

***Taf5* Cas9-KO Strategy**

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Project Overview

Project Name

Taf5

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Taf5* gene. The schematic diagram is as follows:



- The *Taf5* gene has 1 transcript. According to the structure of *Taf5* gene, exon2 of *Taf5-201* (ENSMUST00000026027.6) transcript is recommended as the knockout region. The region contains 238bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Taf5* gene. The brief process is as follows: CRISPR/Cas9 system w

- The N-terminal of *Taf5* gene will remain several amino acids ,it may remain the partial function of *Taf5* gene.
- The *Taf5* gene is located on the Chr19. 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)

Taf5 TATA-box binding protein associated factor 5 [*Mus musculus* (house mouse)]

Gene ID: 226182, updated on 10-Oct-2019

Summary

- Official Symbol** Taf5 provided by [MGI](#)
- Official Full Name** TATA-box binding protein associated factor 5 provided by [MGI](#)
- Primary source** [MGI:MGI:2442144](#)
- See related** [Ensembl:ENSMUSG00000025049](#)
- 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** AV117817; 6330528C20Rik
- Expression** Ubiquitous expression in CNS E14 (RPKM 3.8), CNS E11.5 (RPKM 3.7) and 28 other tissues [See more](#)
- Orthologs** [human](#) [all](#)

Genomic context

Location: 19; 19 C3

See Taf5 in [Genome Data Viewer](#)

Exon count: 11

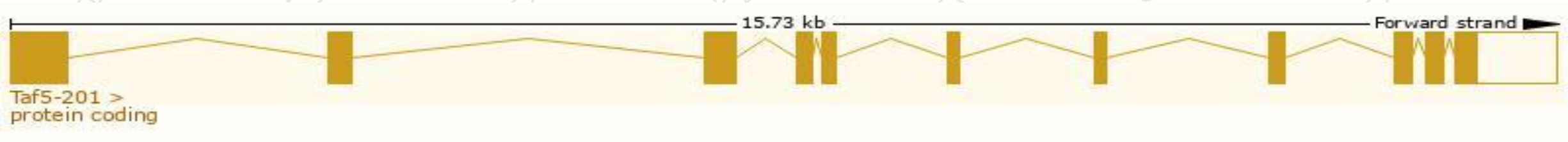
Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	19	NC_000085.6 (47067748..47083479)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	19	NC_000085.5 (47142238..47157969)

Transcript information (Ensembl)

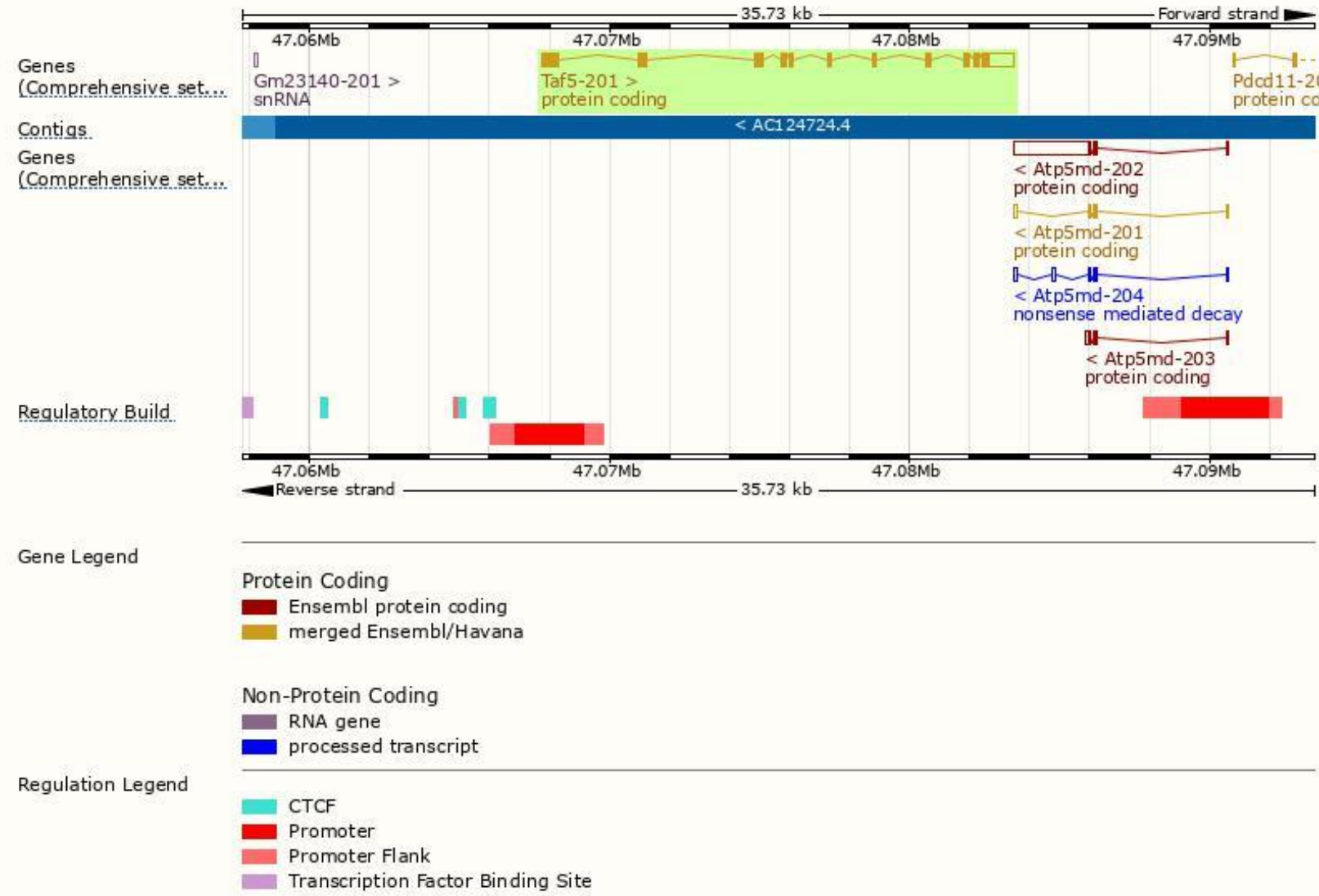
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Taf5-201	ENSMUST00000026027.6	3260	801aa	Protein coding	CCDS29886	F8VPY2	TSL:1 GENCODE basic APPRIS P1

The strategy is based on the design of *Taf5-201* transcript, The transcription is shown below



Genomic location distribution



Protein domain



集萃药康
GemPharmatech



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

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