

Wnt9b Cas9-CKO Strategy

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Reviewer:

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Design Date:

2019-12-11

Project Overview



Project Name

Wnt9b

Project type

Cas9-CKO

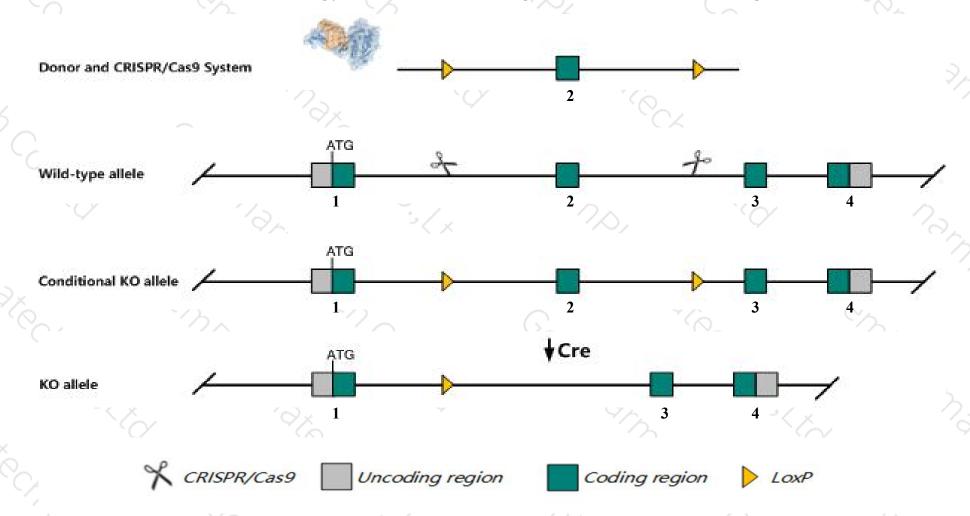
Strain background

C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Wnt9b* gene. The schematic diagram is as follows:



Technical routes



- The *Wnt9b* gene has 1 transcript. According to the structure of *Wnt9b* gene, exon2 of *Wnt9b-201*(ENSMUST00000018630.2) transcript is recommended as the knockout region. The region contains 257bp coding sequence.

 Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Wnt9b* 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



- ➤ According to the existing MGI data, Homozygous null mice display neonatal lethality, disrupted ureteric bud branching, impaired Mullerian duct formation, and incompletely penetrant cleft lip and palate. In mice with alleles that decrease expression kidneys are smaller with fewer mature nephrons.
- > The *Wnt9b* gene is located on the Chr11. 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)



Wnt9b wingless-type MMTV integration site family, member 9B [Mus musculus (house mouse)]

Gene ID: 22412, updated on 19-Mar-2019

Summary

☆ ?

Official Symbol Wnt9b provided by MGI

Official Full Name wingless-type MMTV integration site family, member 9B provided by MGI

Primary source MGI:MGI:1197020

See related Ensembl:ENSMUSG00000018486

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 Wnt14b, Wnt15, clf, clf1, wnt-14b, wnt-15

Expression Broad expression in kidney adult (RPKM 1.2), cerebellum adult (RPKM 0.3) and 17 other tissuesSee more

Orthologs <u>human</u> all

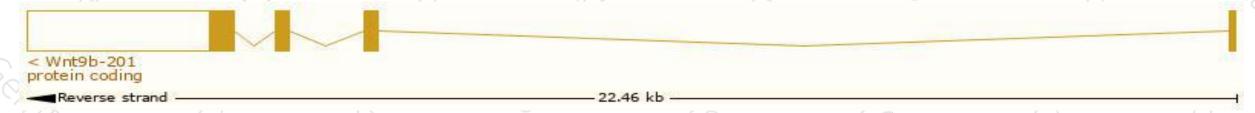
Transcript information (Ensembl)



The gene has 1 transcript, and the transcript is shown below:

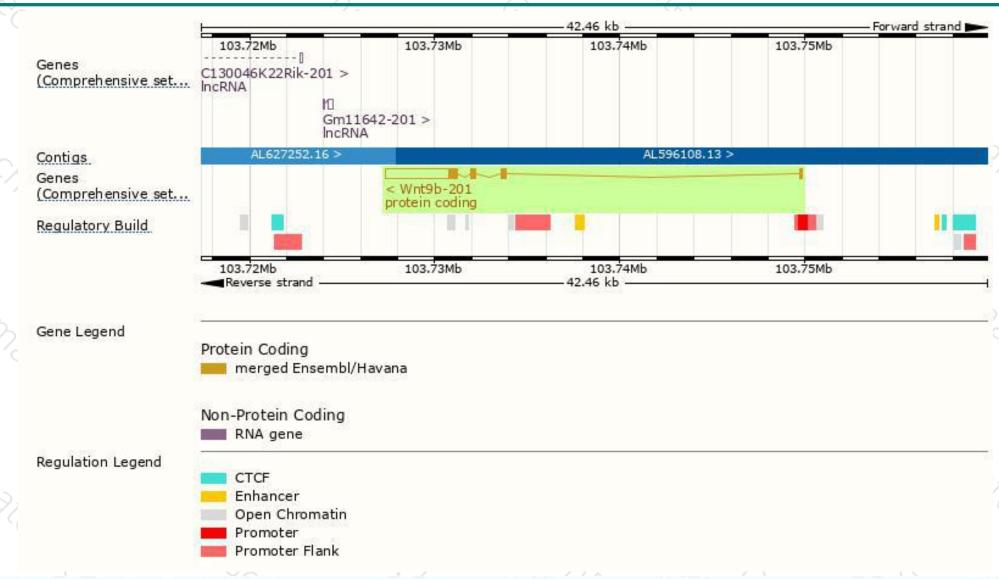
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Wnt9b-201	ENSMUST00000018630.2	4518	<u>359aa</u>	Protein coding	CCDS25522	O35468 Q2TBA6	TSL:1 GENCODE basic APPRIS P1

The strategy is based on the design of Wnt9b-201 transcript, The transcription is shown below



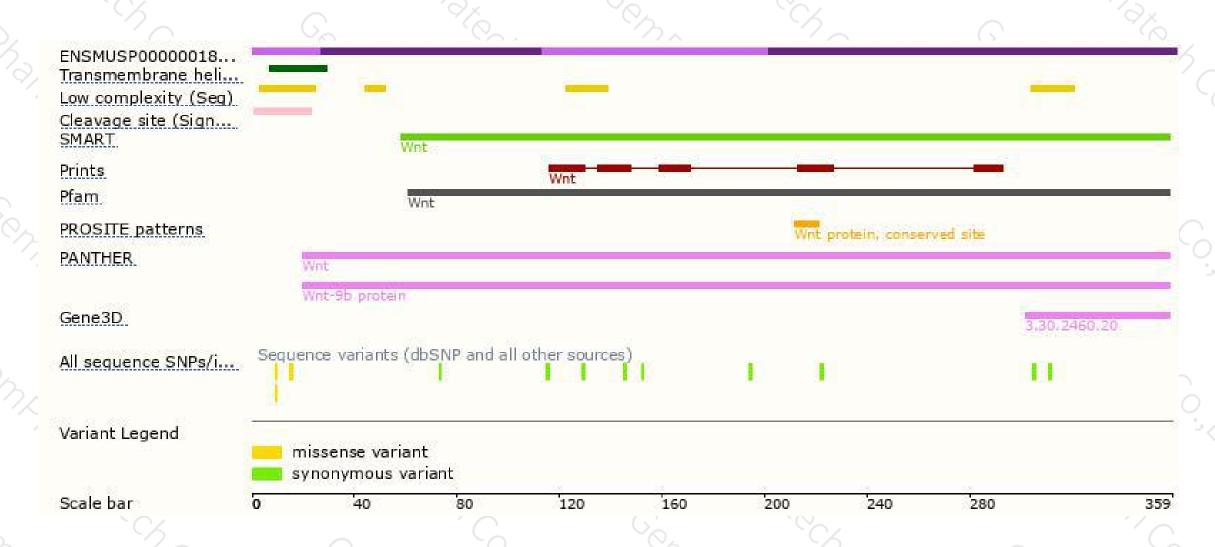
Genomic location distribution





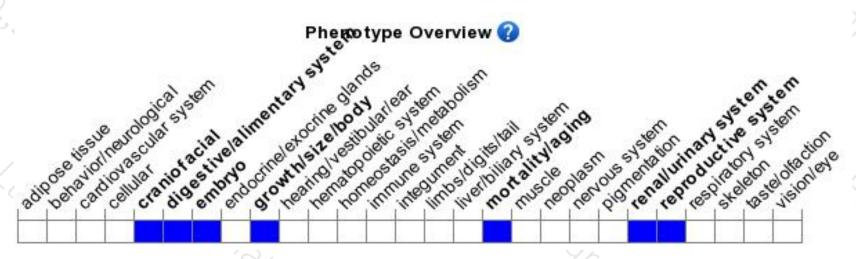
Protein domain





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 null mice display neonatal lethality, disrupted ureteric bud branching, impaired Mullerian duct formation, and incompletely penetrant cleft lip and palate. In mice with alleles that decrease expression kidneys are smaller with fewer mature nephrons.



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





