

Cetn1 Cas9-KO Strategy

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Reviewer:

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Project Overview

Project Name

Cetn1

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Cetn1* gene. The schematic diagram is as follows:



- The *Cetn1* gene has 3 transcripts. According to the structure of *Cetn1* gene, exon1 of *Cetn1-201* (ENSMUST00000062769.6) 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 *Cetn1* gene. The brief process is as follows: CRISPR/Cas9 system

- The *Cetn1* gene is located on the Chr18. 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)

Cetn1 centrin 1 [Mus musculus (house mouse)]

Gene ID: 26369, updated on 9-Apr-2019

Summary



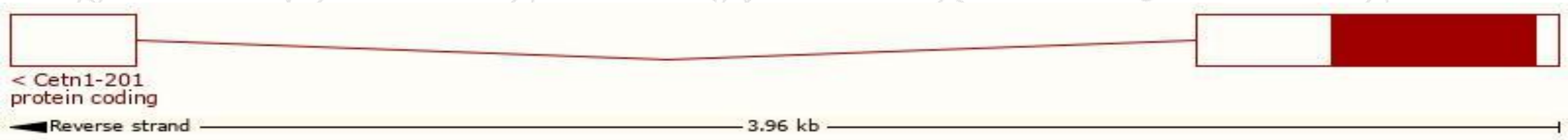
Official Symbol	Cetn1 provided by MGI
Official Full Name	centrin 1 provided by MGI
Primary source	MGI:MGI:1347086
See related	Ensembl:ENSMUSG000000050996
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	caltractin
Orthologs	human all

Transcript information (Ensembl)

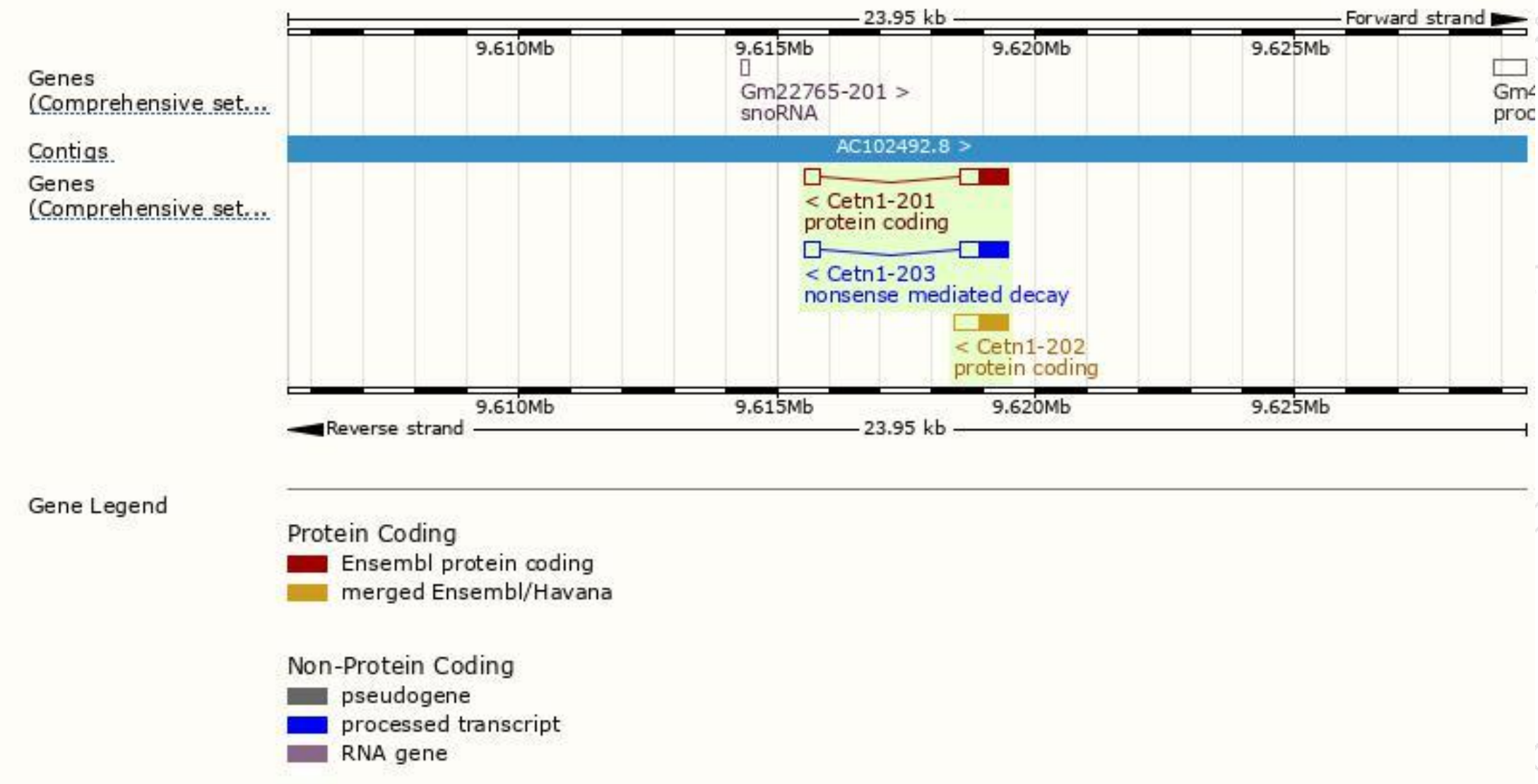
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Cetn1-201	ENSMUST00000062769.6	1247	172aa	Protein coding	CCDS37731	P41209	TSL:1 GENCODE basic APPRIS P1
Cetn1-202	ENSMUST00000234003.1	1051	172aa	Protein coding	CCDS37731	-	GENCODE basic APPRIS P1
Cetn1-203	ENSMUST00000234590.1	1247	172aa	Nonsense mediated decay	CCDS37731	-	

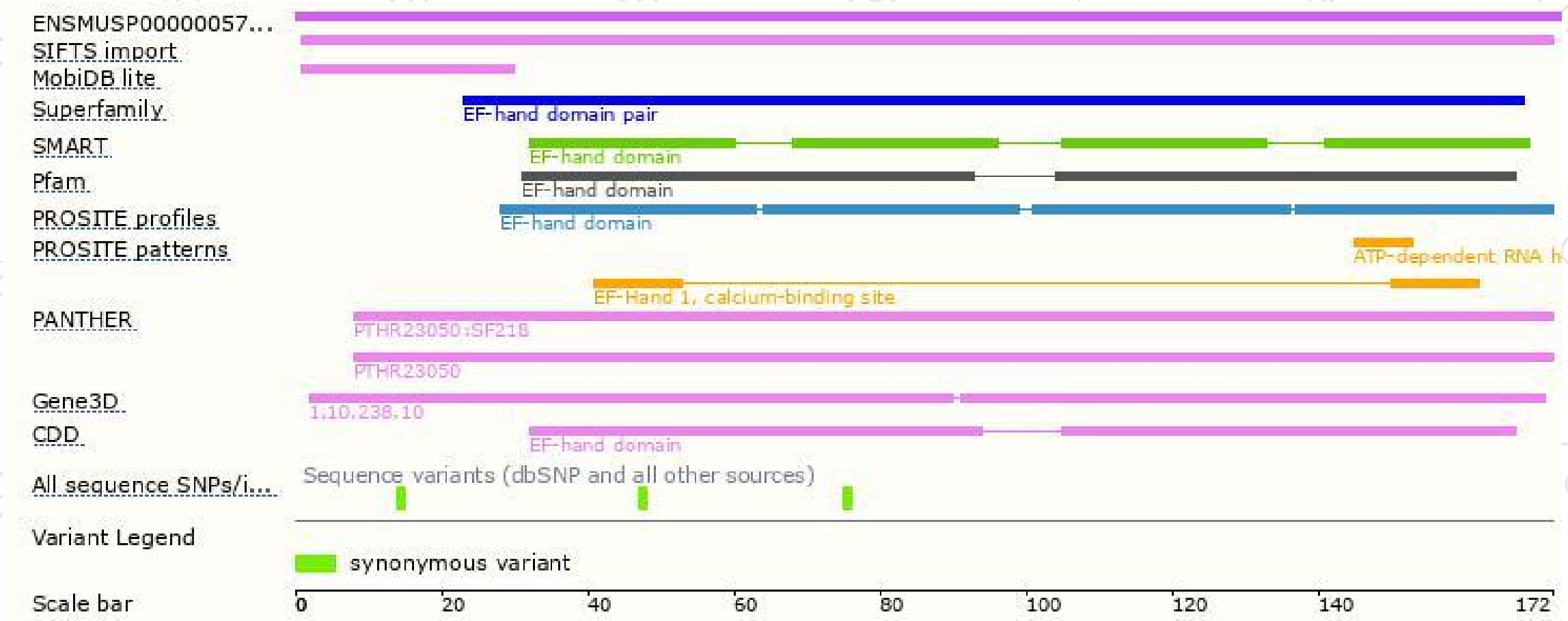
The strategy is based on the design of *Cetn1-201* transcript,The transcription is shown below



Genomic location distribution



Protein domain



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

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