

***Art4* Cas9-KO Strategy**

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Project Overview

Project Name

Art4

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Art4* gene. The schematic diagram is as follows:



- The *Art4* gene has 1 transcript. According to the structure of *Art4* gene, exon2 of *Art4-201* (ENSMUST00000032341.2) transcript is recommended as the knockout region. The region contains 709bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Art4* gene. The brief process is as follows: CRISPR/Cas9 system we

- The *Art4* gene is located on the Chr6. 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)

Art4 ADP-ribosyltransferase 4 [*Mus musculus* (house mouse)]

Gene ID: 109978, updated on 13-Mar-2020

Summary

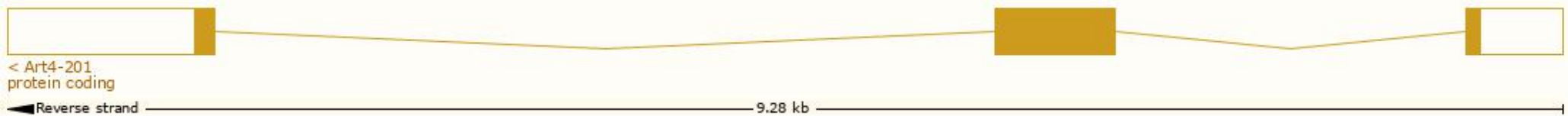
Official Symbol	Art4 provided by MGI
Official Full Name	ADP-ribosyltransferase 4 provided by MGI
Primary source	MGI:MGI:1202710
See related	Ensembl:ENSMUSG00000030217
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	DO; DOK1; ARTC4; 4432404K01Rik
Expression	Biased expression in liver E14.5 (RPKM 15.6), liver E14 (RPKM 12.1) and 12 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

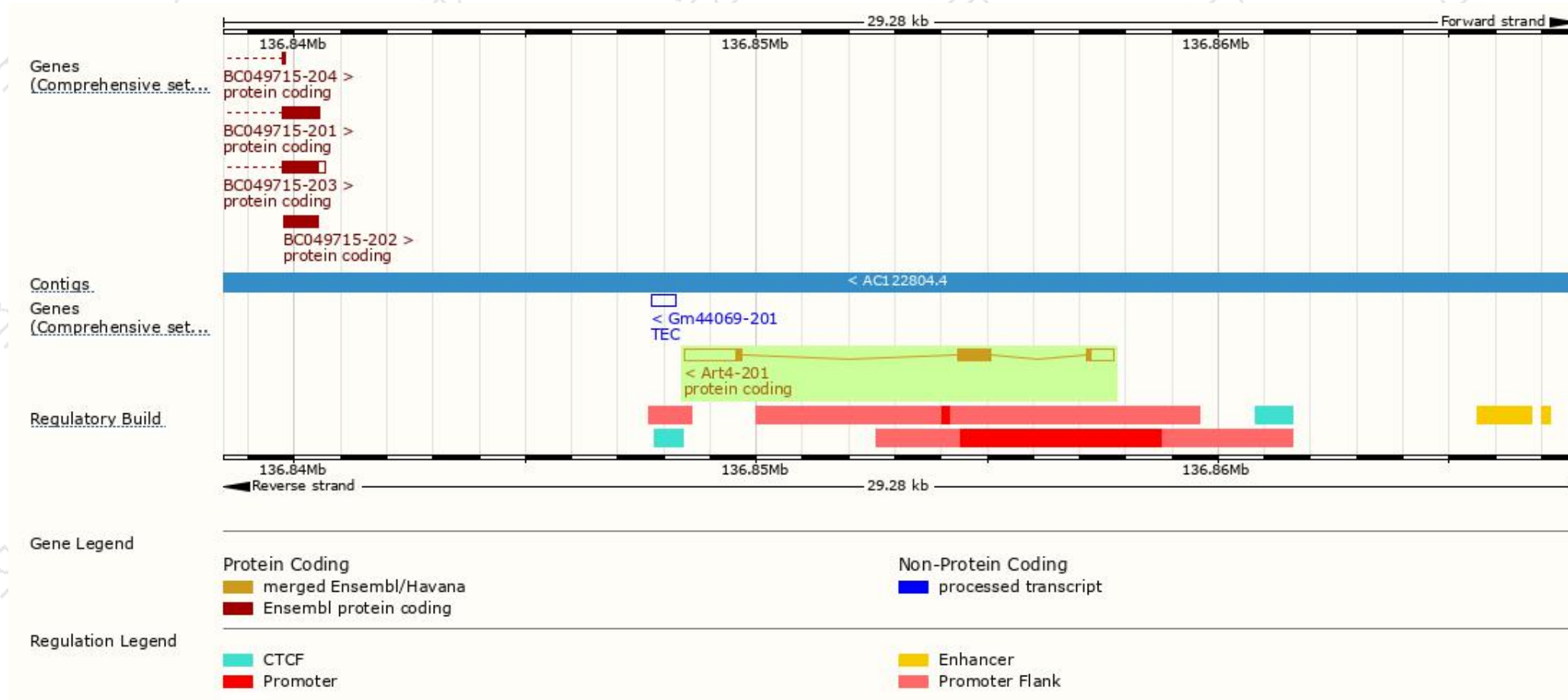
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Art4-201	ENSMUST00000032341.2	2517	300aa	Protein coding	CCDS20658	Q9CRA0	TSL:1 Gencode basic APPRIS P1

The strategy is based on the design of *Art4-201* 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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