

Thada Cas9-CKO Strategy

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Reviewer:

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Project Overview

Project Name

Thada

Project type

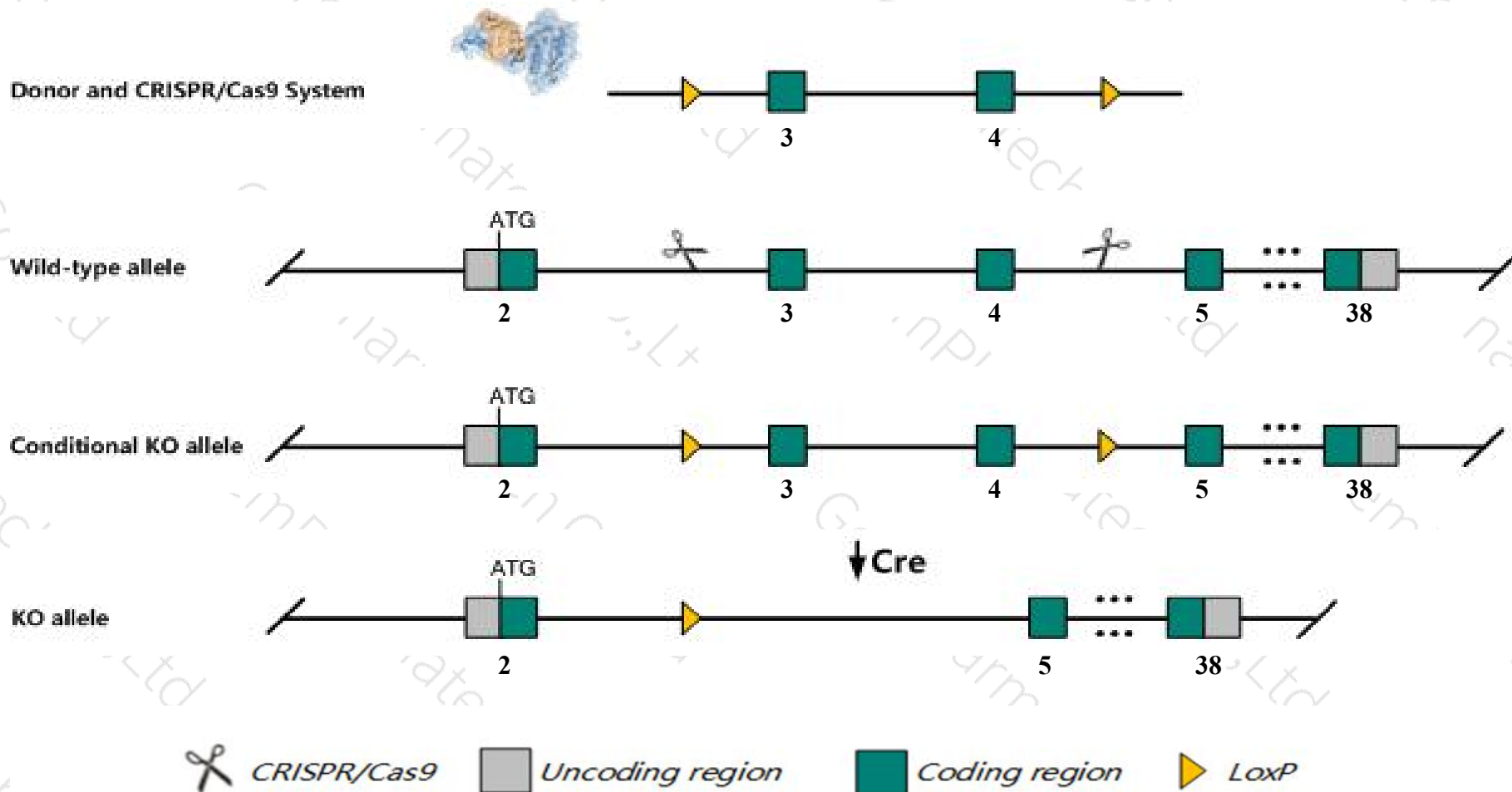
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

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



Technical routes

- The *Thada* gene has 7 transcripts. According to the structure of *Thada* gene, exon3-exon4 of *Thada*-201 (ENSMUST00000047524.9) transcript is recommended as the knockout region. The region contains 226bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Thada* gene. The brief process is as follows: CRISPR/Cas9 system and Donor 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.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

Notice

- The *Thada* gene is located on the Chr17. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)

Thada thyroid adenoma associated [Mus musculus (house mouse)]

Gene ID: 240174, updated on 31-Jan-2019

Summary



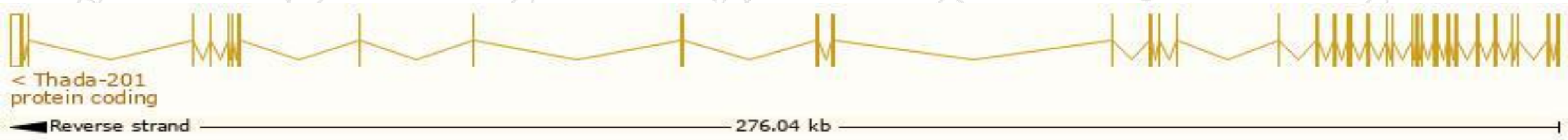
Official Symbol	Thada provided by MGI
Official Full Name	thyroid adenoma associated provided by MGI
Primary source	MGI:MGI:3039623
See related	Ensembl:ENSMUSG00000024251
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	6530405K19, A930021O22, AU019899, BC052885, GITA
Expression	Ubiquitous expression in testis adult (RPKM 2.9), thymus adult (RPKM 2.6) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

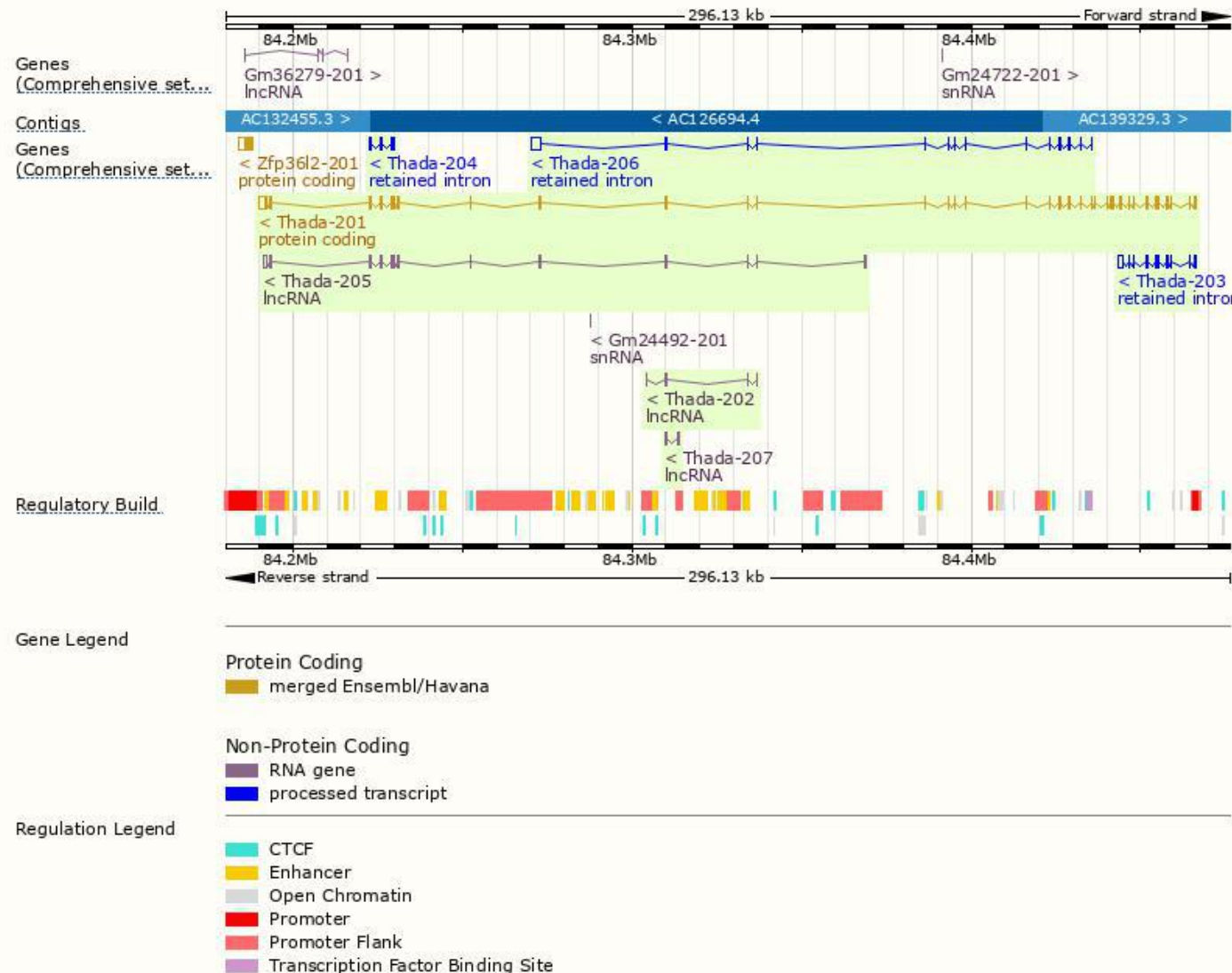
The gene has 7 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Thada-201	ENSMUST00000047524.9	7784	1938aa	Protein coding	CCDS50198	A8C756	TSL:1 GENCODE basic APPRIS P1
Thada-206	ENSMUST00000234724.1	4412	No protein	Retained intron	-	-	
Thada-203	ENSMUST00000234313.1	2397	No protein	Retained intron	-	-	
Thada-204	ENSMUST00000234529.1	608	No protein	Retained intron	-	-	
Thada-205	ENSMUST00000234547.1	2556	No protein	lncRNA	-	-	
Thada-202	ENSMUST00000234134.1	570	No protein	lncRNA	-	-	
Thada-207	ENSMUST00000235126.1	445	No protein	lncRNA	-	-	

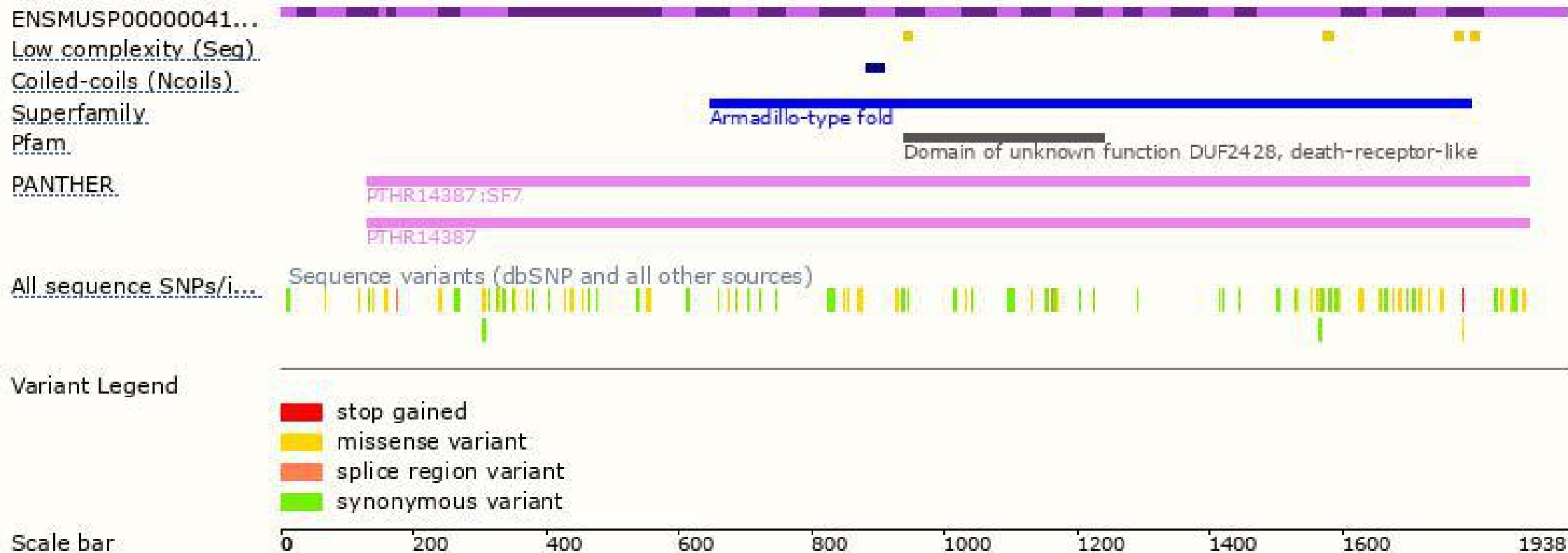
The strategy is based on the design of *Thada-201* transcript,The transcription is shown below



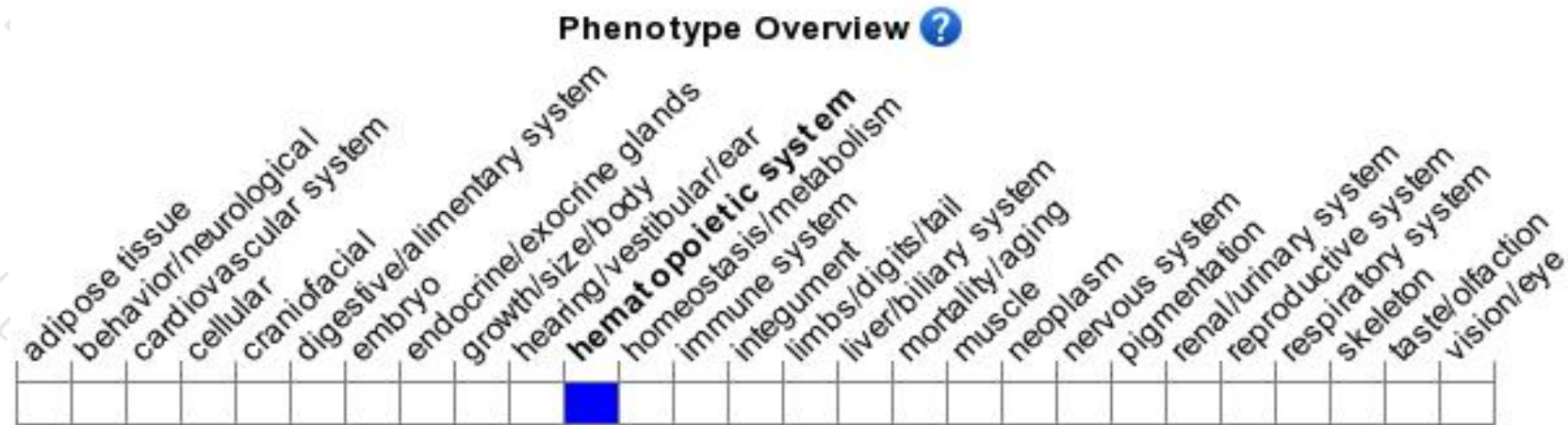
Genomic location distribution



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/>).

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

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