

Catsperd Cas9-KO Strategy

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Project Overview

Project Name

Catsperd

Project type

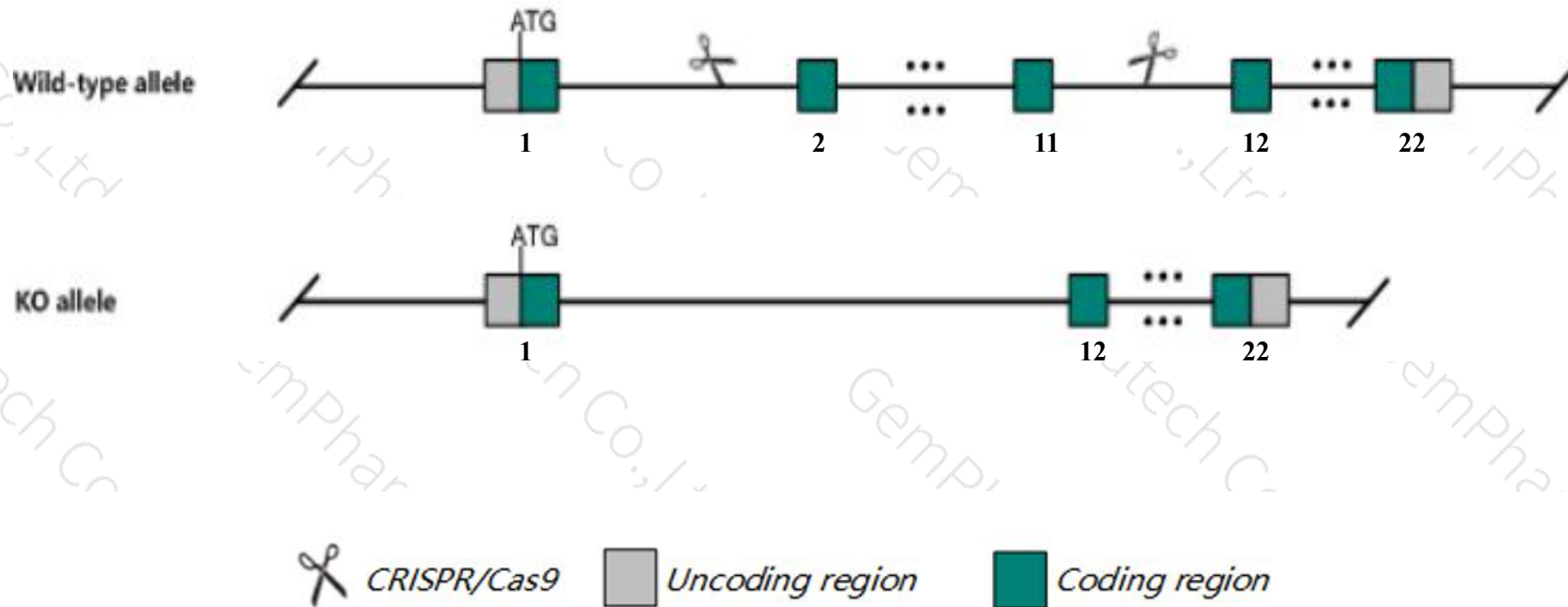
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

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



- The *Catsperd* gene has 3 transcripts. According to the structure of *Catsperd* gene, exon2-exon11 of *Catsperd-201* (ENSMUST00000112979.3) transcript is recommended as the knockout region. The region contains 916bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Catsperd* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, mice homozygous for a deletion in this gene display male infertility. hyperactivity of sperm fails to develop under capacitating conditions.
- The *Catsperd* gene is located on the Chr17. 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)

Catsperd cation channel sperm associated auxiliary subunit delta [Mus musculus (house mouse)]

Gene ID: 106757, updated on 13-Mar-2020

Summary



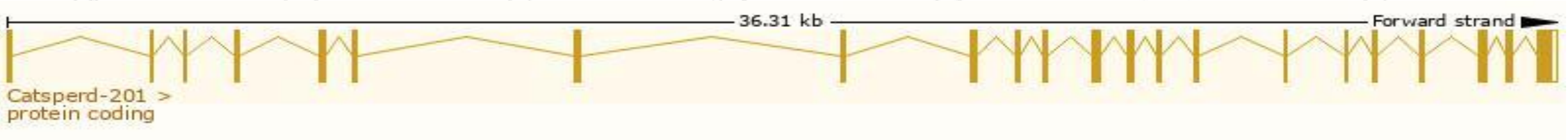
Official Symbol	Catsperd provided by MGI
Official Full Name	cation channel sperm associated auxiliary subunit delta provided by MGI
Primary source	MGI:MGI:2147030
See related	Ensembl:ENSMUSG00000040828
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	4921529N20Rik, 4933402B14Rik, 619718, AW045609, Gm6095, Tmem146
Expression	Biased expression in testis adult (RPKM 33.0), stomach adult (RPKM 5.5) and 1 other tissue See more
Orthologs	human all

Transcript information (Ensembl)

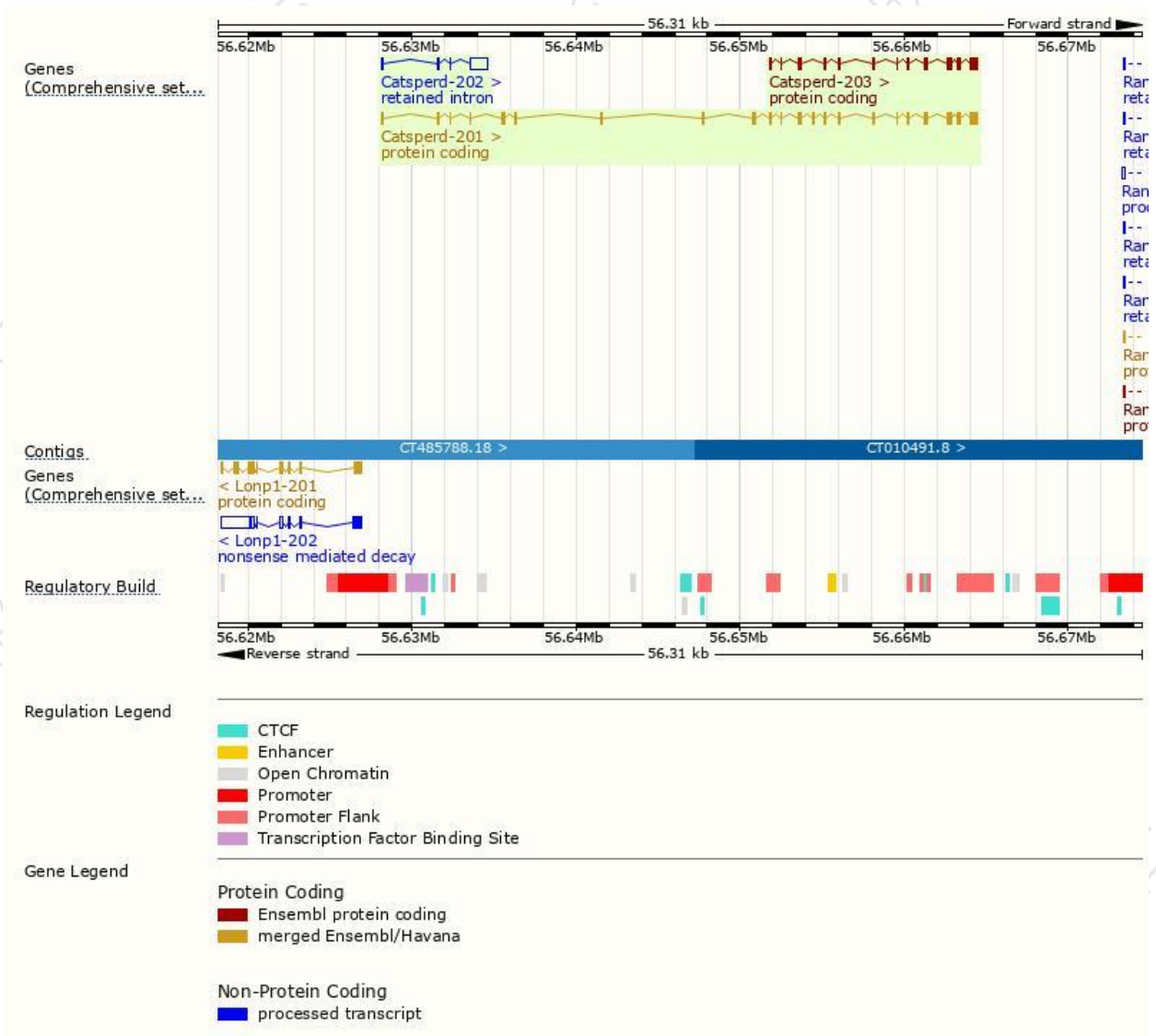
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Catsperd-201	ENSMUST00000112979.3	2557	805aa	Protein coding	CCDS50153	E9Q9F6	TSL:1 GENCODE basic APPRIS P1
Catsperd-203	ENSMUST00000233947.1	1617	505aa	Protein coding	-	A0A3B2WD76	CDS 5' incomplete
Catsperd-202	ENSMUST00000232838.1	1325	No protein	Retained intron	-	-	

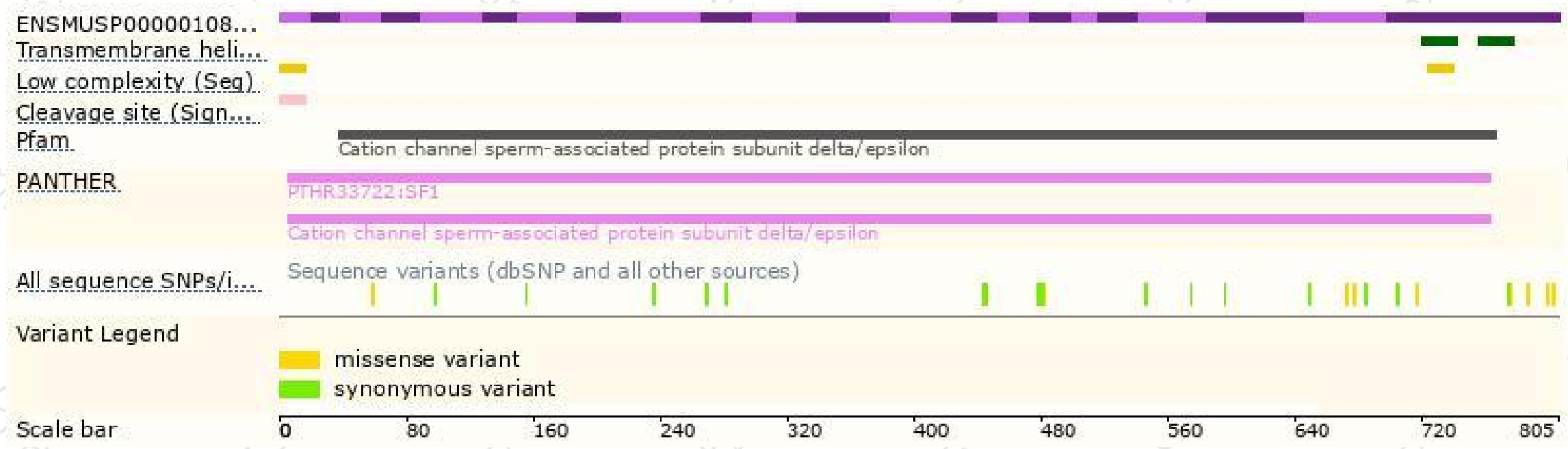
The strategy is based on the design of *Catsperd-201* transcript,the transcription is shown below:



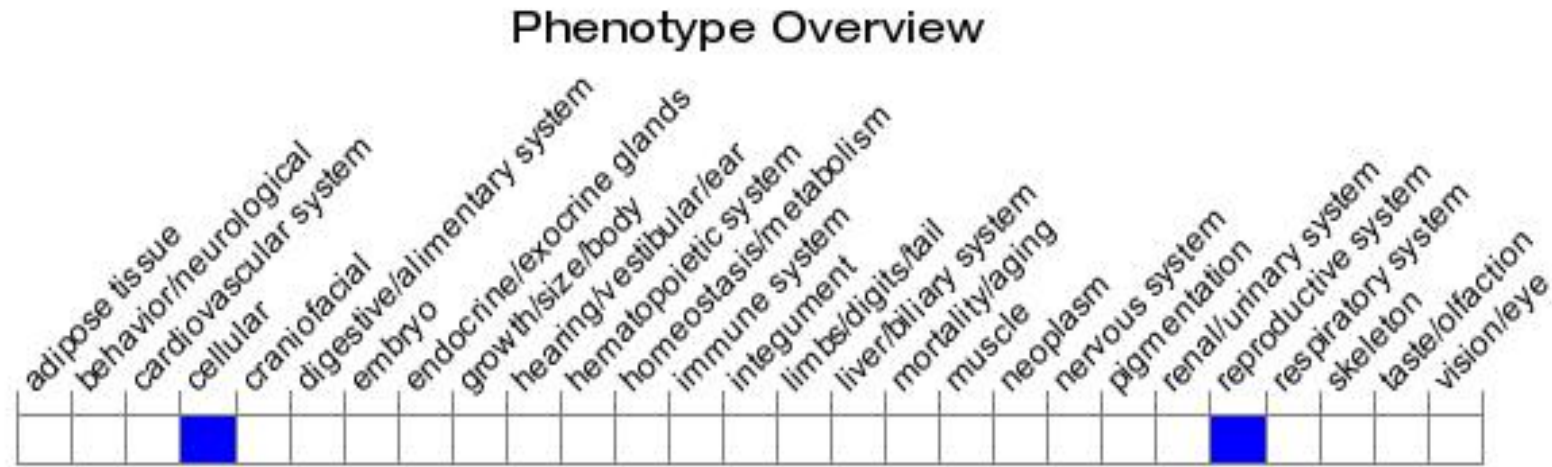
Genomic location distribution



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, mice homozygous for a deletion in this gene display male infertility.

Hyperactivity of sperm fails to develop under capacitating conditions.

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

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