

# ***Dnah9*** Cas9-CKO Strategy

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# Project Overview

**Project Name**

***Dnah9***

**Project type**

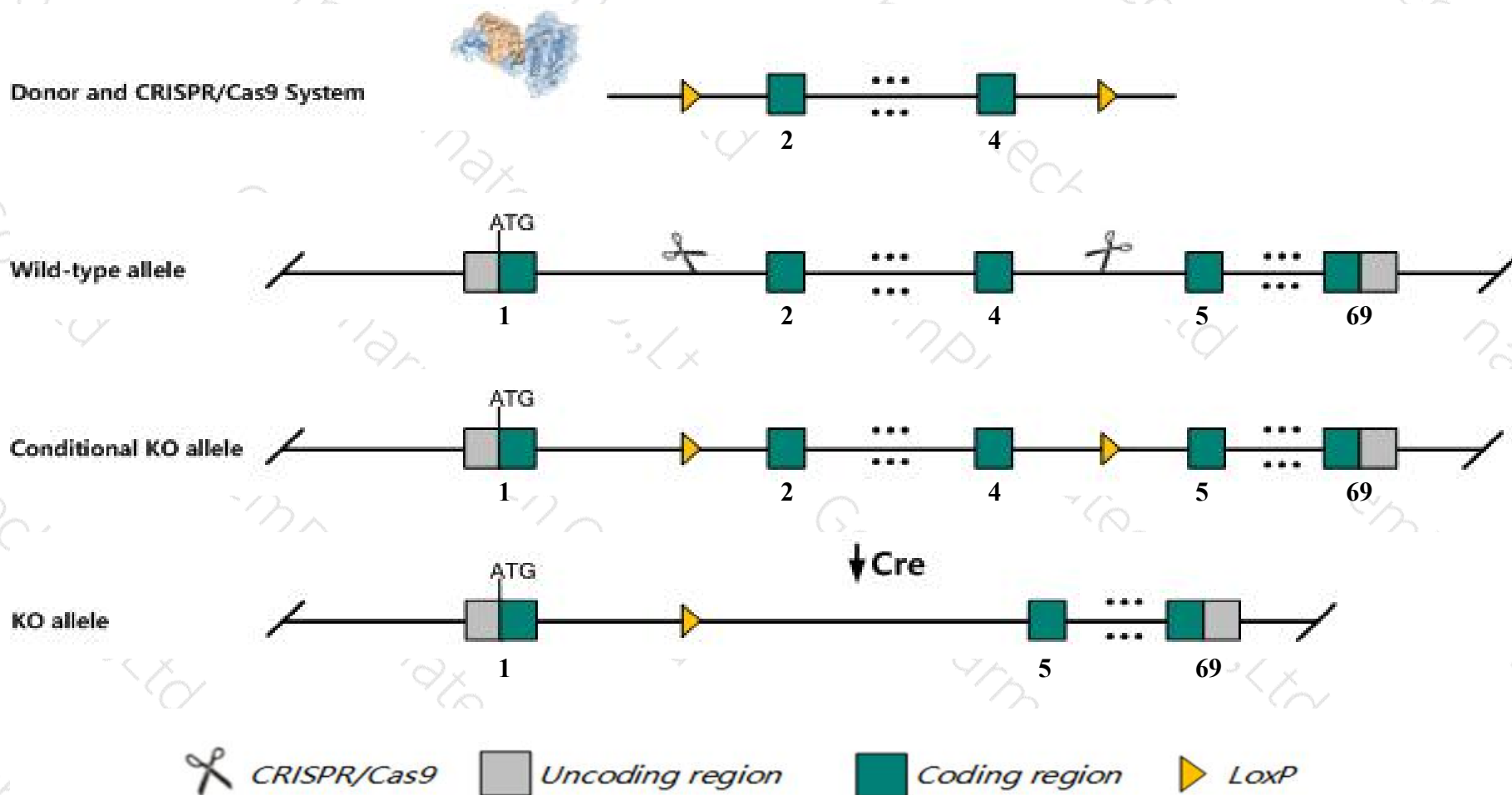
**Cas9-CKO**

**Strain background**

**C57BL/6JGpt**

# Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Dnah9* gene. The schematic diagram is as follows:



- The *Dnah9* gene has 6 transcripts. According to the structure of *Dnah9* gene, exon2-exon4 of *Dnah9*-201 (ENSMUST00000080665.9) transcript is recommended as the knockout region. The region contains 487bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Dnah9* gene. The brief process is as follows: CRISPR/Cas9 system and Donor were microinjected into the fertilized eggs of C57BL/6JGpt mice. Fertilized eggs were transplanted to obtain positive F0 mice which were confirmed by PCR and sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6JGpt mice.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

- Transcript 204 may not be affected. The effect of transcripts 203,206 is unknown.
- The *Dnah9* gene is located on the Chr11. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.



# Gene information (NCBI)

## Dnah9 dynein, axonemal, heavy chain 9 [ *Mus musculus* (house mouse) ]

Gene ID: 237806, updated on 10-Oct-2019

### Summary



Official Symbol	Dnah9 provided by <a href="#">MGI</a>
Official Full Name	dynein, axonemal, heavy chain 9 provided by <a href="#">MGI</a>
Primary source	<a href="#">MGI:MGI:1289279</a>
See related	<a href="#">Ensembl:ENSMUSG00000056752</a>
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	<a href="#">Mus musculus</a>
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Dnahc9; mKIAA0357; 9030022M04; A230091C01; C230051G16; D11Ert686e
Expression	Low expression observed in reference dataset <a href="#">See more</a>
Orthologs	<a href="#">human</a> <a href="#">all</a>

### Genomic context



Location: 11 B3; 11 40.53 cM

Exon count: 69

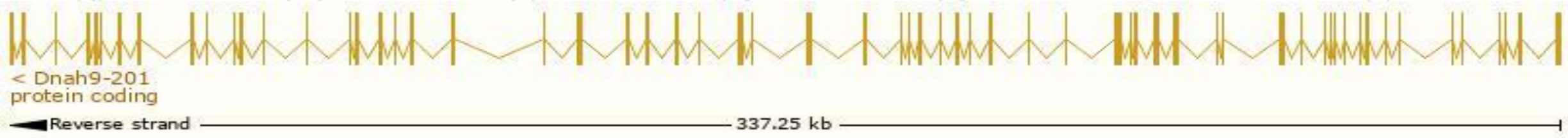
See Dnah9 in [Genome Data Viewer](#)

# Transcript information (Ensembl)

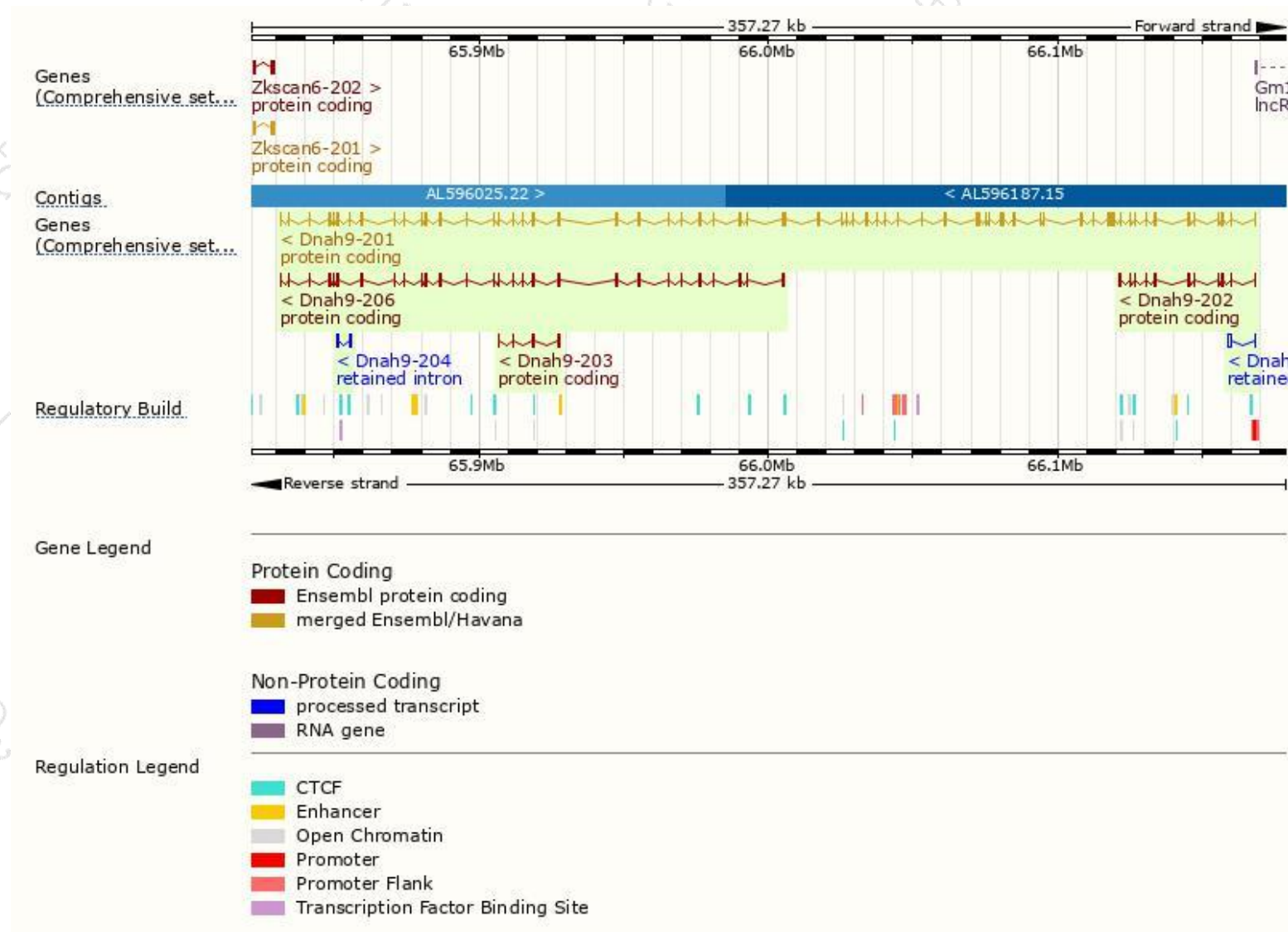
The gene has 6 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Dnah9-201	<a href="#">ENSMUST00000080665.9</a>	13798	<a href="#">4484aa</a>	Protein coding	<a href="#">CCDS36182</a>	<a href="#">B1AR51</a>	TSL:5 GENCODE basic APPRIS P1
Dnah9-206	<a href="#">ENSMUST00000152386.7</a>	6031	<a href="#">1958aa</a>	Protein coding	-	<a href="#">F7DC70</a>	CDS 5' incomplete TSL:5
Dnah9-202	<a href="#">ENSMUST00000108691.1</a>	2671	<a href="#">457aa</a>	Protein coding	-	<a href="#">Q5SX07</a>	TSL:1 GENCODE basic
Dnah9-203	<a href="#">ENSMUST00000130739.1</a>	837	<a href="#">279aa</a>	Protein coding	-	<a href="#">B7FAT5</a>	5' and 3' truncations in transcript evidence prevent annotation of the start and the end of the CDS. CDS 5' and 3' incomplete TSL:5
Dnah9-205	<a href="#">ENSMUST00000142836.1</a>	1797	No protein	Retained intron	-	-	TSL:1
Dnah9-204	<a href="#">ENSMUST00000136124.1</a>	532	No protein	Retained intron	-	-	TSL:5

The strategy is based on the design of *Dnah9-201* transcript,The transcription is shown below

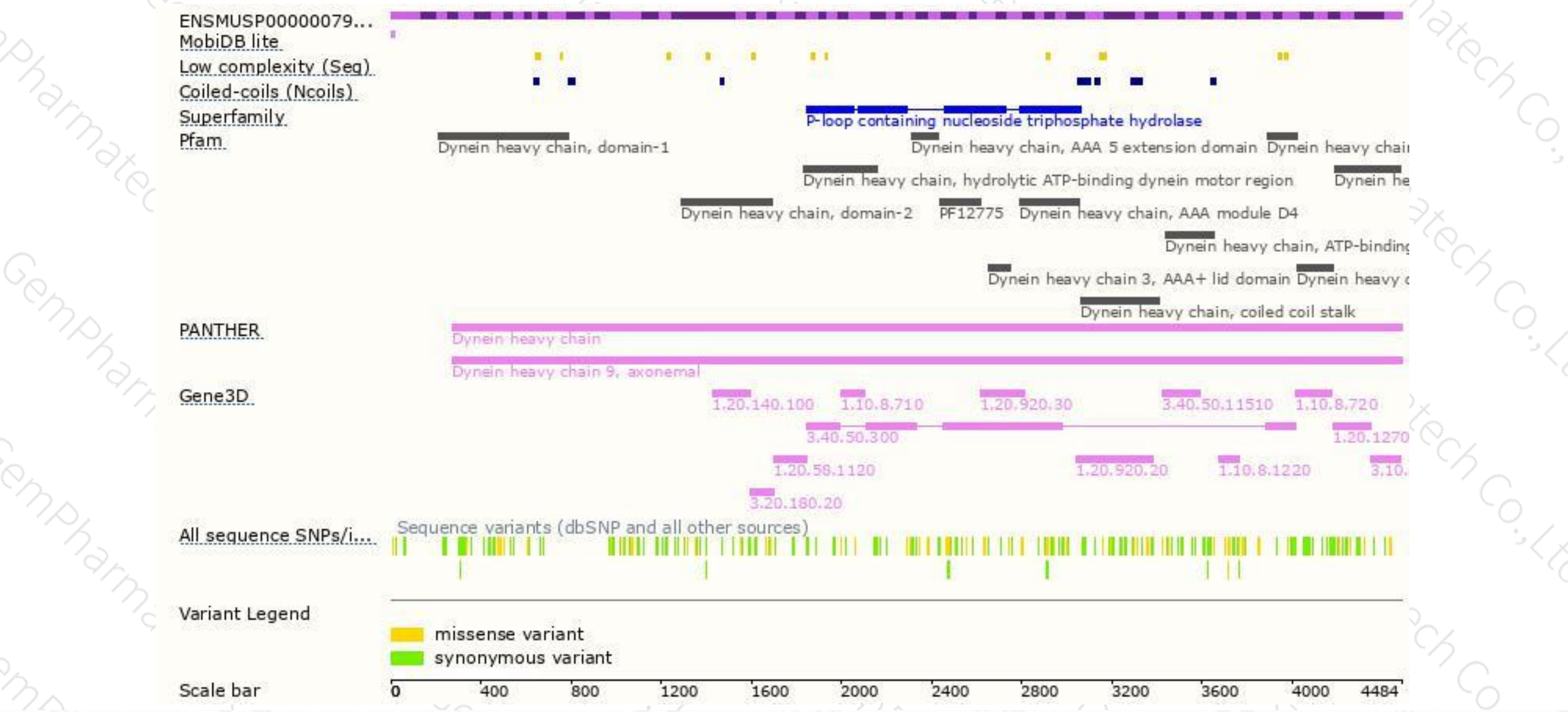


# Genomic location distribution





# Protein domain



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

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