

Wnt1 Cas9-KO Strategy

Designer:

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Project Overview

Project Name

Wnt1

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Wnt1* gene. The schematic diagram is as follows:



- The *Wnt1* gene has 1 transcript. According to the structure of *Wnt1* gene, exon1-exon4 of *Wnt1-201* (ENSMUST00000023734.7) transcript is recommended as the knockout region. The region contains all of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Wnt1* gene. The brief process is as follows: CRISPR/Cas9 system v

- According to the existing MGI data, In mild form, homozygotes have ataxia and hypertonia, with malformation of anterior cerebellum, deep midline fissure, and impaired fertility. In the severe form, there is virtually no midbrain and cerebellum and mutants die within hours of birth.
- The *Wnt1* gene is located on the Chr15. 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)

Wnt1 wingless-type MMTV integration site family, member 1 [Mus musculus (house mouse)]

Gene ID: 22408, updated on 12-Mar-2019

Summary



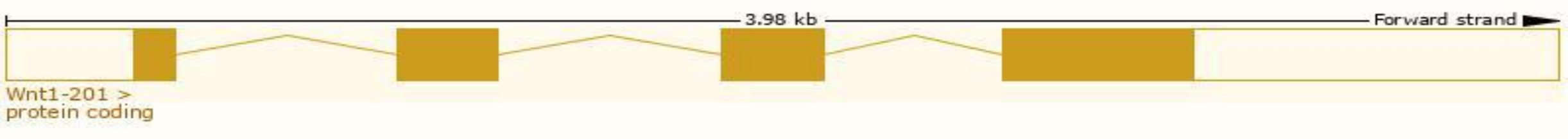
Official Symbol	Wnt1 provided by MGI
Official Full Name	wingless-type MMTV integration site family, member 1 provided by MGI
Primary source	MGI:MGI:98953
See related	Ensembl:ENSMUSG00000022997
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Int-1, Wnt-1, sw, swaying
Expression	Biased expression in CNS E11.5 (RPKM 2.1), testis adult (RPKM 1.8) and 5 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

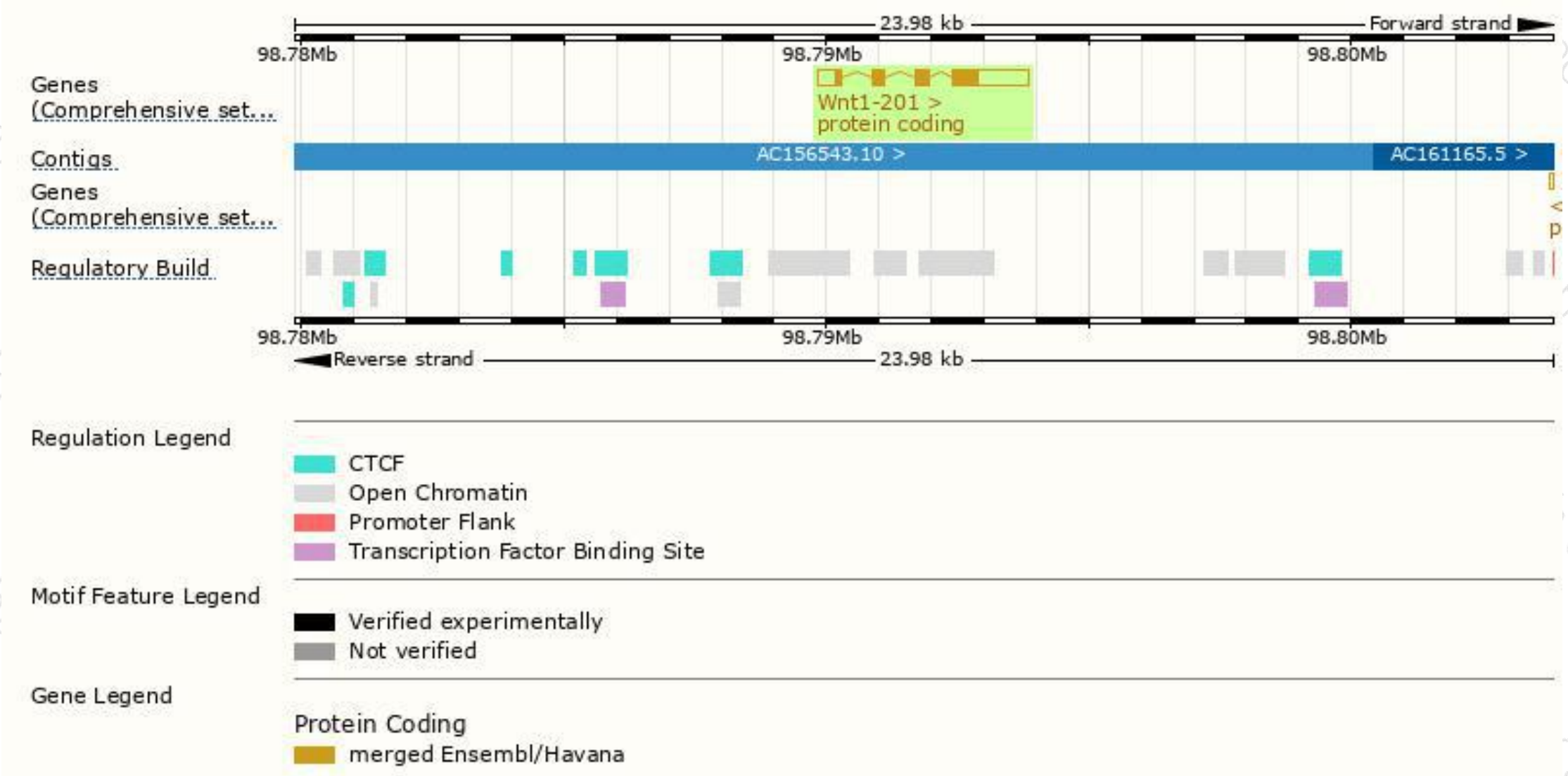
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Wnt1-201	ENSMUST00000023734.7	2378	370aa	Protein coding	CCDS27807	P04426 Q3UR96	TSL:1 GENCODE basic APPRIS P1

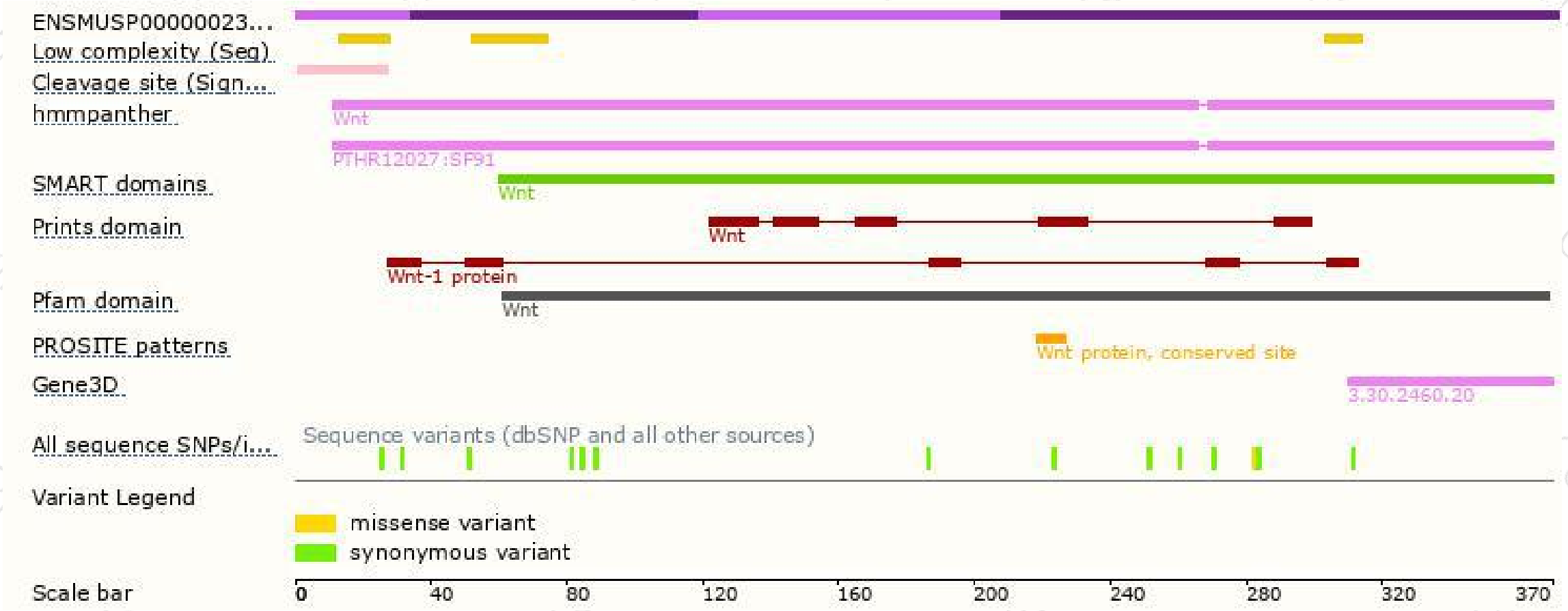
The strategy is based on the design of *Wnt1-201* transcript, The transcription is shown below



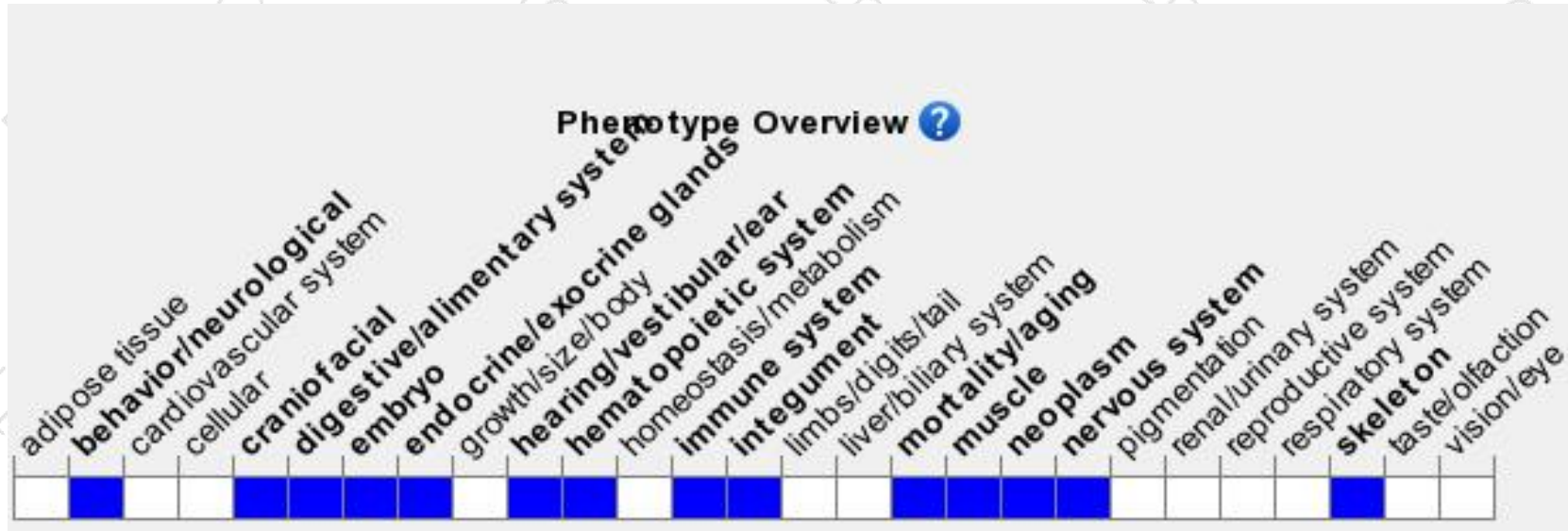
Genomic location distribution



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, In mild form, homozygotes have ataxia and hypertonia, with malformation of anterior cerebellum, deep midline fissure, and impaired fertility. In the severe form, there is virtually no midbrain and cerebellum and mutants die within hours of birth.

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

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