

Hs6st3 Cas9-KO Strategy

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Reviewer:

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Project Overview

Project Name

Hs6st3

Project type

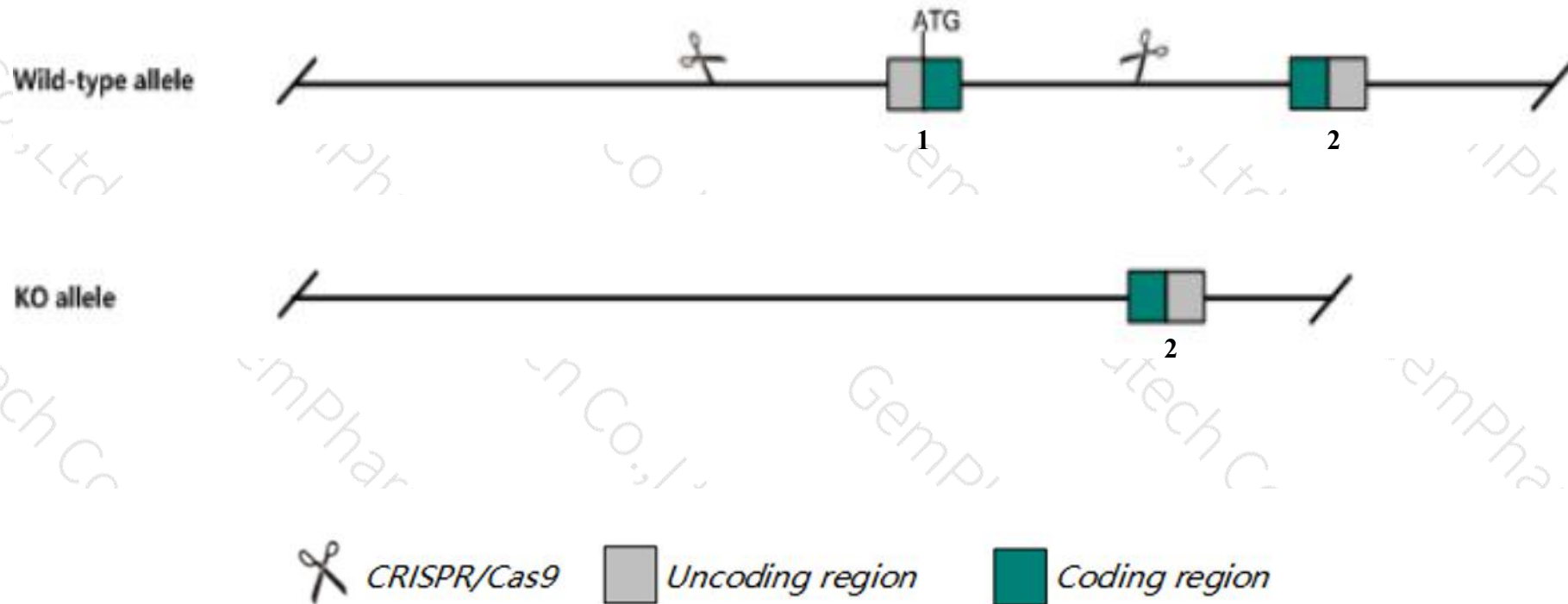
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Hs6st3* gene. The schematic diagram is as follows:



- The *Hs6st3* gene has 1 transcript. According to the structure of *Hs6st3* gene, exon1 of *Hs6st3-201* (ENSMUST00000065904.4) transcript is recommended as the knockout region. The region contains start codon ATG. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Hs6st3* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Hs6st3* gene is located on the Chr14. 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)

Hs6st3 heparan sulfate 6-O-sulfotransferase 3 [*Mus musculus* (house mouse)]

Gene ID: 50787, updated on 13-Mar-2020

Summary

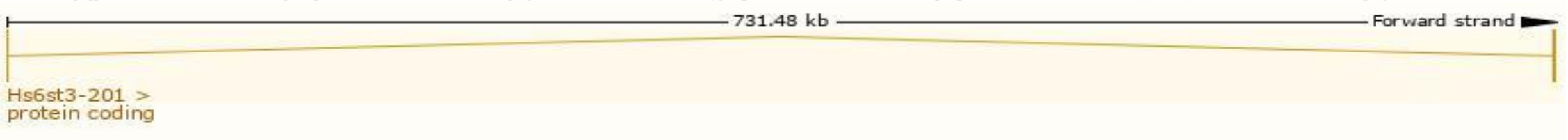
Official Symbol	Hs6st3 provided by MGI
Official Full Name	heparan sulfate 6-O-sulfotransferase 3 provided by MGI
Primary source	MGI:MGI:1354960
See related	Ensembl:ENSMUSG00000053465
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	6OST3; HS6ST-3
Expression	Biased expression in cortex adult (RPKM 1.3), frontal lobe adult (RPKM 0.8) and 7 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

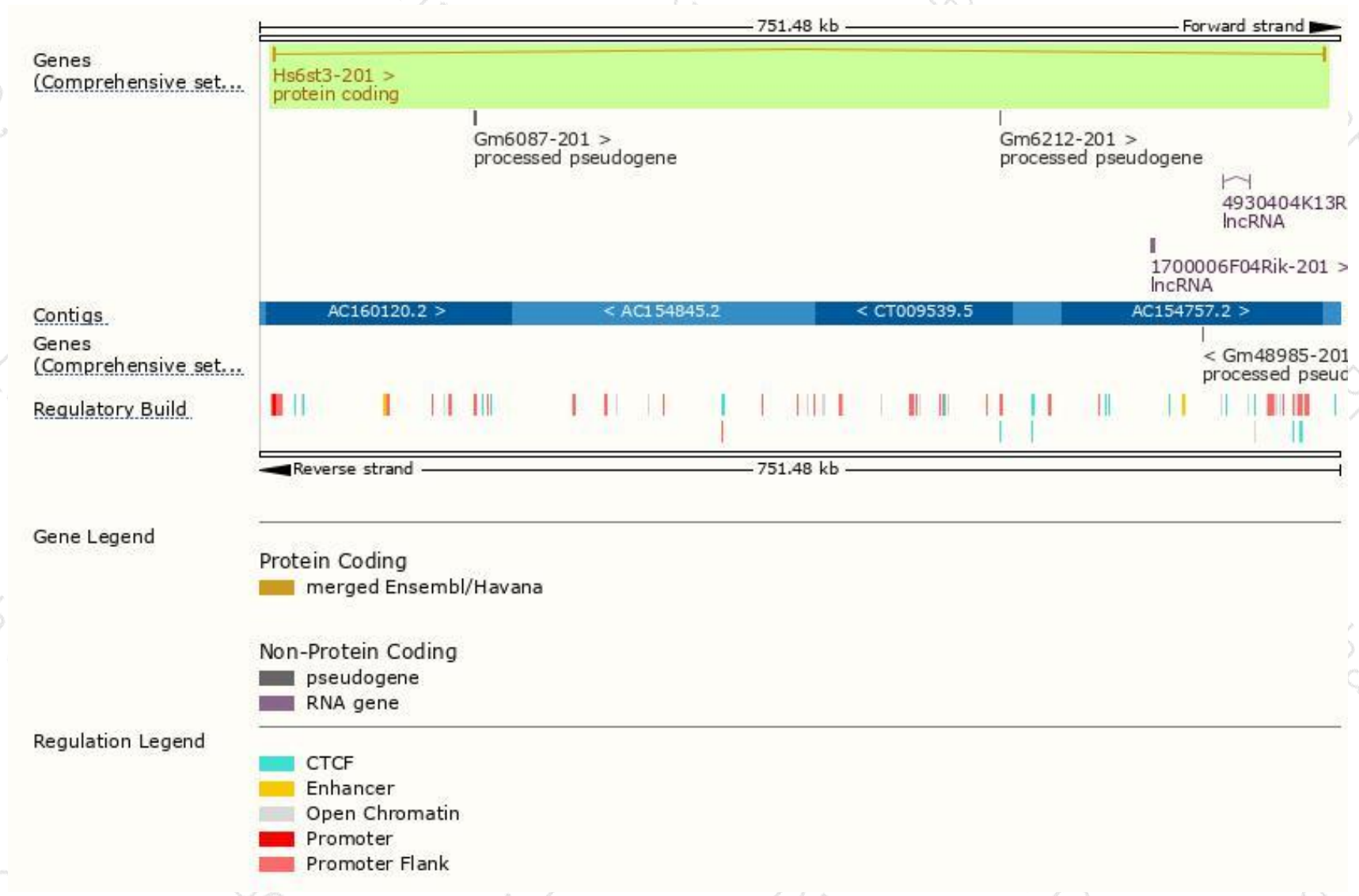
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Hs6st3-201	ENSMUST00000065904.4	1708	470aa	Protein coding	CCDS27339	Q9QYK4	TSL:1 Gencode basic APPRIS P1

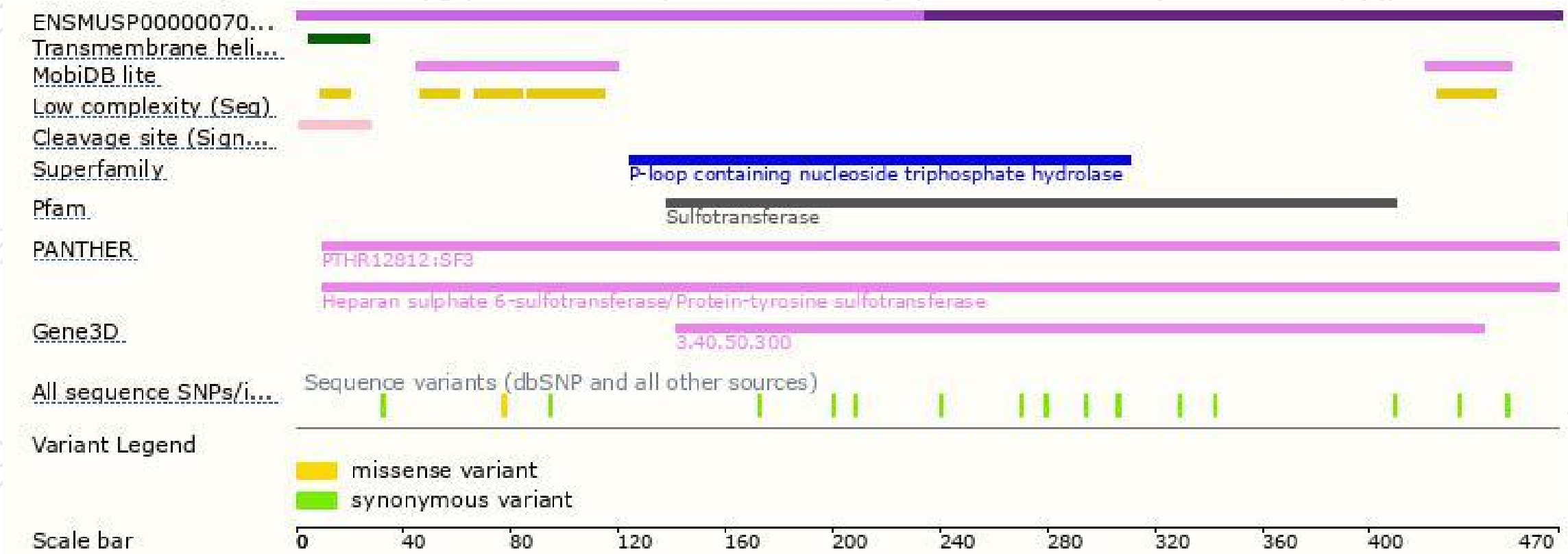
The strategy is based on the design of *Hs6st3-201* transcript, The transcription is shown below



Genomic location distribution



Protein domain



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

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