

Wnt7a Cas9-CKO Strategy

Designer:

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Design Date:

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Project Overview

Project Name

Wnt7a

Project type

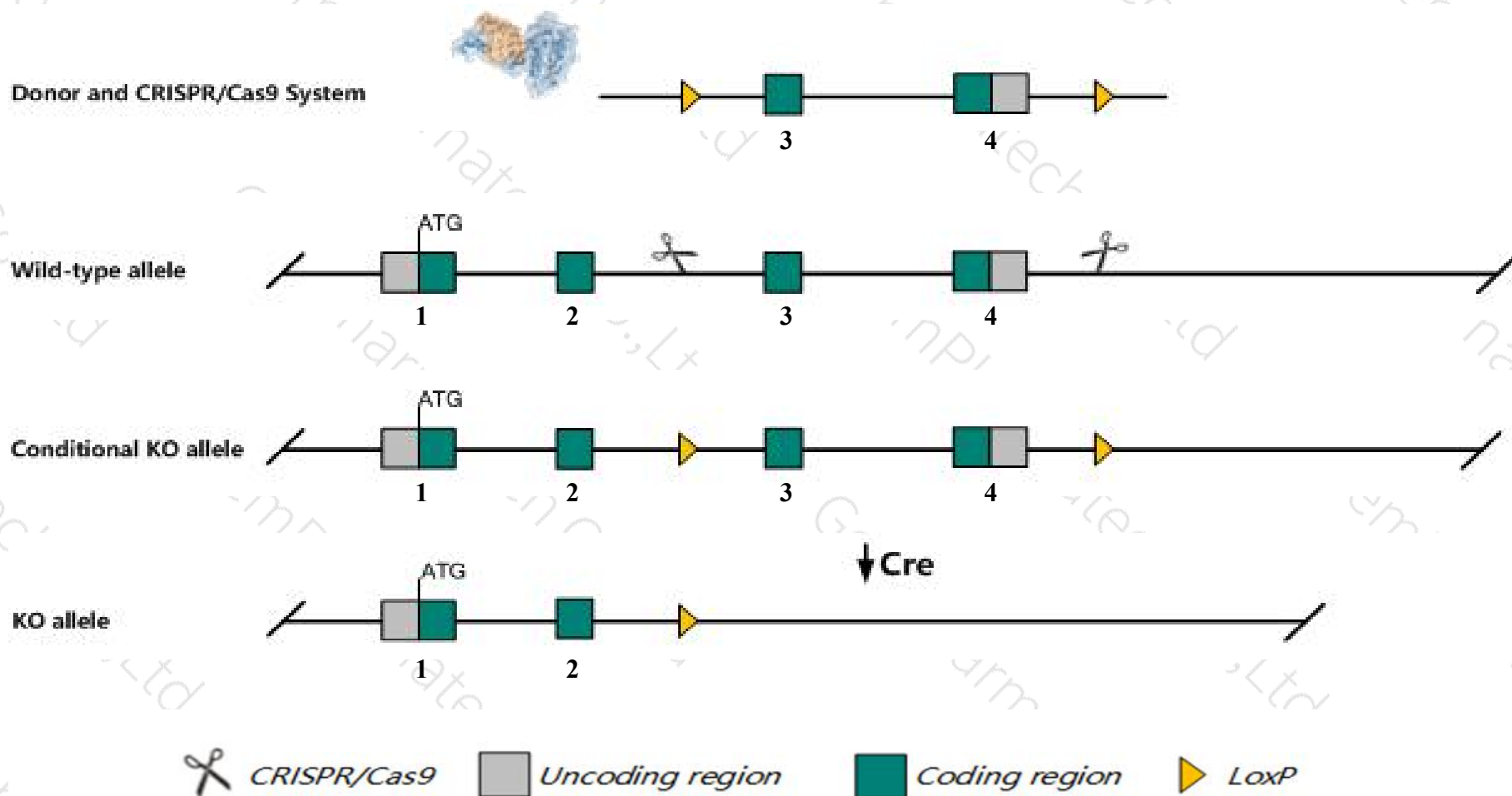
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Wnt7a* gene. The schematic diagram is as follows:



- The *Wnt7a* gene has 3 transcripts. According to the structure of *Wnt7a* gene, exon3-exon4 of *Wnt7a-201* (ENSMUST00000032180.6) transcript is recommended as the knockout region. The region contains 752bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Wnt7a* 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.

- According to the existing MGI data, Homozygous mutants have skeletal abnormalities including absence of digits, especially of the forelimb, and sometimes absence of the ulna. Occasionally, there is an extra set of ribs. Both sexes are sterile due to abnormalities of the Mullerian duct.
- The *Wnt7a* gene is located on the Chr6. 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)

Wnt7a wingless-type MMTV integration site family, member 7A [Mus musculus (house mouse)]

Gene ID: 22421, updated on 5-Mar-2019

Summary



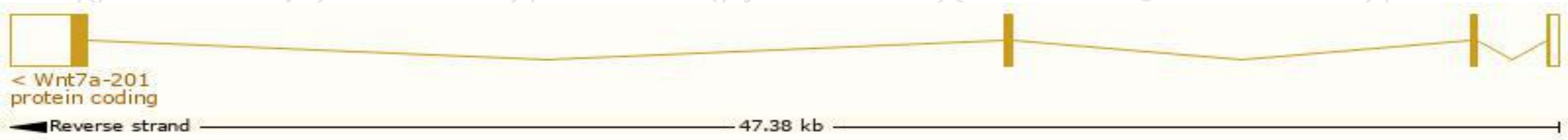
Official Symbol	Wnt7a provided by MGI
Official Full Name	wingless-type MMTV integration site family, member 7A provided by MGI
Primary source	MGI:MGI:98961
See related	Ensembl:ENSMUSG00000030093
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	AI849442, Wnt-7a, px, tw
Expression	Biased expression in cerebellum adult (RPKM 11.5), CNS E18 (RPKM 6.0) and 10 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

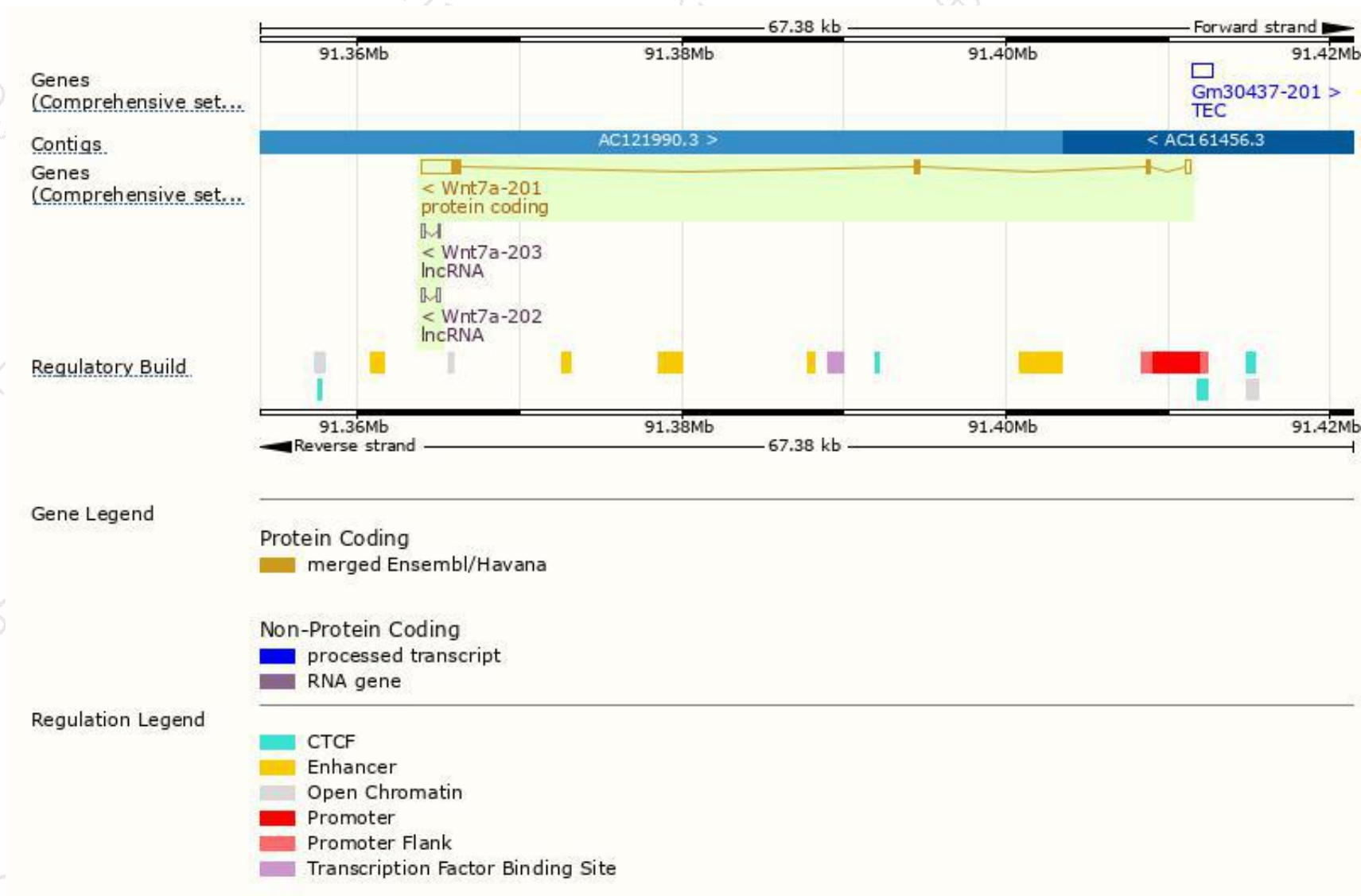
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Wnt7a-201	ENSMUST00000032180.6	3172	349aa	Protein coding	CCDS39568	P24383	TSL:1 GENCODE basic APPRIS P1
Wnt7a-202	ENSMUST00000132936.2	297	No protein	lncRNA	-	-	TSL:5
Wnt7a-203	ENSMUST00000133092.2	293	No protein	lncRNA	-	-	TSL:3

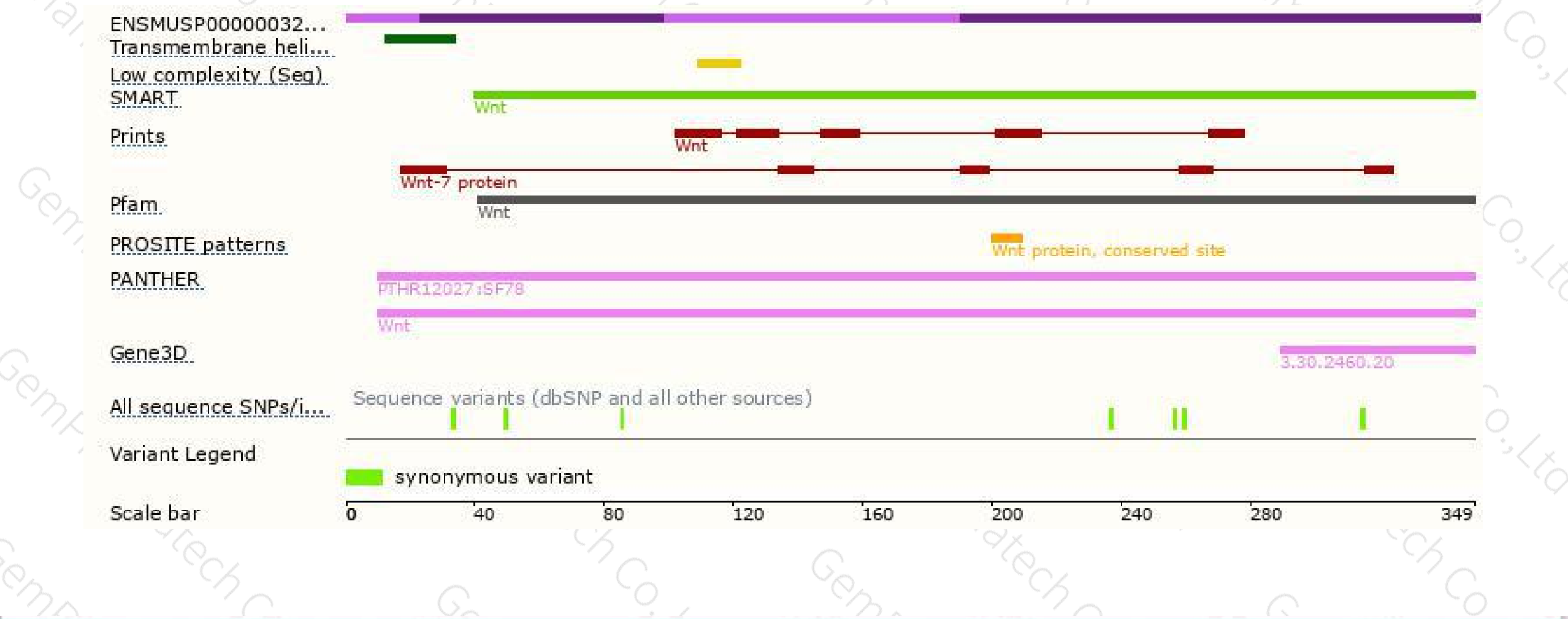
The strategy is based on the design of *Wnt7a-201* transcript,The transcription is shown below



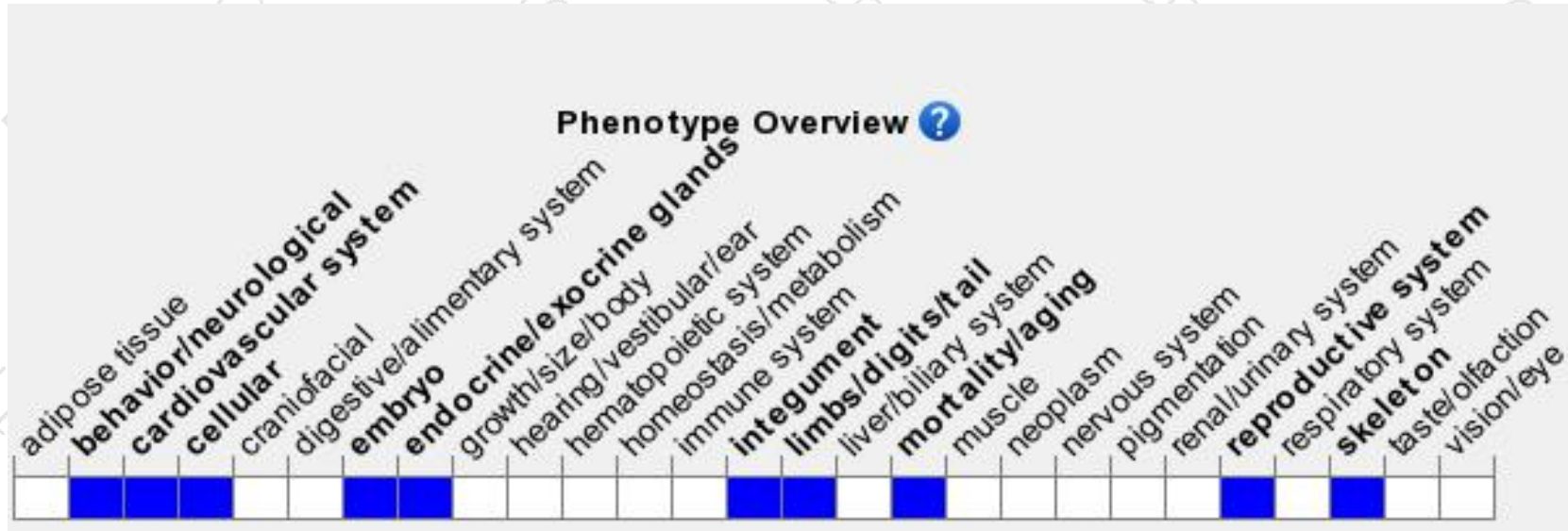
Genomic location distribution



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 skeletal abnormalities including absence of digits, especially of the forelimb, and sometimes absence of the ulna. Occasionally, there is an extra set of ribs. Both sexes are sterile due to abnormalities of the Mullerian duct.

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

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