

***Zfp324* Cas9-CKO Strategy**

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Project Overview

Project Name

Zfp324

Project type

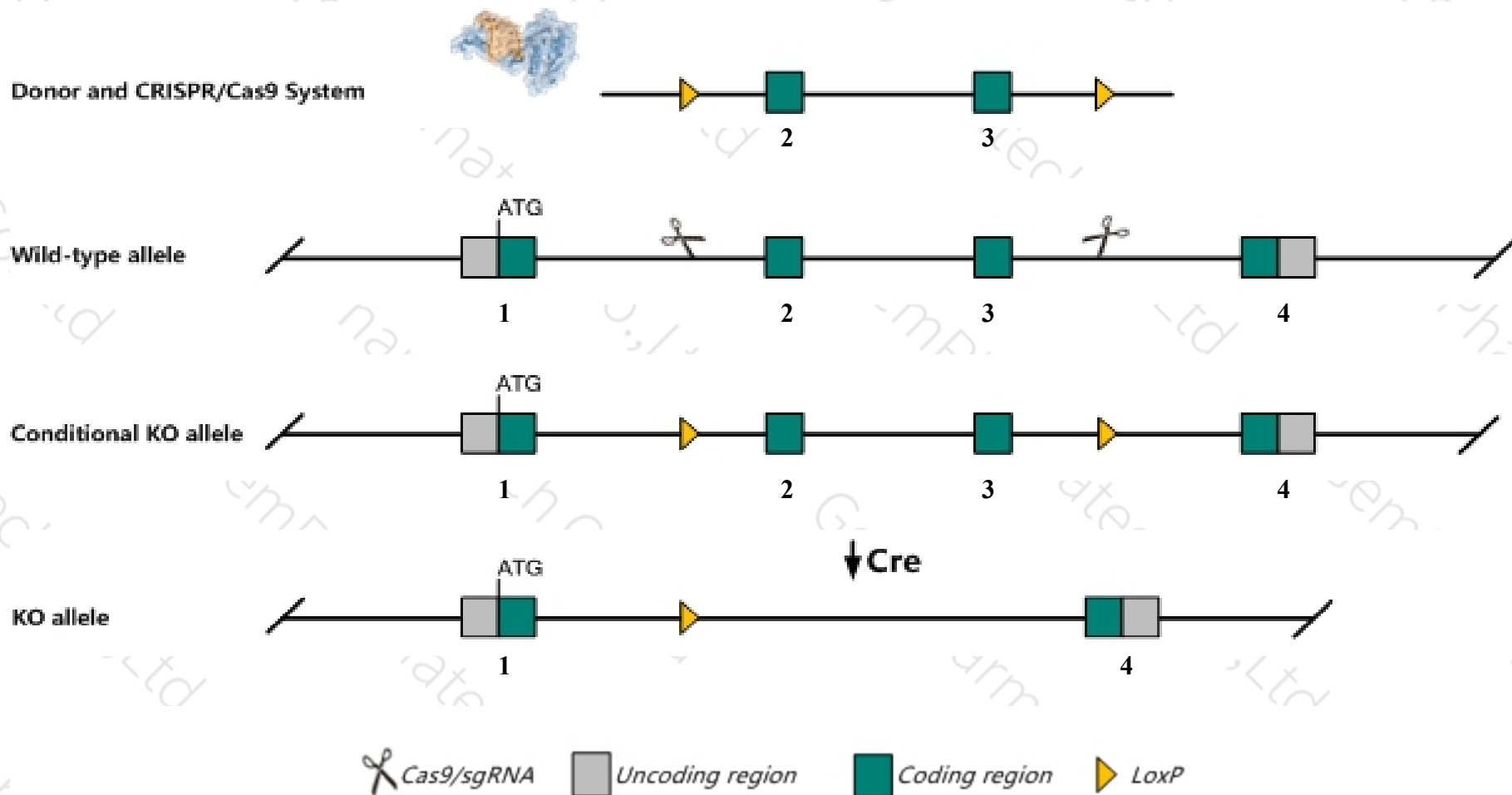
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Zfp324* gene. The schematic diagram is as follows:



Technical routes

- The *Zfp324* gene has 4 transcripts. According to the structure of *Zfp324* gene, exon2-exon3 of *Zfp324-201*(ENSMUST00000038701.7) transcript is recommended as the knockout region. The region contains 244bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Zfp324* 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.

Notice

- The *Zfp324* gene is located on the Chr7. 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)

Zfp324 zinc finger protein 324 [Mus musculus (house mouse)]

Gene ID: 243834, updated on 13-Mar-2020

Summary



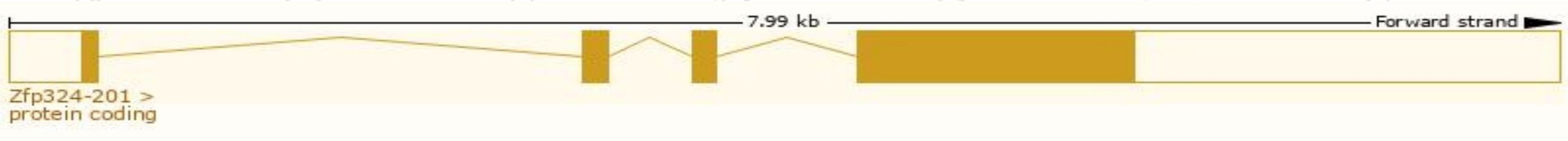
Official Symbol	Zfp324 provided by MGI
Official Full Name	zinc finger protein 324 provided by MGI
Primary source	MGI:MGI:2444641
See related	Ensembl:ENSMUSG00000004500
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	A930002M05, D430030K24Rik, ZF5128
Expression	Ubiquitous expression in ovary adult (RPKM 6.1), adrenal adult (RPKM 3.9) and 28 other tissues See more

Transcript information (Ensembl)

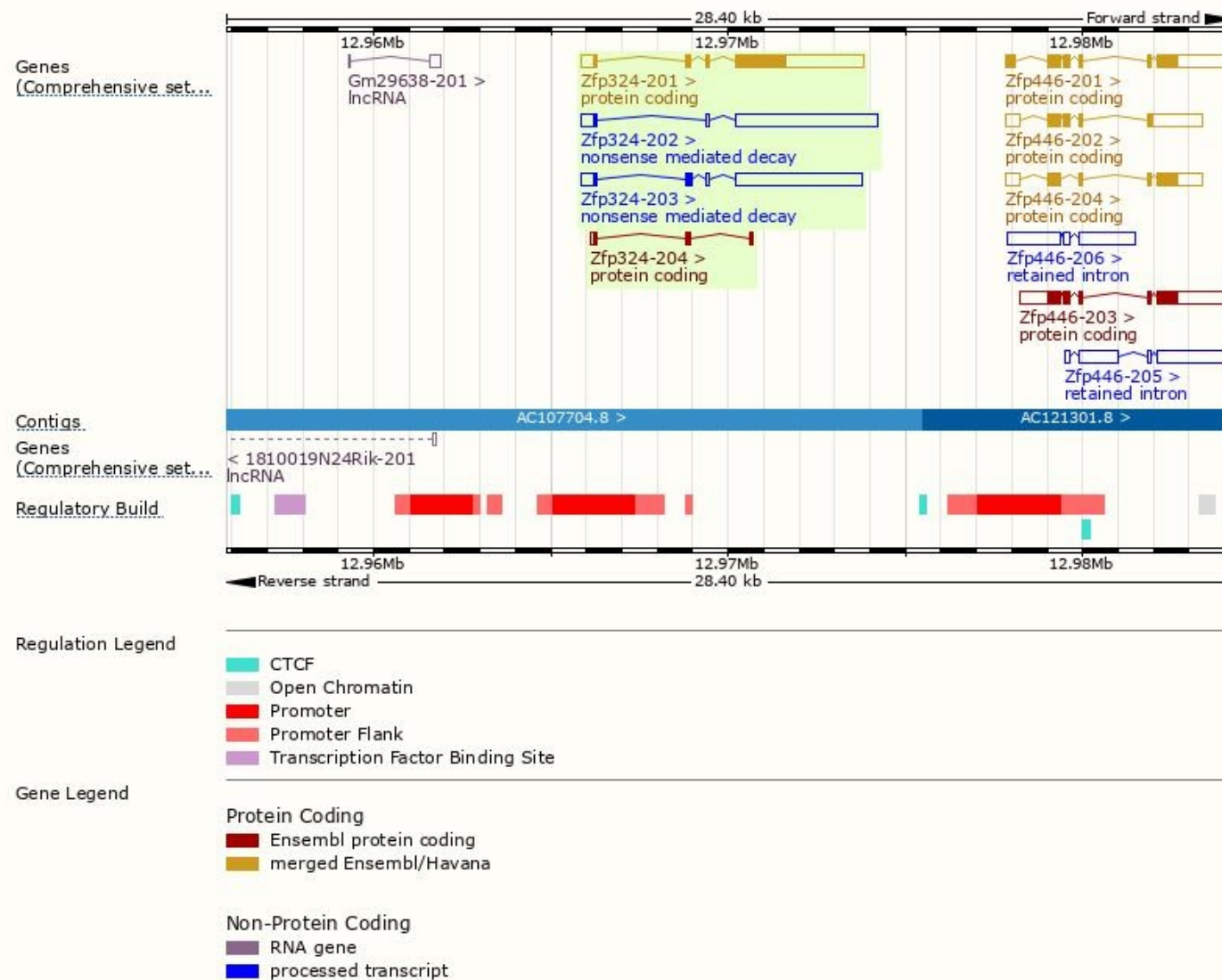
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Zfp324-201	ENSMUST00000038701.7	4315	583aa	Protein coding	CCDS20819	Q78F42	TSL:1 GENCODE basic APPRIS P1
Zfp324-204	ENSMUST00000210619.1	447	87aa	Protein coding	-	A0A1B0GSA6	TSL:3 GENCODE basic
Zfp324-202	ENSMUST00000124387.1	4595	36aa	Nonsense mediated decay	-	E0CYR3	TSL:1
Zfp324-203	ENSMUST00000128293.1	4310	75aa	Nonsense mediated decay	-	E0CXY6	TSL:1

The strategy is based on the design of *Zfp324-201* transcript,the transcription is shown below:



Genomic location distribution



Protein domain



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

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