
Official Symbol	Zcchc14 provided by MGI
Official Full Name	zinc finger, CCHC domain containing 14 provided by MGI
Primary source	MGI:MGI:2159407
See related	Ensembl:ENSMUSG00000061410
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	AA792890, BDG29
Expression	Ubiquitous expression in ovary adult (RPKM 21.7), adrenal adult (RPKM 17.7) and 26 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

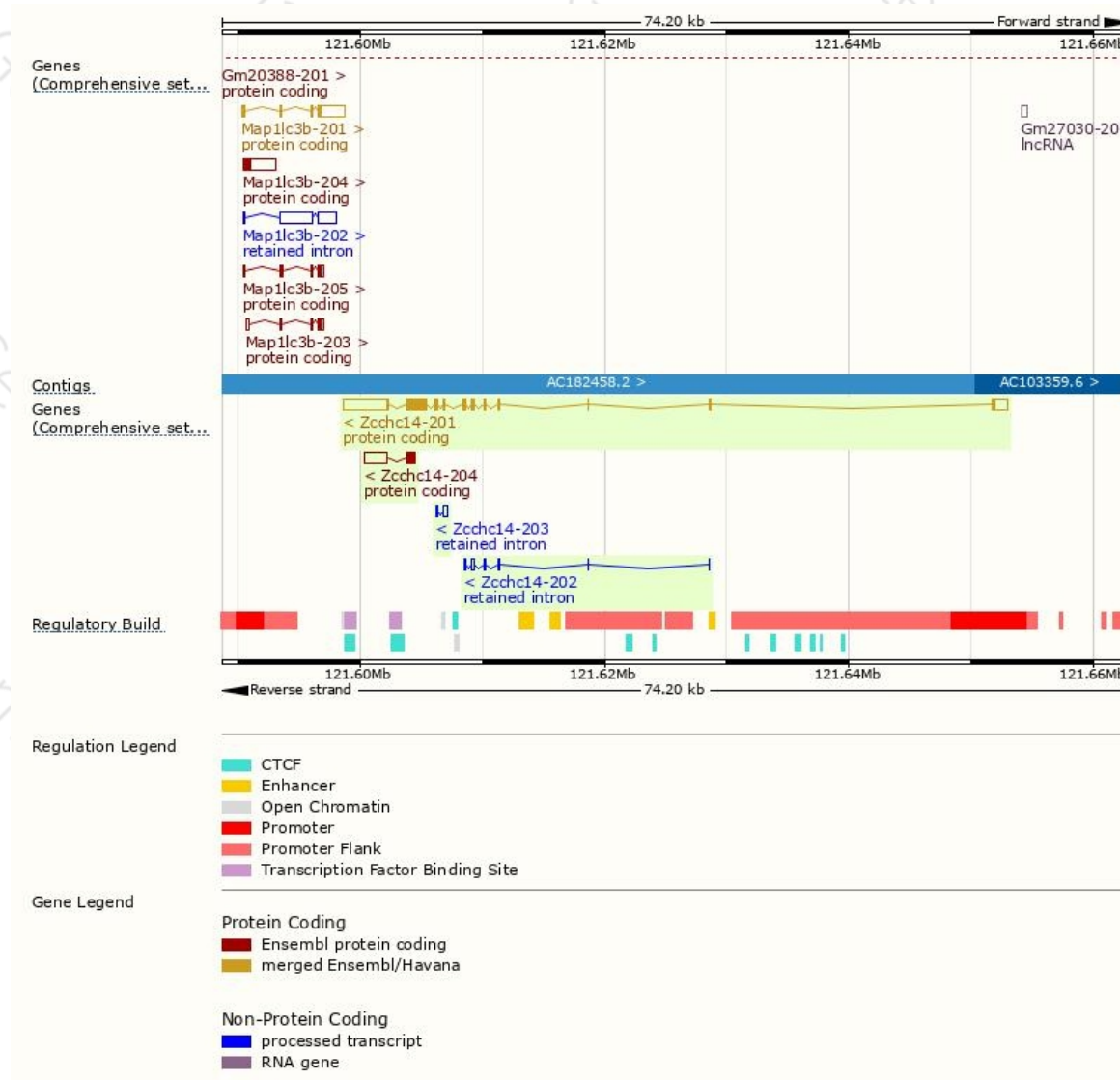
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Zcchc14-201	ENSMUST00000046386.4	7444	956aa	Protein coding	CCDS22727	A2RTC2 Q8VIG0	TSL:1 GENCODE basic APPRIS P1
Zcchc14-204	ENSMUST00000154725.1	2607	242aa	Protein coding	-	F6XMW0	CDS 5' incomplete TSL:1
Zcchc14-202	ENSMUST00000134212.1	625	No protein	Retained intron	-	-	TSL:3
Zcchc14-203	ENSMUST00000139379.1	446	No protein	Retained intron	-	-	TSL:2

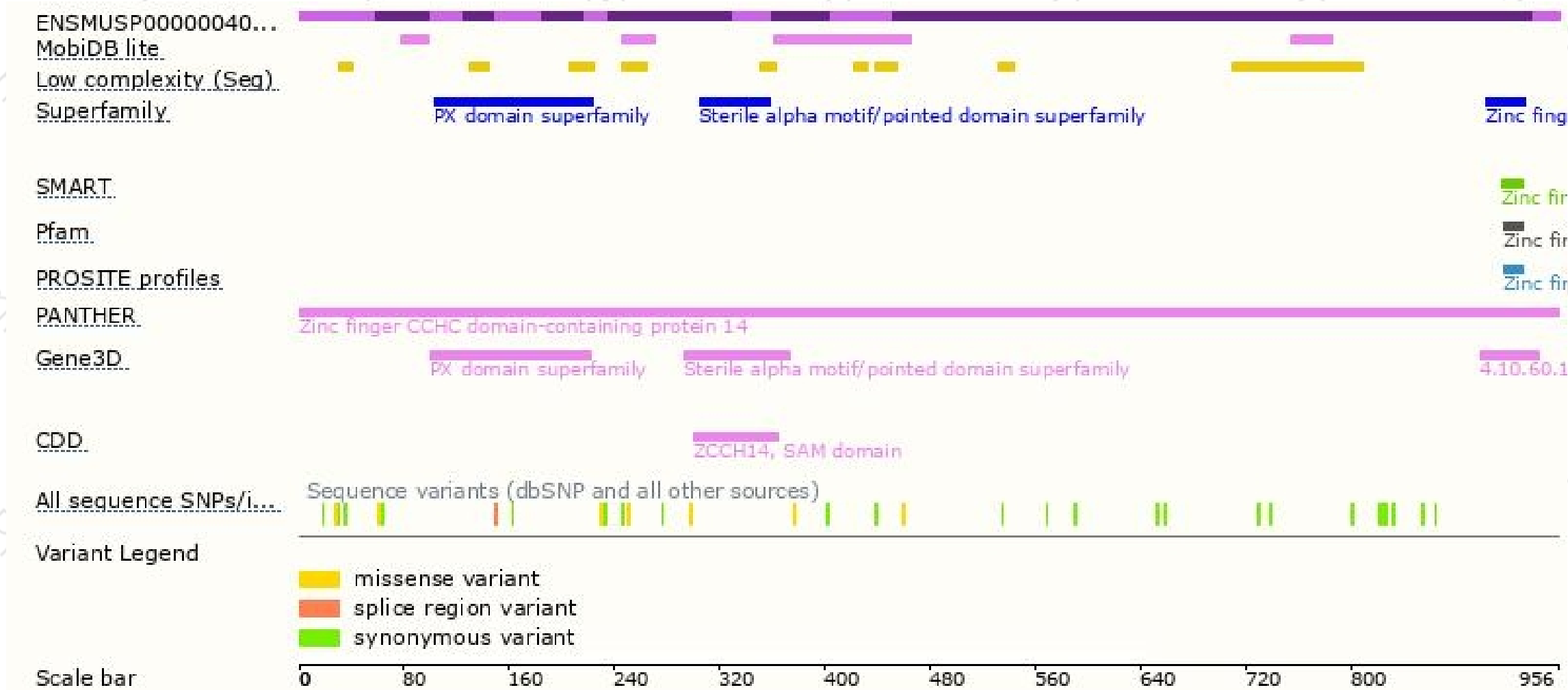
The strategy is based on the design of *Zcchc14-201* transcript,the transcription is shown below:



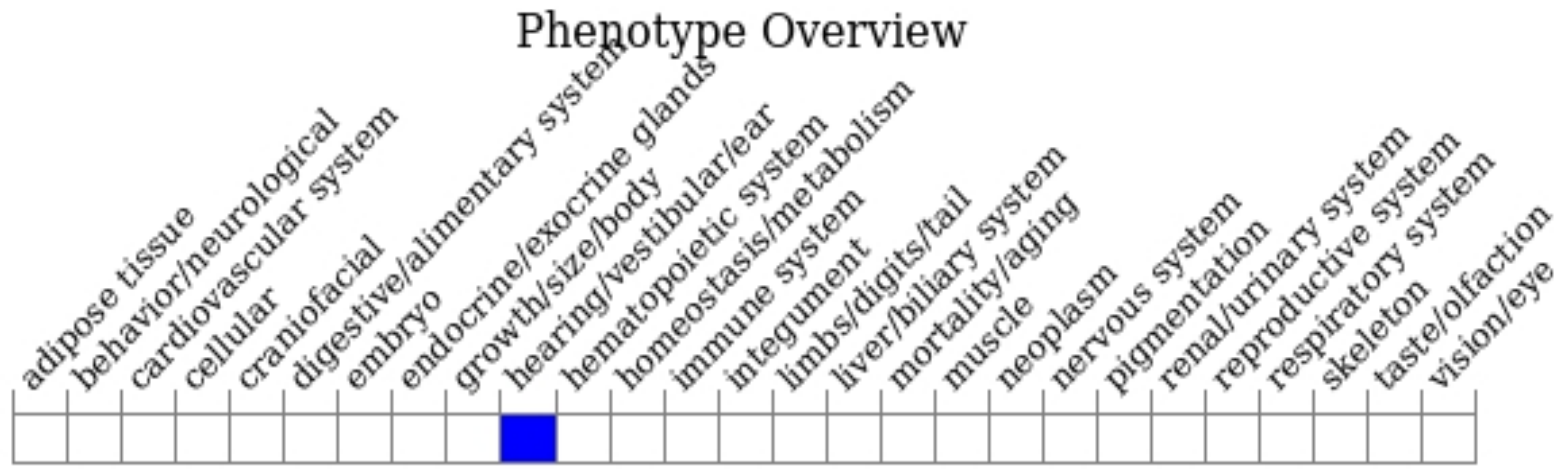
Genomic location distribution



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/>).

If you have any questions, you are welcome to inquire.

Tel: 025-5864 1534

