

# ***Tbx15*** Cas9-KO Strategy

**Designer:**

**Daohua Xu**

**Reviewer:**

**Huimin Su**

**Design Date:**

**2019-8-30**

# Project Overview

**Project Name**

***Tbx15***

**Project type**

**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Tbx15* gene. The schematic diagram is as follows:



- The *Tbx15* gene has 3 transcripts. According to the structure of *Tbx15* gene, exon2 of *Tbx15-201* (ENSMUST00000029462.9) transcript is recommended as the knockout region. The region contains 214bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Tbx15* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Homozygous mutants have low set ears that project laterally, skeletal abnormalities and distinctive dorsoventral coat color patterning.
- The *Tbx15* gene is located on the Chr3. 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)

## Tbx15 T-box 15 [Mus musculus (house mouse)]

Gene ID: 21384, updated on 31-Jan-2019

### Summary



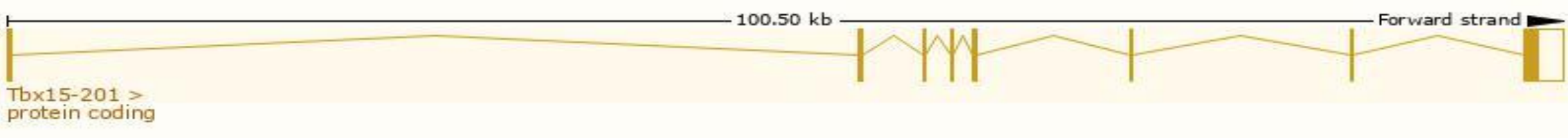
<b>Official Symbol</b>	Tbx15 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	T-box 15 provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:1277234</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000027868</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	VALIDATED
<b>Organism</b>	<a href="#">Mus musculus</a>
<b>Lineage</b>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
<b>Also known as</b>	Tbx14, Tbx8, de
<b>Expression</b>	Biased expression in limb E14.5 (RPKM 21.8), mammary gland adult (RPKM 8.5) and 2 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

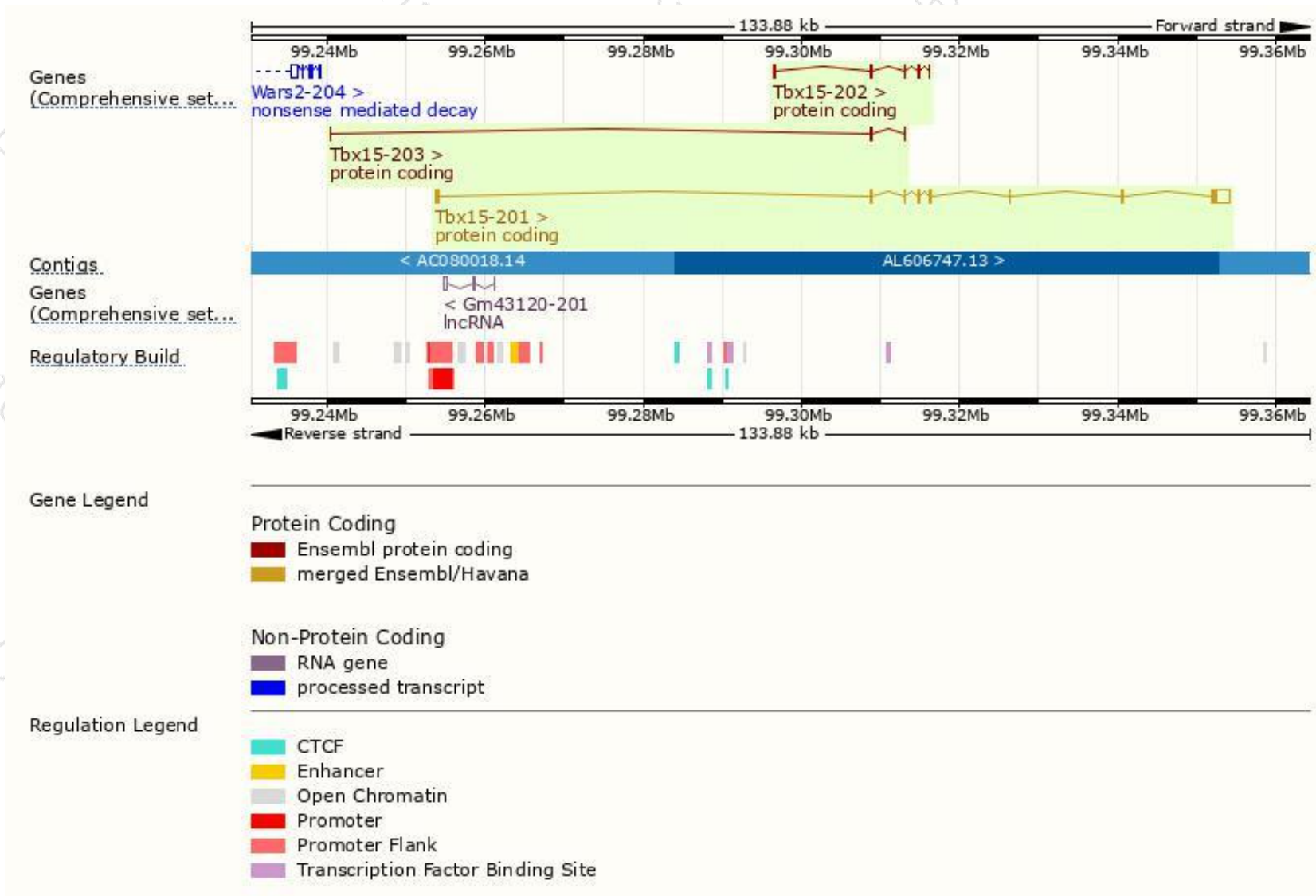
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Tbx15-201	<a href="#">ENSMUST00000029462.9</a>	3566	<a href="#">602aa</a>	Protein coding	<a href="#">CCDS17673</a>	<a href="#">O70306</a>	TSL:1 Gencode basic APPRIS P1
Tbx15-202	<a href="#">ENSMUST00000150756.2</a>	653	<a href="#">141aa</a>	Protein coding	-	<a href="#">A0A0G2JDG4</a>	CDS 3' incomplete TSL:3
Tbx15-203	<a href="#">ENSMUST00000151606.7</a>	333	<a href="#">51aa</a>	Protein coding	-	<a href="#">A0A0G2JG43</a>	CDS 3' incomplete TSL:3

The strategy is based on the design of *Tbx15-201* transcript,The transcription is shown below

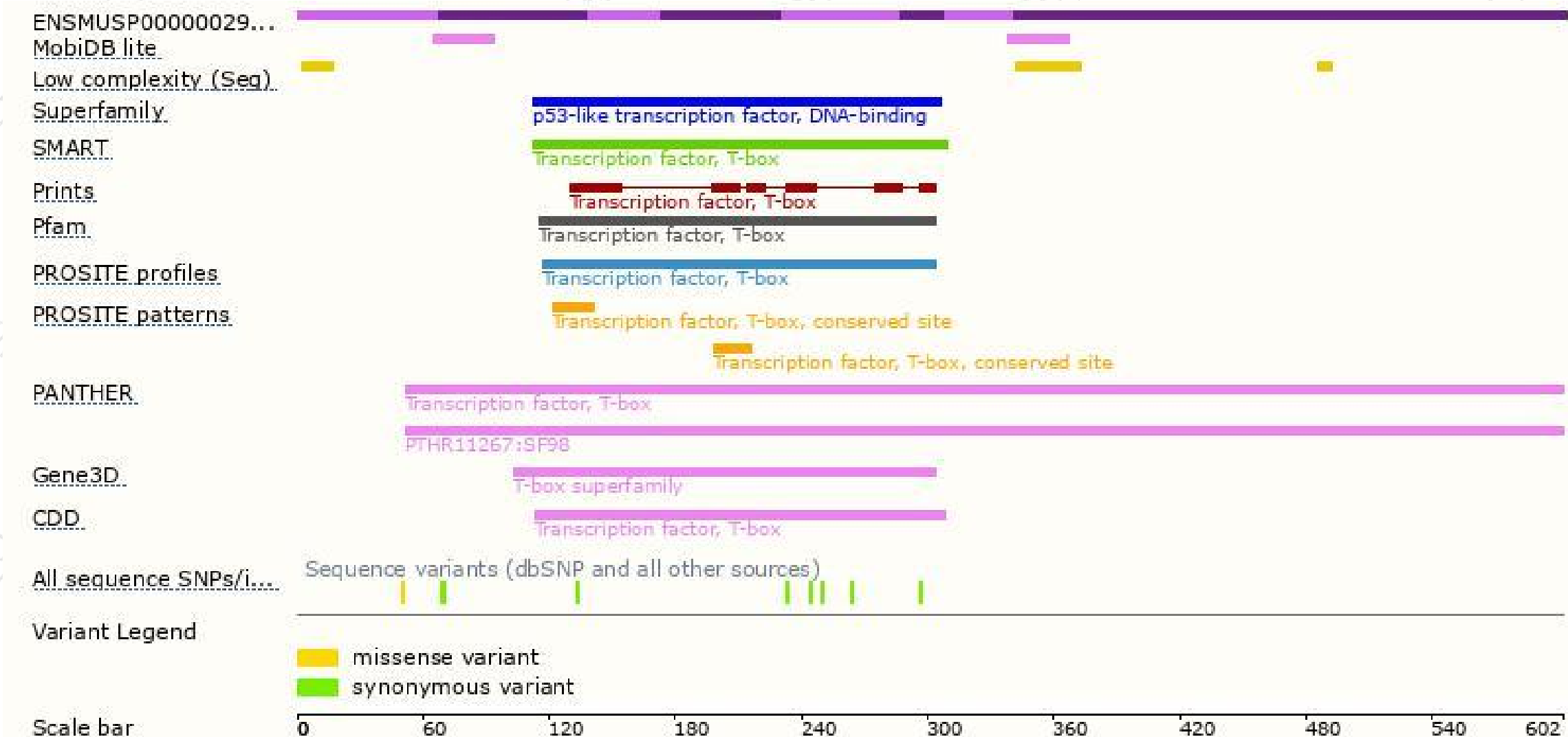


# Genomic location distribution

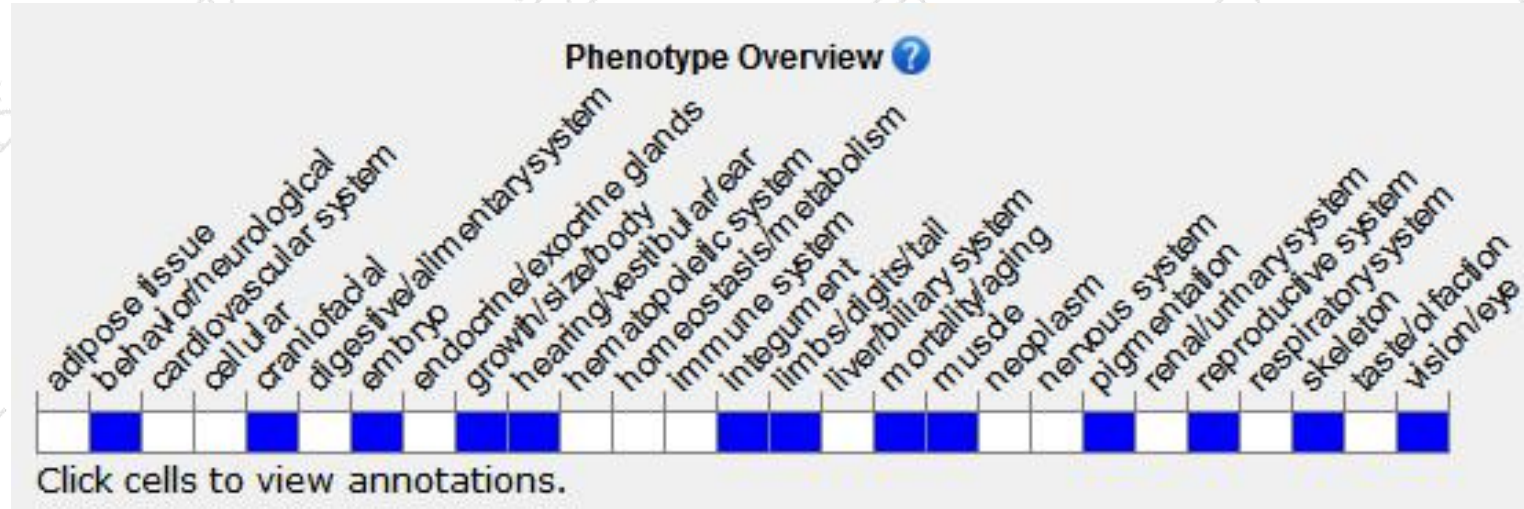




# Protein domain



# 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, Homozygous mutants have low set ears that project laterally, skeletal abnormalities and distinctive dorsoventral coat color patterning.

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

Tel: 400-9660890

