

# Tnks1bp1 Cas9-KO Strategy

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Reviewer: Xiaojing Li

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# **Project Overview**



**Project Name** 

Tnks1bp1

**Project type** 

Cas9-KO

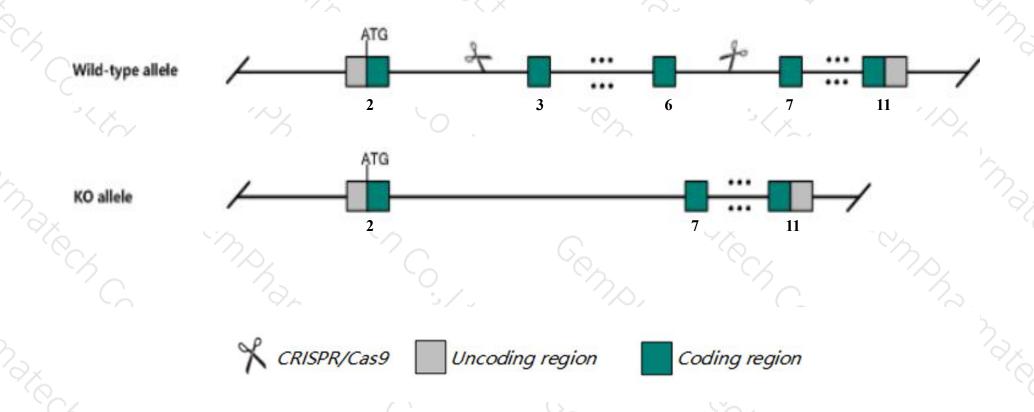
Strain background

C57BL/6JGpt

# **Knockout strategy**



This model will use CRISPR/Cas9 technology to edit the *Tnks1bp1* gene. The schematic diagram is as follows:



### **Technical routes**



- The *Tnks1bp1* gene has 8 transcripts. According to the structure of *Tnks1bp1* gene, exon3-exon6 of *Tnks1bp1-202*(ENSMUST00000111605.8) transcript is recommended as the knockout region. The region contains 4192bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Tnks1bp1* gene. The brief process is as follows: CRISPR/Cas9 system were microinjected into the fertilized eggs of C57BL/6JGpt mice. Fertilized eggs were transplanted to obtain positive F0 mice which were confirmed by PCR and sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6JGpt mice.

### **Notice**



- > The *Tnks1bp1* gene is located on the Chr2. 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)



#### Tnks1bp1 tankyrase 1 binding protein 1 [ Mus musculus (house mouse) ]

Gene ID: 228140, updated on 25-Sep-2020

#### Summary

☆ ?

Official Symbol Tnks1bp1 provided by MGI

Official Full Name tankyrase 1 binding protein 1 provided by MGI

Primary source MGI:MGI:2446193

See related Ensembl: ENSMUSG00000033955

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 TAB18; Tab182; mKIAA1741

Expression Ubiquitous expression in lung adult (RPKM 19.5), colon adult (RPKM 15.5) and 28 other tissues See more

Orthologs <u>human</u> all

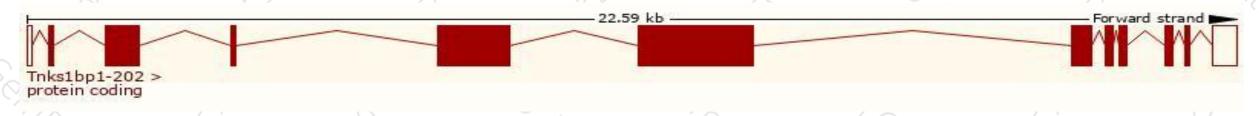
## Transcript information (Ensembl)



The gene has 8 transcripts, all transcripts are shown below:

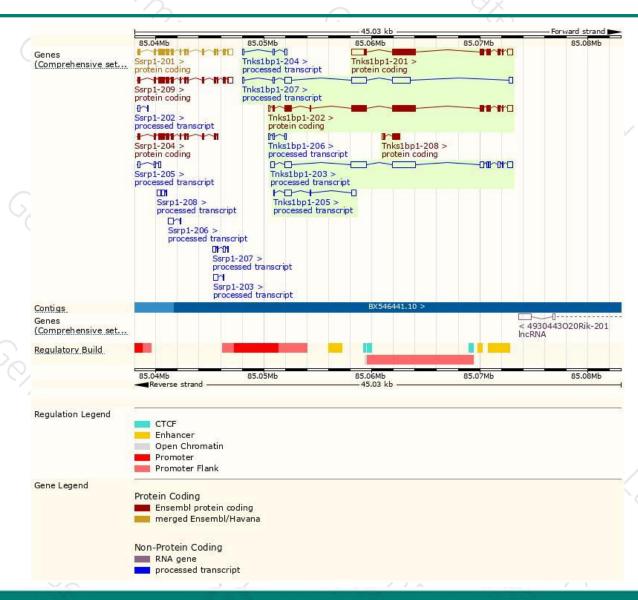
Niema	Transcript ID	h	Destric	Distress	CCDC	UniProt	Flore
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Tnks1bp1-202	ENSMUST00000111605.8	5747	<u>1720aa</u>	Protein coding	CCDS38166	P58871	TSL:1 GENCODE basic APPRIS P2
Tnks1bp1-201	ENSMUST00000048400.3	4799	<u>1058aa</u>	Protein coding	-	Z4YJL4	TSL:1 GENCODE basic APPRIS ALT2
Tnks1bp1-208	ENSMUST00000238769.1	939	276aa	Protein coding	10	2	CDS 3' incomplete
Tnks1bp1-203	ENSMUST00000126309.7	5885	No protein	Processed transcript	15	-	TSL:5
Tnks1bp1-207	ENSMUST00000151092.7	4199	No protein	Processed transcript	12	-	TSL:5
Tnks1bp1-205	ENSMUST00000148682.1	1214	No protein	Processed transcript	150	-	TSL:5
Tnks1bp1-204	ENSMUST00000139915.1	433	No protein	Processed transcript	-	-	TSL:2
Tnks1bp1-206	ENSMUST00000150200.7	342	No protein	Processed transcript	2	ġ.	TSL:2
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The strategy is based on the design of *Tnks1bp1-202* transcript, the transcription is shown below:



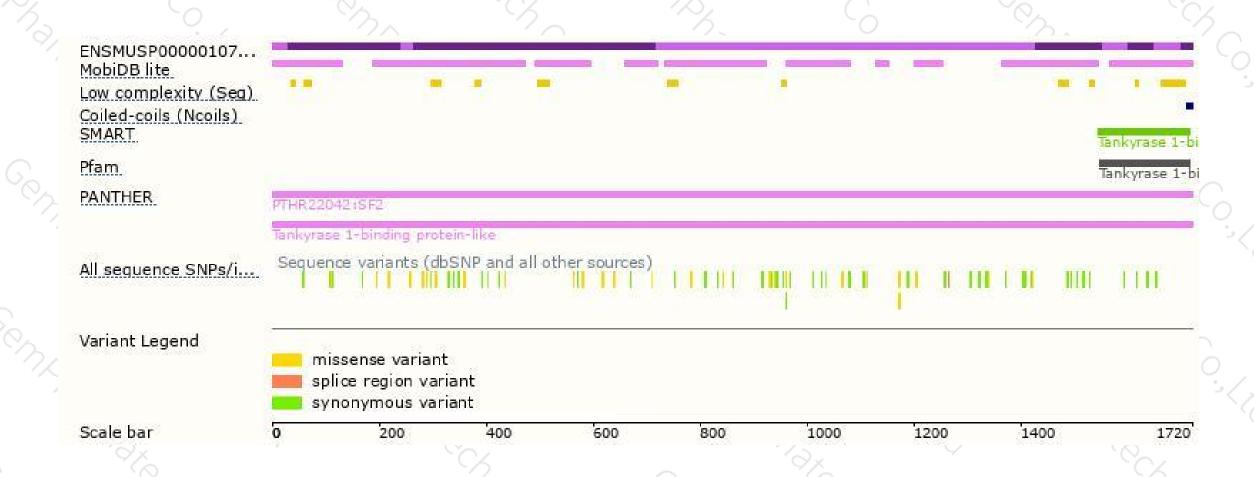
### Genomic location distribution





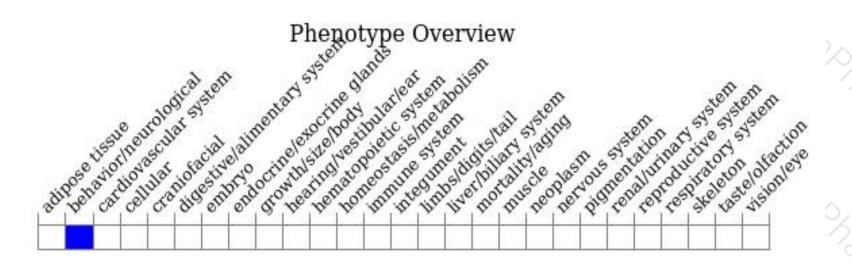
### Protein domain





# Mouse phenotype description(MGI)





Phenotypes affected by the gene are marked in blue.Data quoted from MGI database(http://www.informatics.jax.org/).



If you have any questions, you are welcome to inquire. Tel: 400-9660890





