Polr2a Cas9-KO Strategy

Designer: Daohua Xu
Reviewer: Huimin Su

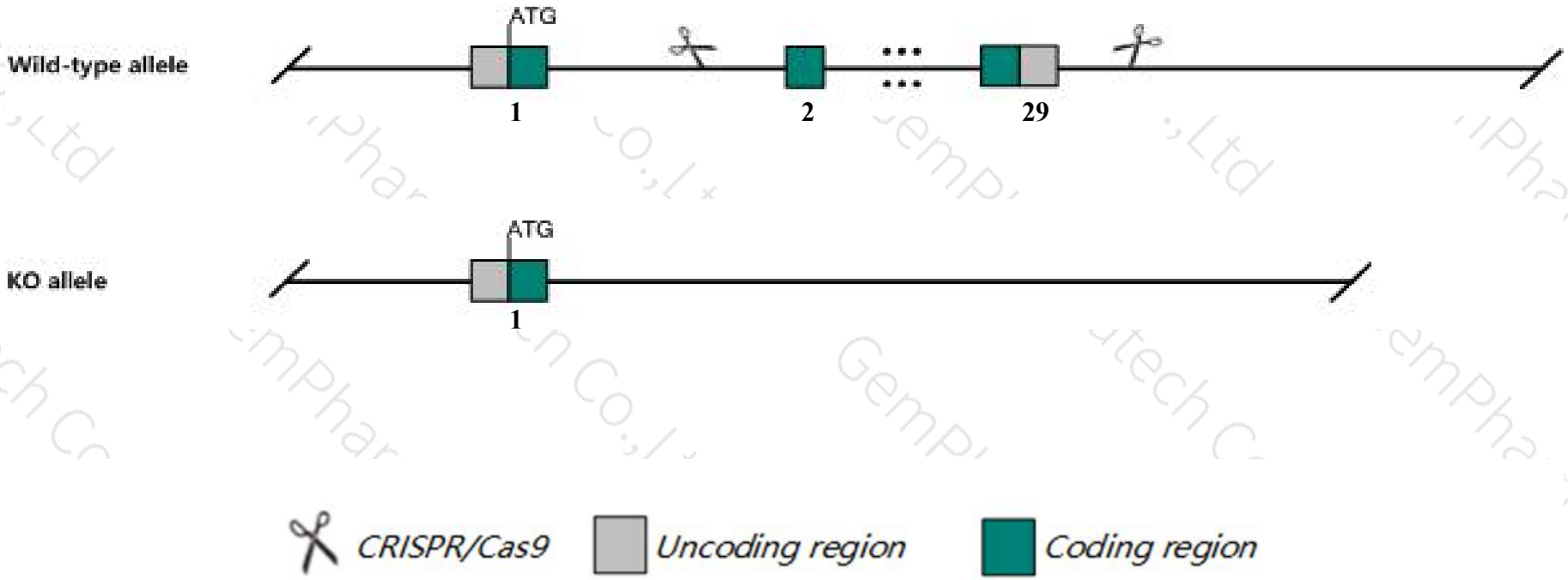
Design Date: 2019-11-22

Project Overview

Project Name	***Polr2a***
Project type	**Cas9-KO**
Strain background	**C57BL/6JGpt**

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Polr2a* gene. The schematic diagram is as follows:



Technical routes

➢ The *Polr2a* gene has 4 transcripts. According to the structure of *Polr2a* gene, exon2-exon29 of *Polr2a-201* (ENSMUST00000058470.15) 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 *Polr2a* gene. The brief process is as follows: CRISPR/Cas9 systen

Notice

- According to the existing MGI data,Homozygotes for a reporter allele show prenatal lethality. Homozygotes for a small deletion in the C-terminal domain are viable, fertile and developmentally normal. Homozygotes for a larger deletion show reduced fetal size and partial postnatal lethality; survivors are small but otherwise normal.

- The *Polr2a* gene is located on the Chr11. 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information（NCBI）

集萃药康 GemPharmatech

Polr2a polymerase (RNA) II (DNA directed) polypeptide A [Mus musculus (house mouse)]

Gene ID: 20020, updated on 5-Mar-2019

Summary

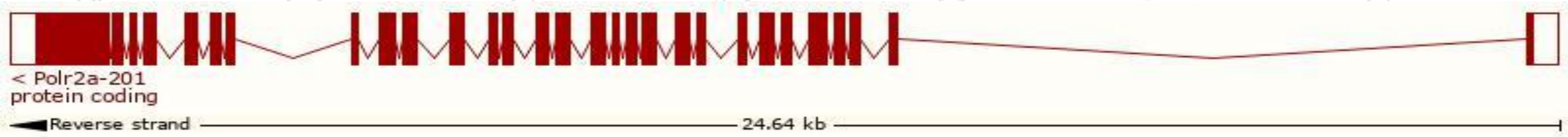
Official Symbol	Polr2a provided by MGI
Official Full Name	polymerase (RNA) II (DNA directed) polypeptide A provided by MGI
Primary source	MGI:MGI:98086
See related	Ensembl:ENSMUSG00000005198
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	220kDa, Rpb1, Rpo2-1
Expression	Ubiquitous expression in thymus adult (RPKM 39.8), limb E14.5 (RPKM 30.7) and 28 other tissues See more
Orthologs	human all

Transcript information（Ensembl）

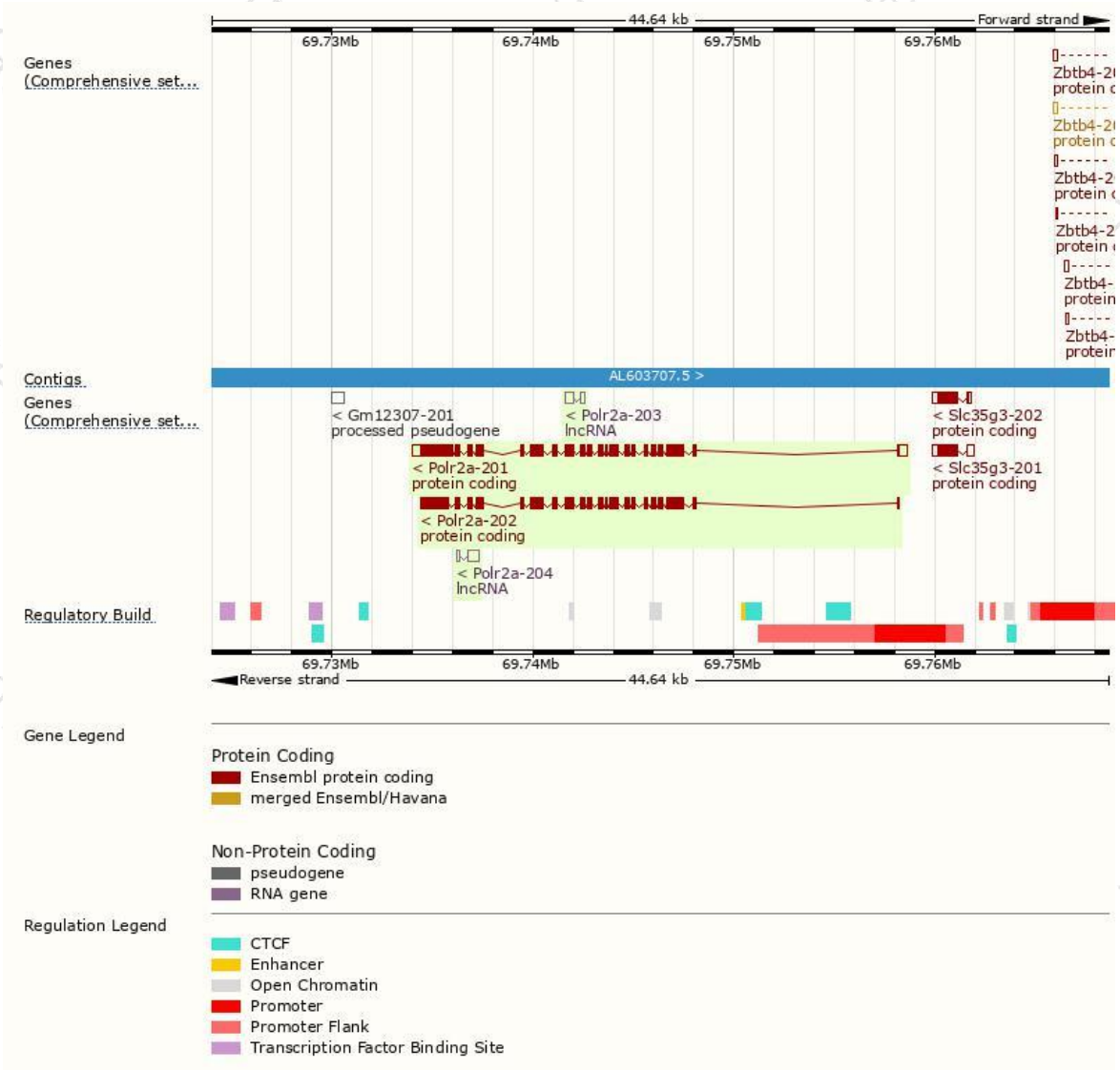
The gene has 4 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Polr2a-201	ENSMUST00000058470.15	6740	1970aa	Protein coding	CCDS70217	P08775	TSL:1 GENCODE basic APPRIS P1
Polr2a-202	ENSMUST00000071213.3	5799	1932aa	Protein coding	-	A0A0R4J0V5	TSL:5 GENCODE basic
Polr2a-204	ENSMUST00000156588.1	689	No protein	lncRNA	-	-	TSL:3
Polr2a-203	ENSMUST00000151586.1	586	No protein	lncRNA	-	-	TSL:2

The strategy is based on the design of *Polr2a-201* transcript,The transcription is shown below

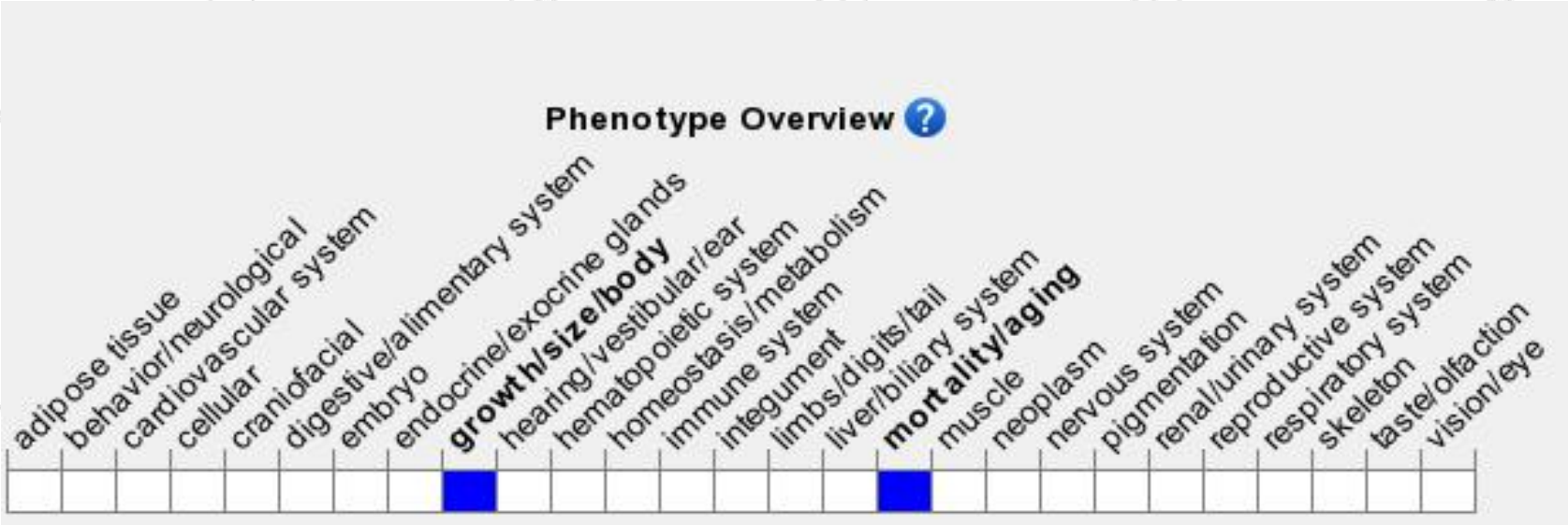


Genomic location distribution



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,Homozygotes for a reporter allele show prenatal lethality. Homozygotes for a small deletion in the C-terminal domain are viable, fertile and developmentally normal. Homozygotes for a larger deletion show reduced fetal size and partial postnatal lethality; survivors are small but otherwise normal.

集萃药康
GemPharmatech

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

Tel: 400-9660890

集萃药康生物科技
GemPharmatech Co.,Ltd