

# *Jmjd1c* Cas9-KO Strategy

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**Reviewer:**

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**Design Date:**

**2020-3-6**

# Project Overview

**Project Name**

***Imjd1c***

**Project type**

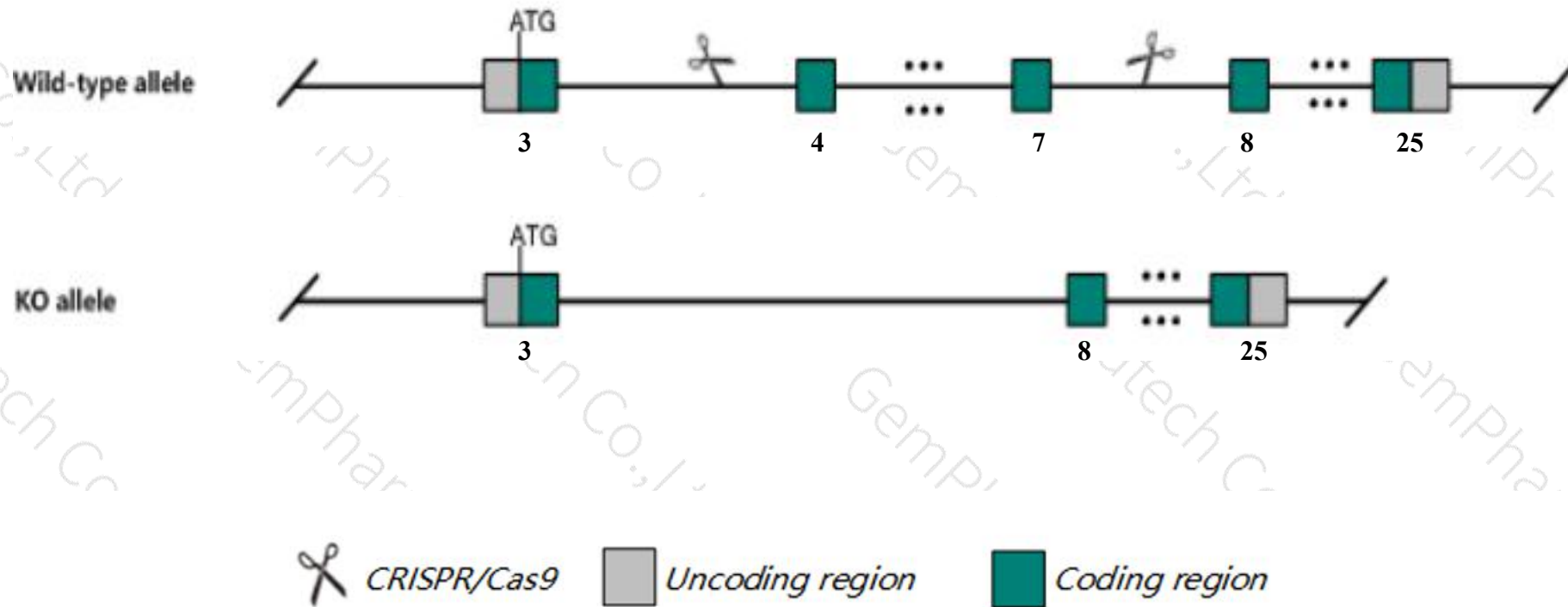
**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Jmjd1c* gene. The schematic diagram is as follows:



- The *Jmjd1c* gene has 12 transcripts. According to the structure of *Jmjd1c* gene, exon4-exon7 of *Jmjd1c*-206 (ENSMUST00000173689.7) transcript is recommended as the knockout region. The region contains 2144bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Jmjd1c* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice homozygous for a null allele exhibit an age-dependent male infertility phenotype, characterized by early loss of undifferentiated spermatogonia, and a progressive reduction in testis size/weight and male germ cells, partly due to increased male germ cell apoptosis.
- The *Jmjd1c* gene is located on the Chr10. 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)

## Jmjd1c jumonji domain containing 1C [Mus musculus (house mouse)]

Gene ID: 108829, updated on 7-Apr-2019

### Summary



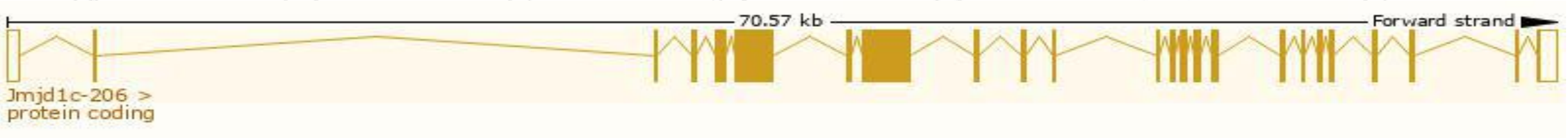
<b>Official Symbol</b>	Jmjd1c provided by <a href="#">MGI</a>
<b>Official Full Name</b>	jumonji domain containing 1C provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:1918614</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000037876</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	VALIDATED
<b>Organism</b>	<a href="#">Mus musculus</a>
<b>Lineage</b>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
<b>Also known as</b>	5430433L24Rik, D630035I23Rik, Jmjd1c, TRIP8
<b>Expression</b>	Ubiquitous expression in cerebellum adult (RPKM 6.3), CNS E14 (RPKM 5.6) and 28 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

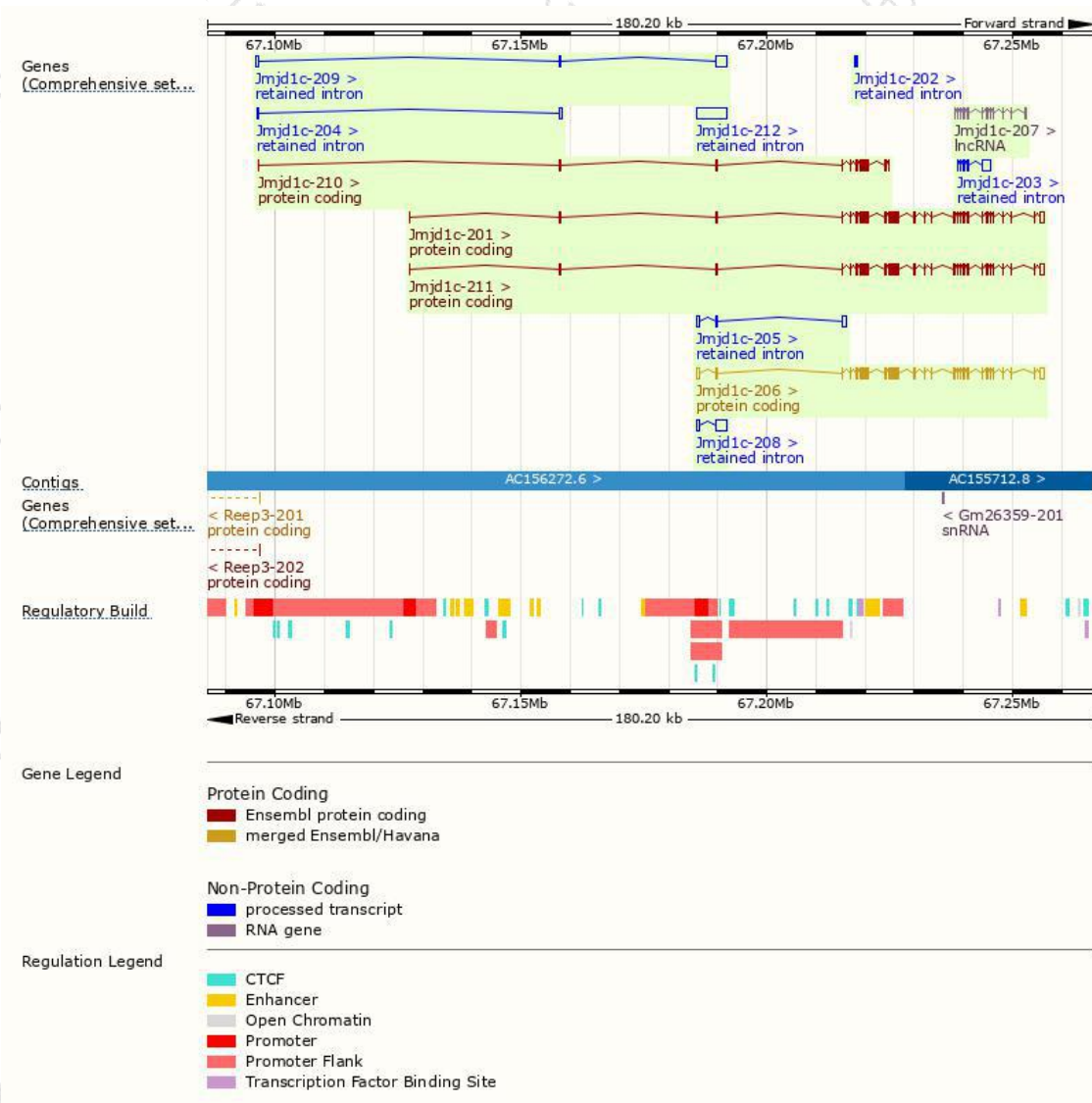
The gene has 12 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Jmjd1c-206	<a href="#">ENSMUST00000173689.7</a>	8609	<a href="#">2350aa</a>	Protein coding	<a href="#">CCDS56708</a>	<a href="#">Q69ZK6</a>	TSL:2 GENCODE basic APPRIS ALT2
Jmjd1c-201	<a href="#">ENSMUST00000051446.15</a>	8382	<a href="#">2530aa</a>	Protein coding	<a href="#">CCDS56707</a>	<a href="#">A0A0A0MQ98</a>	TSL:5 GENCODE basic APPRIS P4
Jmjd1c-211	<a href="#">ENSMUST00000174408.7</a>	8377	<a href="#">2531aa</a>	Protein coding	-	<a href="#">G3UZM1</a>	TSL:5 GENCODE basic APPRIS ALT2
Jmjd1c-210	<a href="#">ENSMUST00000174317.7</a>	3076	<a href="#">744aa</a>	Protein coding	-	<a href="#">G3UYW3</a>	CDS 3' incomplete TSL:1
Jmjd1c-212	<a href="#">ENSMUST00000219563.1</a>	6003	No protein	Retained intron	-	-	TSL:NA
Jmjd1c-209	<a href="#">ENSMUST00000174062.7</a>	2772	No protein	Retained intron	-	-	TSL:1
Jmjd1c-208	<a href="#">ENSMUST00000173949.1</a>	2574	No protein	Retained intron	-	-	TSL:1
Jmjd1c-203	<a href="#">ENSMUST00000173236.1</a>	2365	No protein	Retained intron	-	-	TSL:1
Jmjd1c-205	<a href="#">ENSMUST00000173661.7</a>	1579	No protein	Retained intron	-	-	TSL:1
Jmjd1c-204	<a href="#">ENSMUST00000173295.1</a>	651	No protein	Retained intron	-	-	TSL:2
Jmjd1c-202	<a href="#">ENSMUST00000173187.1</a>	522	No protein	Retained intron	-	-	TSL:3
Jmjd1c-207	<a href="#">ENSMUST00000173762.1</a>	2015	No protein	lncRNA	-	-	TSL:1

The strategy is based on the design of *Jmjd1c-206* transcript,The transcription is shown below

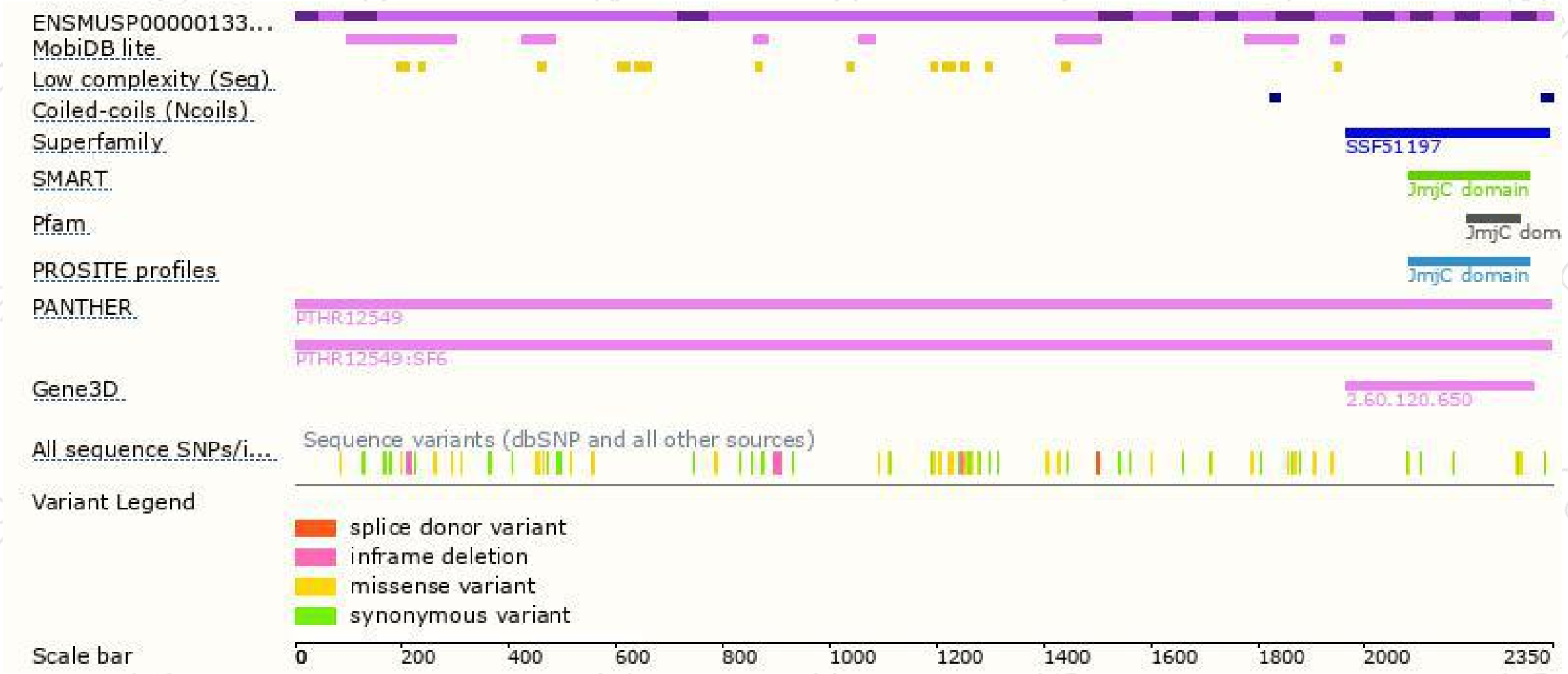


# Genomic location distribution

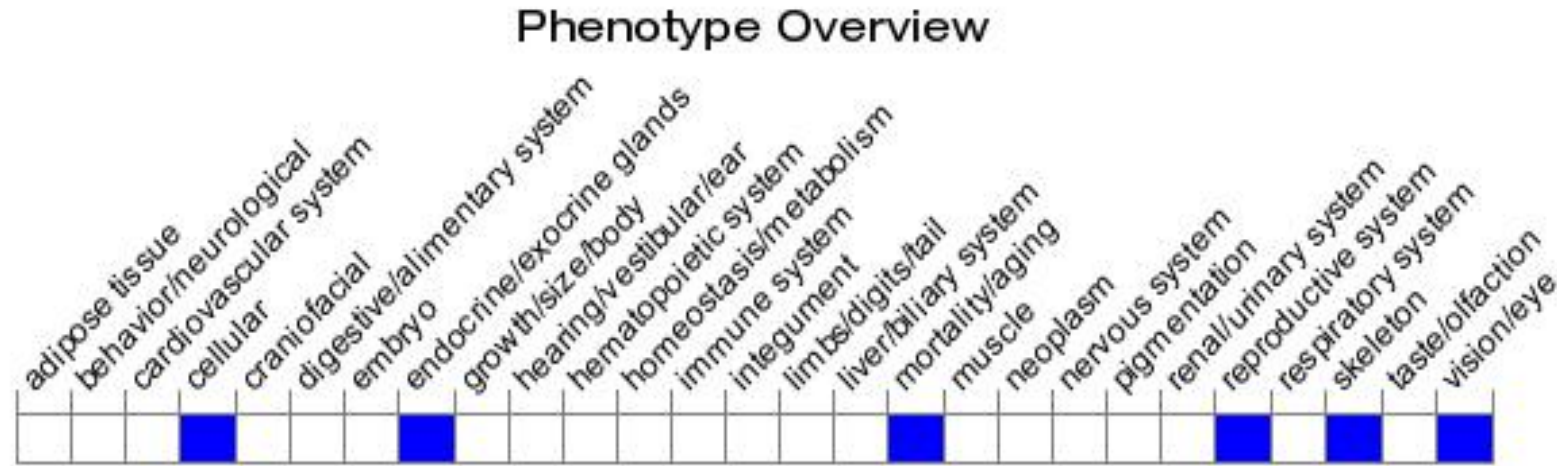




# Protein domain



# 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 null allele exhibit an age-dependent male infertility phenotype, characterized by early loss of undifferentiated spermatogonia, and a progressive reduction in testis size/weight and male germ cells, partly due to increased male germ cell apoptosis.

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

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