

# ***Krt13* Cas9-CKO Strategy**

**Designer:**

**JiaYu**

**Reviewer:**

**Xiaojing Li**

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# Project Overview

**Project Name**

***Krt13***

**Project type**

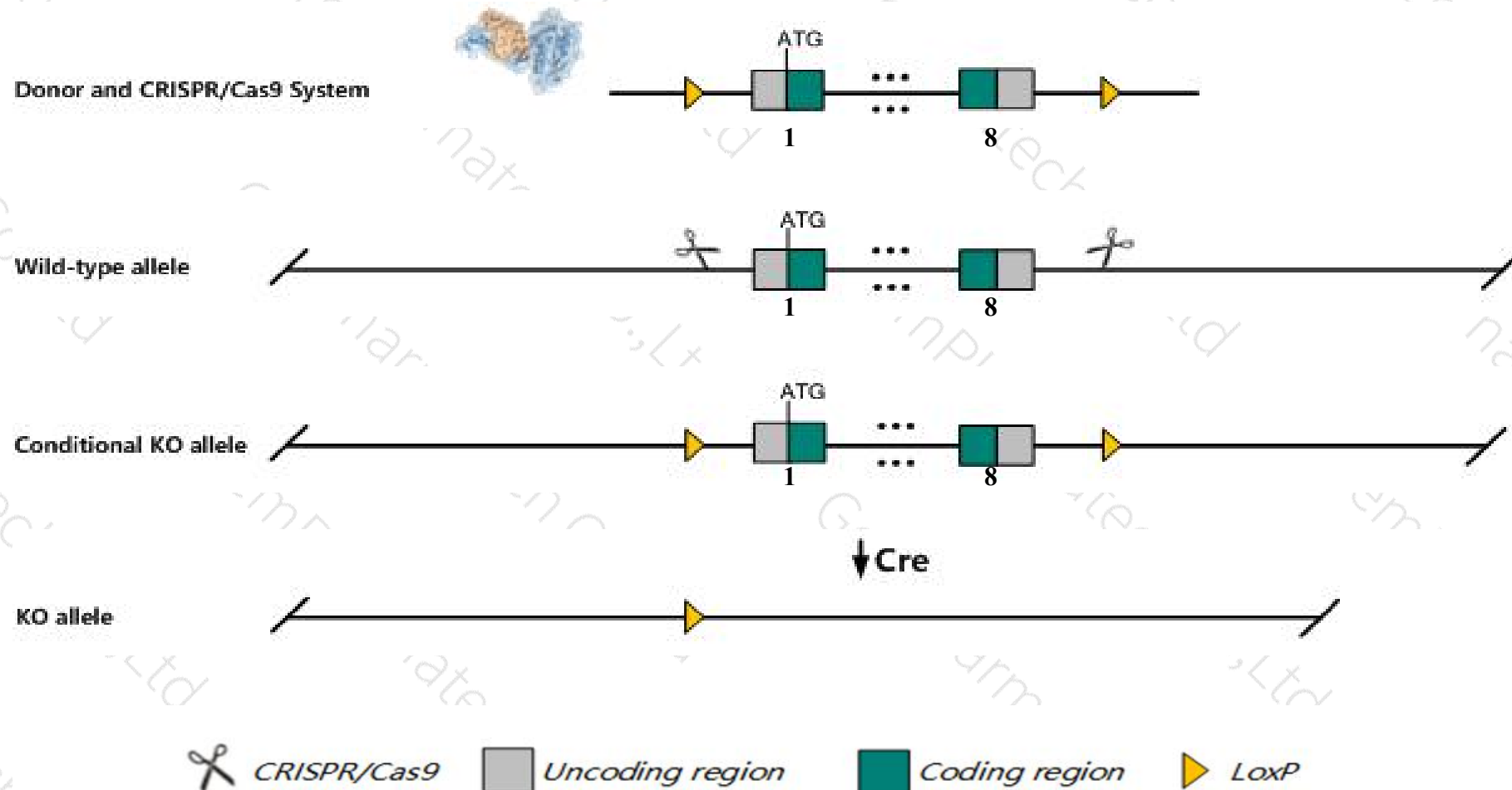
**Cas9-CKO**

**Strain background**

**C57BL/6JGpt**

# Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Krt13* gene. The schematic diagram is as follows:



# Technical routes

- The *Krt13* gene has 3 transcripts. According to the structure of *Krt13* gene, exon1-exon8 of *Krt13-201* (ENSMUST00000007275.2) 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 *Krt13* 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.

# Notice

- The *Krt13* 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)

## Krt13 keratin 13 [Mus musculus (house mouse)]

Gene ID: 16663, updated on 31-Jan-2019

### Summary



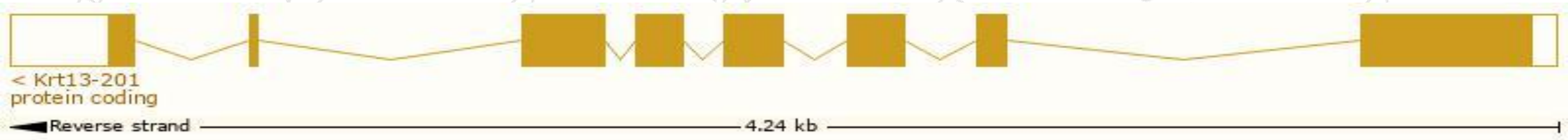
<b>Official Symbol</b>	Krt13 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	keratin 13 provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:101925</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000044041</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	REVIEWED
<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>	CK13, K13, Krt-1.13, Krt1-13
<b>Summary</b>	The protein encoded by this gene is a member of the keratin gene family. The keratins are intermediate filament proteins responsible for the structural integrity of epithelial cells and are subdivided into cytokeratins and hair keratins. Most of the type I cytokeratins consist of acidic proteins which are arranged in pairs of heterotypic keratin chains. This type I cytokeratin is paired with keratin 4 and expressed in the suprabasal layers of non-cornified stratified epithelia. Two transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Sep 2015]
<b>Expression</b>	Biased expression in lung adult (RPKM 380.7) and stomach adult (RPKM 364.4) <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

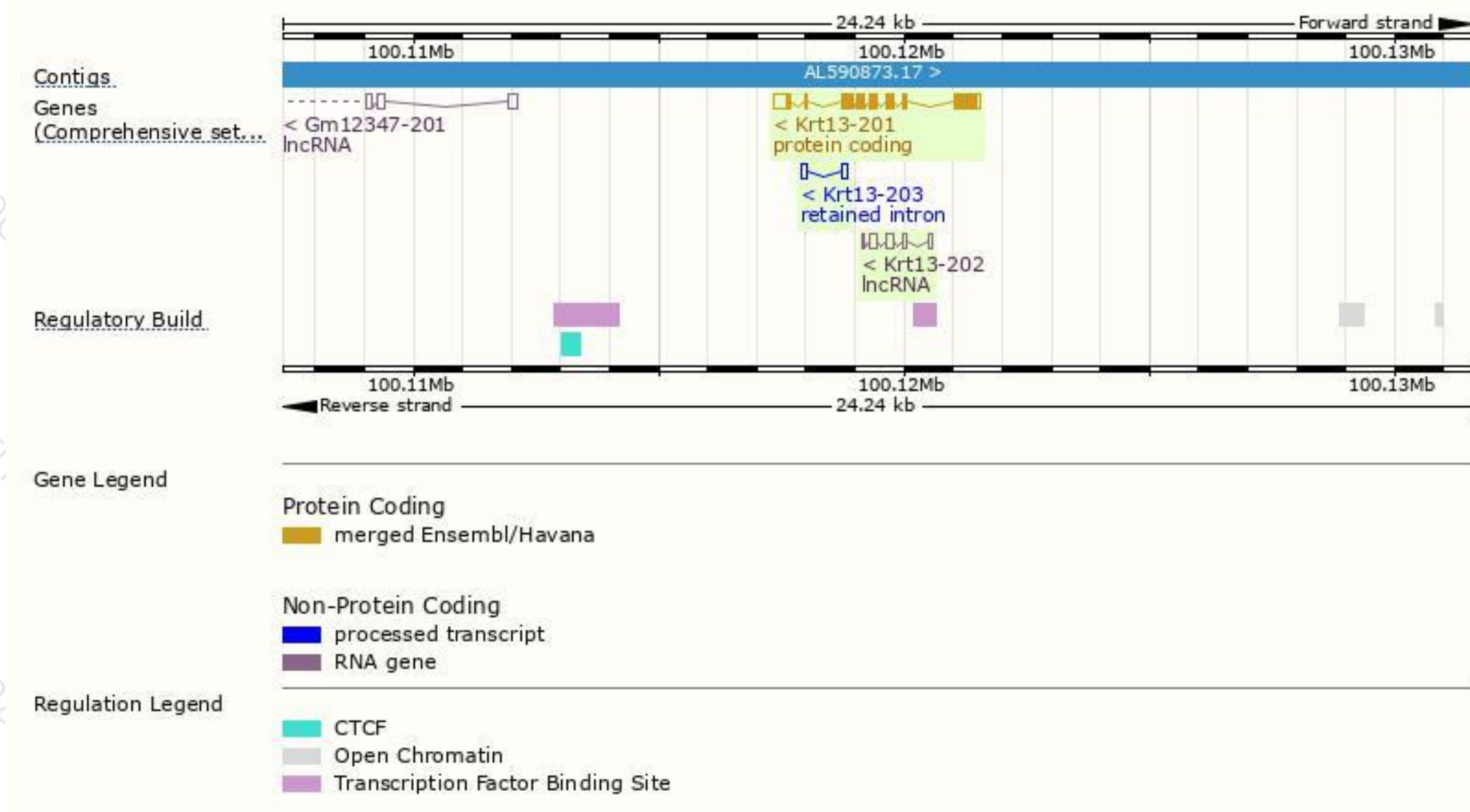
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Krt13-201	<a href="#">ENSMUST00000007275.2</a>	1654	<a href="#">437aa</a>	Protein coding	<a href="#">CCDS25409</a>	<a href="#">P08730</a>	TSL:1 GENCODE basic APPRIS P1
Krt13-203	<a href="#">ENSMUST00000134282.1</a>	207	No protein	Retained intron	-	-	TSL:3
Krt13-202	<a href="#">ENSMUST00000128934.1</a>	512	No protein	lncRNA	-	-	TSL:3

The strategy is based on the design of *Krt13-201* transcript,The transcription is shown below

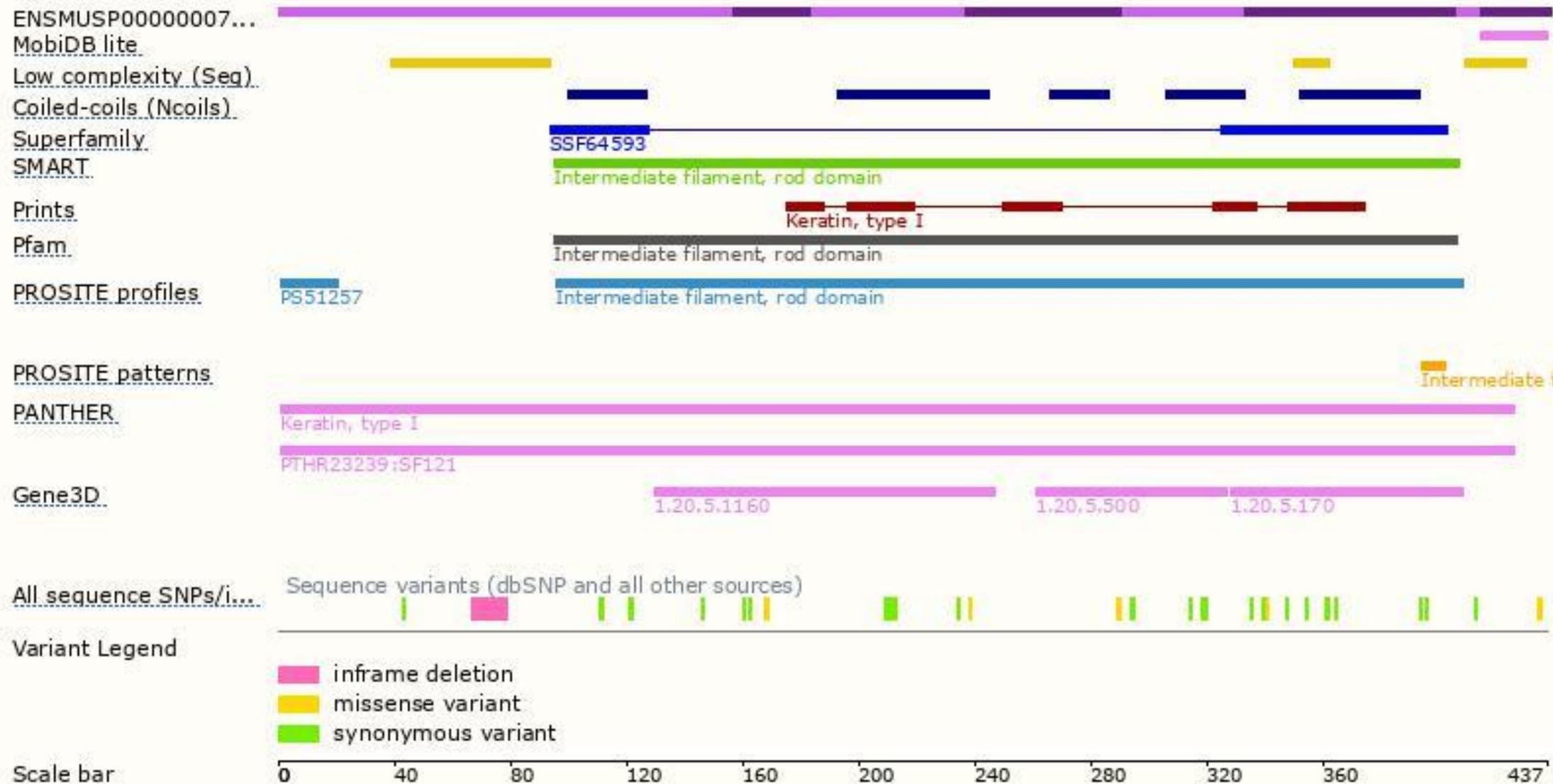


# Genomic location distribution





# Protein domain



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

Tel: 400-9660890

