

***Tex101* Cas9-KO Strategy**

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Project Overview

Project Name

Tex101

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Tex101* gene. The schematic diagram is as follows:



- The *Tex101* gene has 3 transcripts. According to the structure of *Tex101* gene, exon2-exon4 of *Tex101-201* (ENSMUST00000078001.6) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Tex101* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Male mice homozygous for a knockout allele of this marker are infertile due to the failure of sperm to migrate into the oviduct.
- The *Tex101* gene is located on the Chr7. 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)

Tex101 testis expressed gene 101 [Mus musculus (house mouse)]

Gene ID: 56746, updated on 31-Jan-2019

Summary



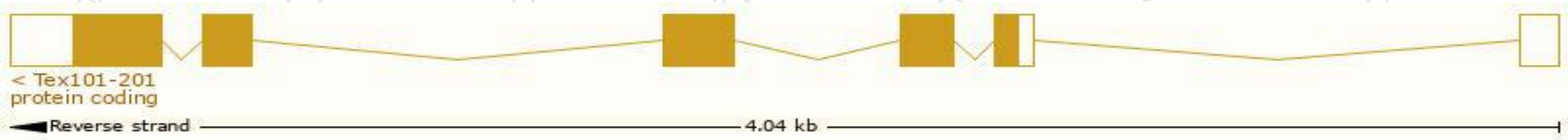
Official Symbol	Tex101 provided by MGI
Official Full Name	testis expressed gene 101 provided by MGI
Primary source	MGI:MGI:1930791
See related	Ensembl:ENSMUSG00000062773
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	1700008H15Rik, AI429076, TES101RP
Expression	Restricted expression toward testis adult (RPKM 185.4) See more
Orthologs	human all

Transcript information (Ensembl)

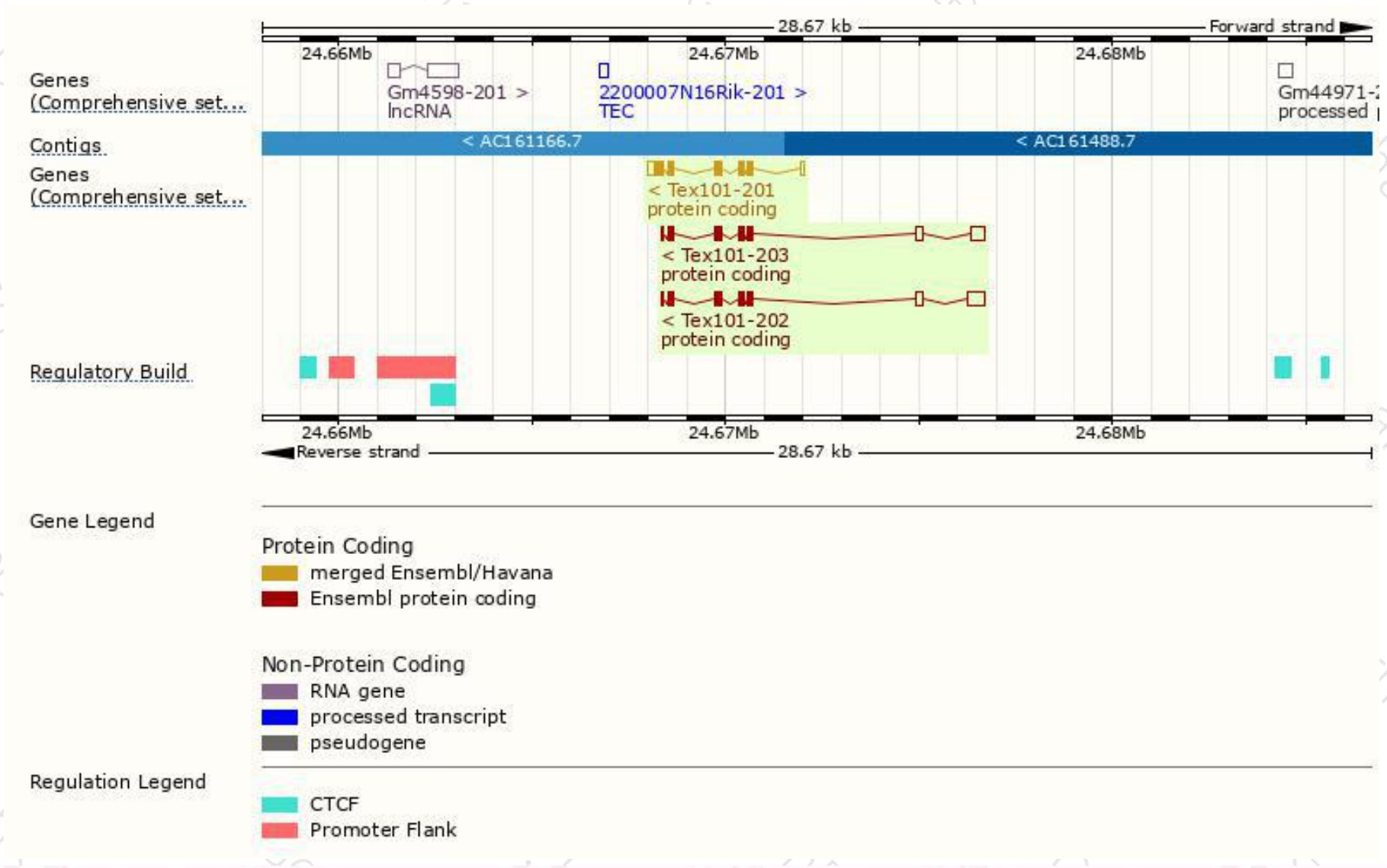
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Tex101-201	ENSMUST00000078001.6	1059	250aa	Protein coding	CCDS20959	Q9JMI7	TSL:1 GENCODE basic APPRIS P1
Tex101-202	ENSMUST00000205488.1	1220	194aa	Protein coding	-	Q076U5	CDS 3' incomplete TSL:1
Tex101-203	ENSMUST00000206390.1	1161	194aa	Protein coding	-	Q076U5	CDS 3' incomplete TSL:1

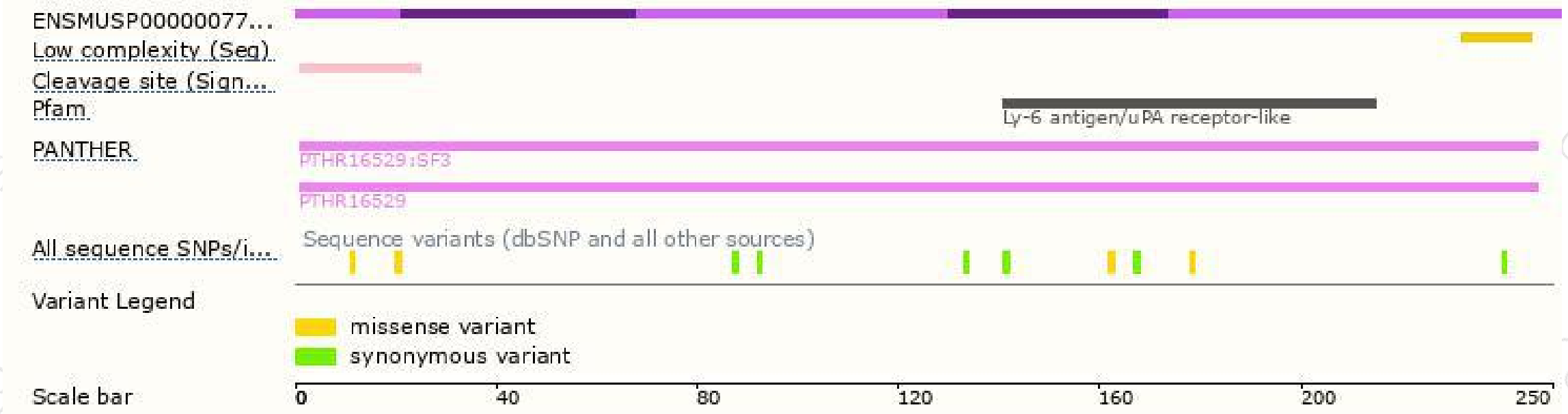
The strategy is based on the design of *Tex101-201* transcript,The transcription is shown below



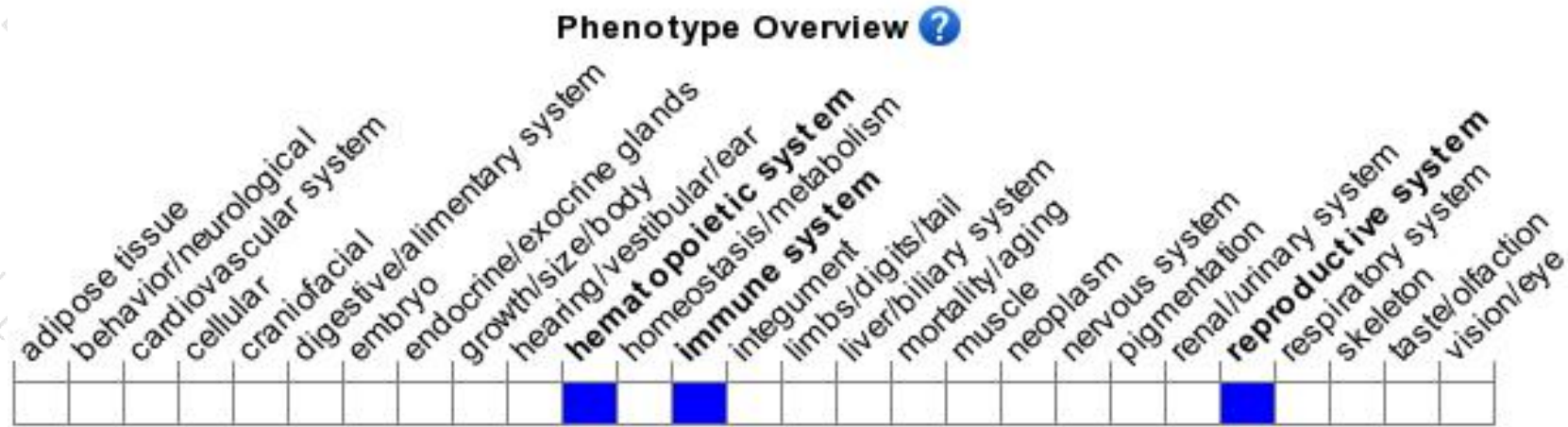
Genomic location distribution



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, Male mice homozygous for a knockout allele of this marker are infertile due to the failure of sperm to migrate into the oviduct.

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

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