

# ***Rps24* Cas9-CKO Strategy**

Designer: Xueting Zhang  
reviewer: Yanhua Shen  
Date: 2020-02-26

# Project Overview

**Project Name**

***Rps24***

**Project type**

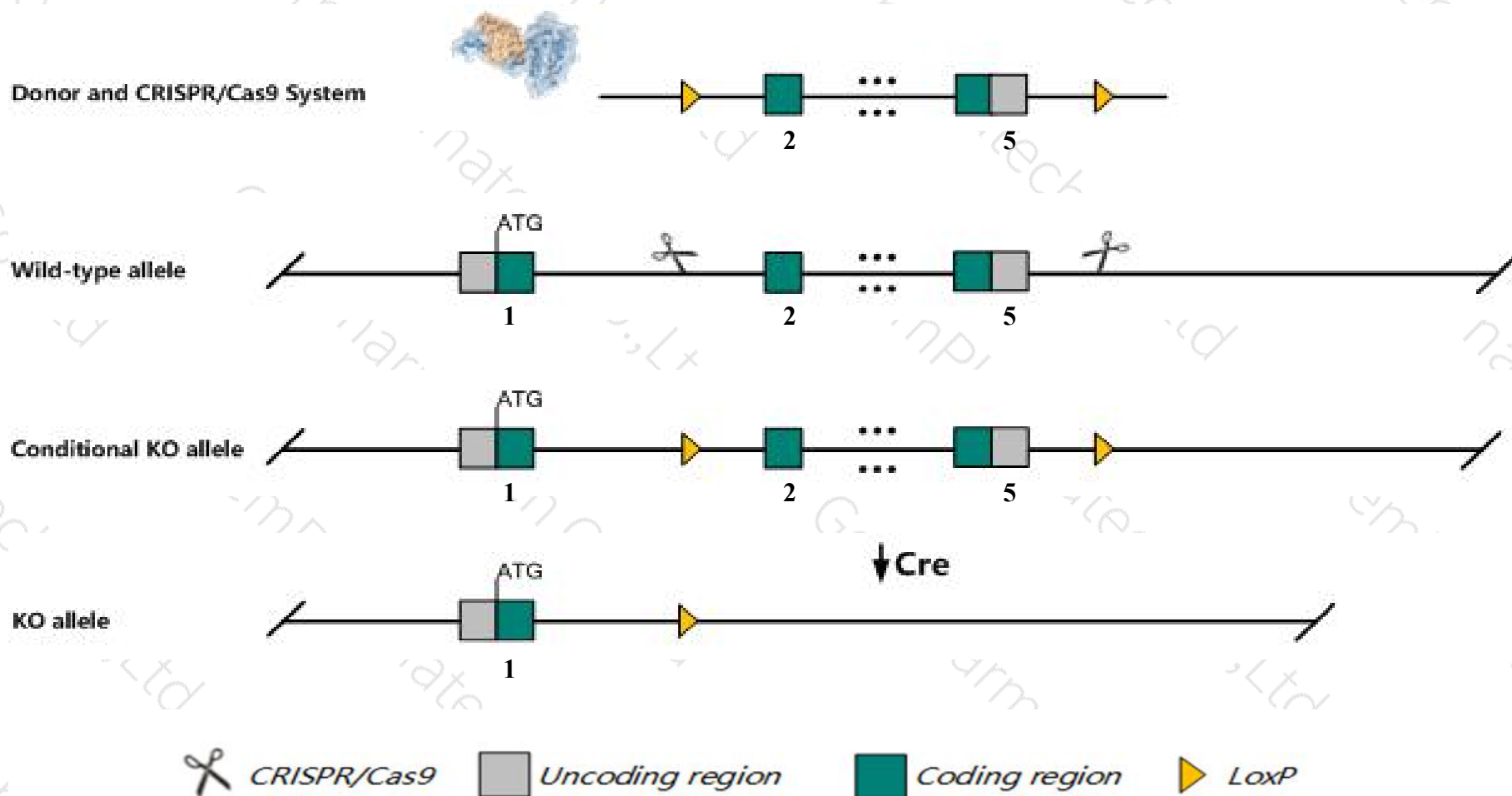
**Cas9-CKO**

**Strain background**

**C57BL/6JGpt**

# Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Rps24* gene. The schematic diagram is as follows:



- The *Rps24* gene has 12 transcripts. According to the structure of *Rps24* gene, exon2-exon5 of *Rps24-210* (ENSMUST00000225023.1) transcript is recommended as the knockout region. The region contains most of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Rps24* 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

- Transcript *Rps24-206&208&209* may be destroyed directly.
- The floxed region is near to the N-terminal of *Polr3a* gene, this strategy may influence the regulatory function of the N-terminal of *Polr3a* gene.
- The *Rps24* gene is located on the Chr14. 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)

## Rps24 ribosomal protein S24 [ *Mus musculus* (house mouse) ]

Gene ID: 20088, updated on 24-Oct-2019

Summary

- Official Symbol** Rps24 provided by [MGI](#)
- Official Full Name** ribosomal protein S24 provided by [MGI](#)
- Primary source** [MGI:MGI:98147](#)
- See related** [Ensembl:ENSMUSG00000025290](#)
- 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
- Expression** Ubiquitous expression in CNS E11.5 (RPKM 223.6), bladder adult (RPKM 143.0) and 24 other tissues [See more](#)
- Orthologs** [human](#) [all](#)

Genomic context

**Location:** 14; 14 A3

**Exon count:** 8

See Rps24 in [Genome Data Viewer](#)

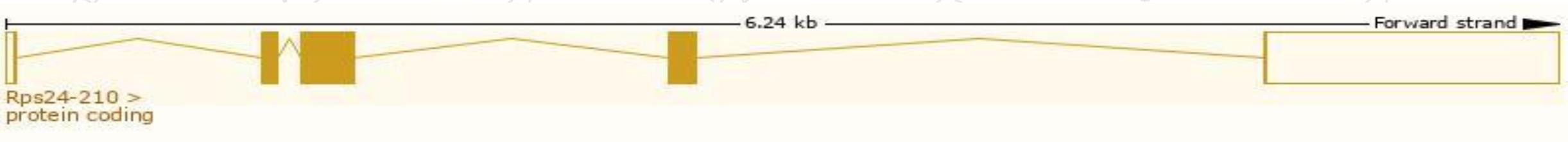
Annotation release	Status	Assembly	Chr	Location
<a href="#">108</a>	current	GRCm38.p6 ( <a href="#">GCF_000001635.26</a> )	14	NC_000080.6 (24490678..24496960)
Build 37.2	previous assembly	MGSCv37 ( <a href="#">GCF_000001635.18</a> )	14	NC_000080.5 (25309903..25315368)

# Transcript information (Ensembl)

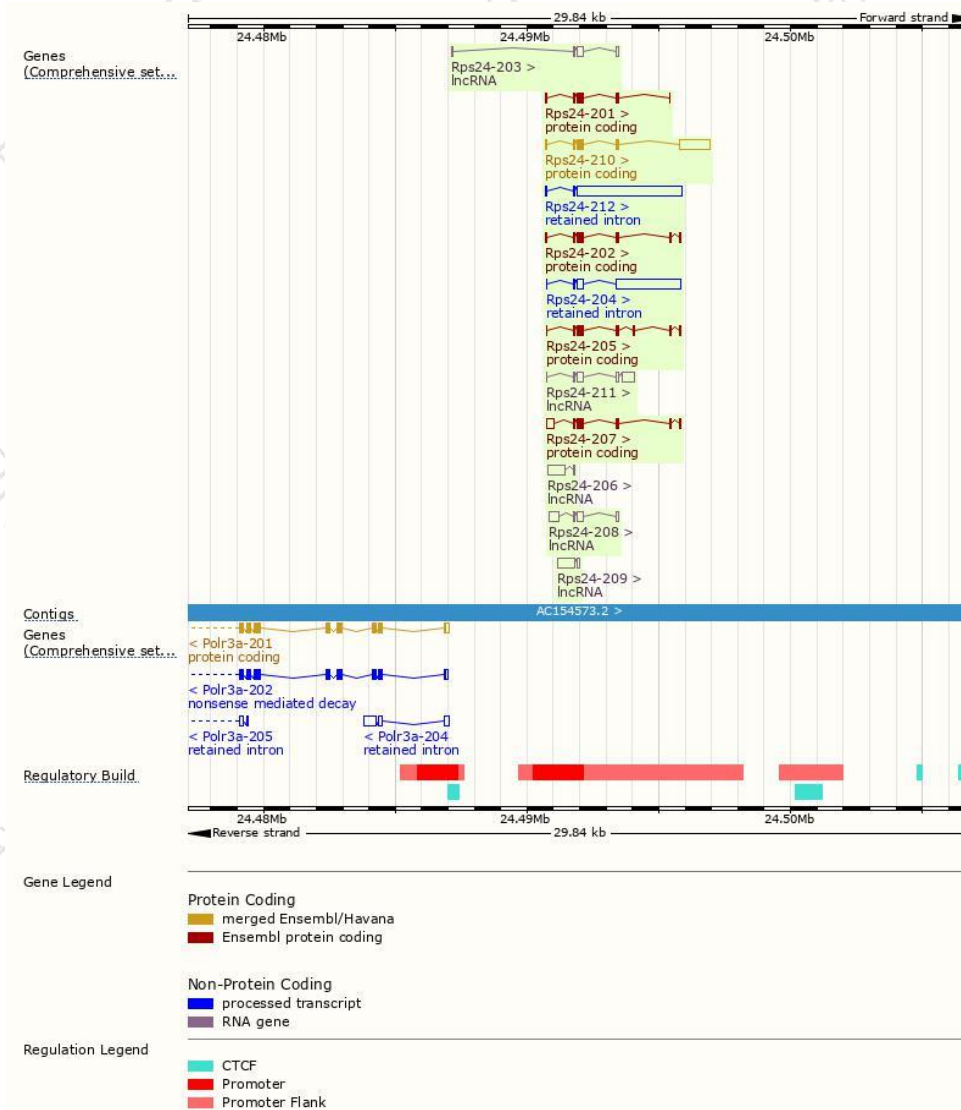
The gene has 12 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Rps24-210	<a href="#">ENSMUST00000225023.1</a>	1613	<a href="#">133aa</a>	Protein coding	<a href="#">CCDS36829</a>	<a href="#">P62849 Q5M9M7</a>	GENCODE basic APPRIS P4
Rps24-202	<a href="#">ENSMUST00000169826.2</a>	519	<a href="#">130aa</a>	Protein coding	<a href="#">CCDS36830</a>	<a href="#">P62849</a>	TSL:1 GENCODE basic APPRIS ALT 1
Rps24-201	<a href="#">ENSMUST00000112384.9</a>	468	<a href="#">130aa</a>	Protein coding	<a href="#">CCDS36830</a>	<a href="#">P62849</a>	TSL:5 GENCODE basic APPRIS ALT 1
Rps24-207	<a href="#">ENSMUST00000224568.1</a>	714	<a href="#">118aa</a>	Protein coding	-	<a href="#">A0A286YEB7</a>	GENCODE basic
Rps24-205	<a href="#">ENSMUST00000223999.1</a>	525	<a href="#">131aa</a>	Protein coding	-	<a href="#">P62849</a>	GENCODE basic APPRIS ALT 1
Rps24-212	<a href="#">ENSMUST00000225994.1</a>	4059	No protein	Retained intron	-	-	
Rps24-204	<a href="#">ENSMUST00000223939.1</a>	2769	No protein	Retained intron	-	-	
Rps24-211	<a href="#">ENSMUST00000225117.1</a>	887	No protein	lncRNA	-	-	
Rps24-209	<a href="#">ENSMUST00000224699.1</a>	750	No protein	lncRNA	-	-	
Rps24-206	<a href="#">ENSMUST00000224549.1</a>	745	No protein	lncRNA	-	-	
Rps24-208	<a href="#">ENSMUST00000224569.1</a>	686	No protein	lncRNA	-	-	
Rps24-203	<a href="#">ENSMUST00000223931.1</a>	461	No protein	lncRNA	-	-	

The strategy is based on the design of *Rps24-210* transcript,The transcription is shown below

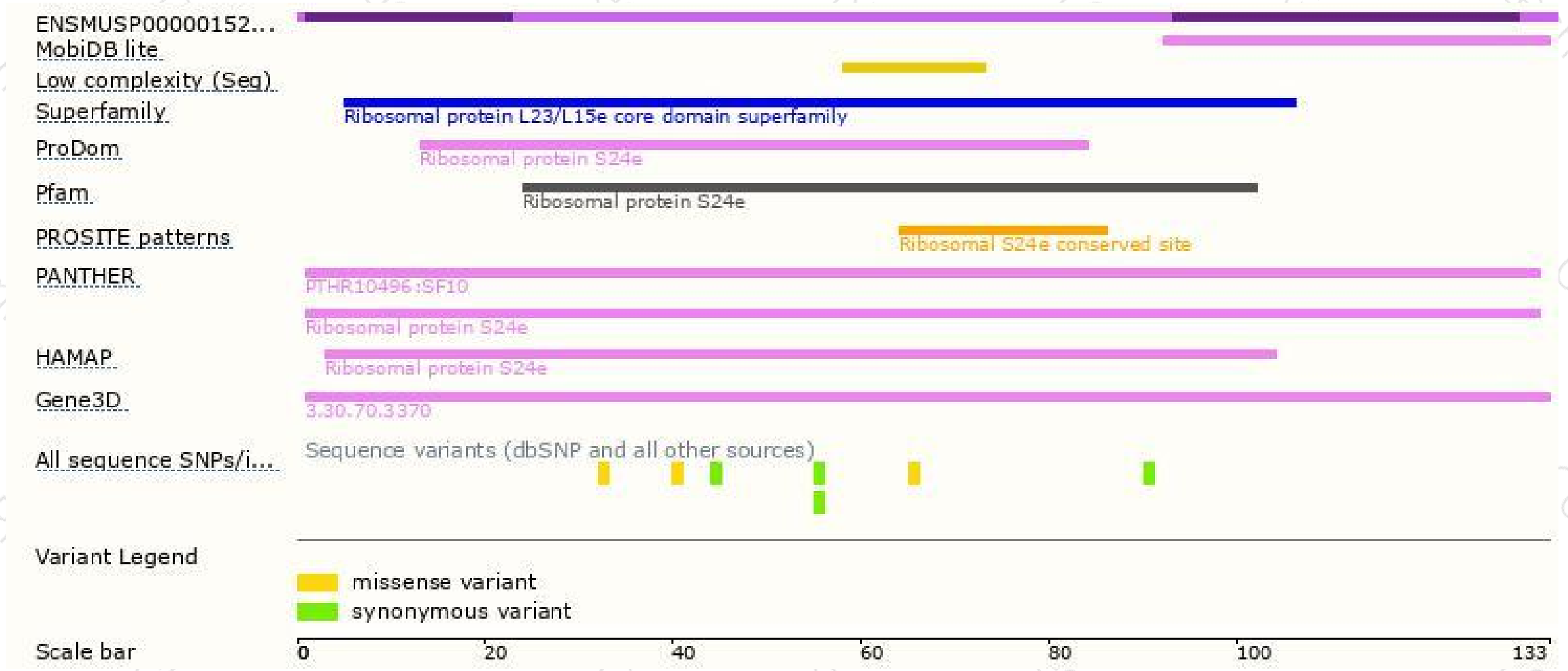


# Genomic location distribution





# Protein domain



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

Tel: 400-9660890

