

# ***Lyzl6*** Cas9-KO Strategy

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# Project Overview

**Project Name**

***Lyz16***

**Project type**

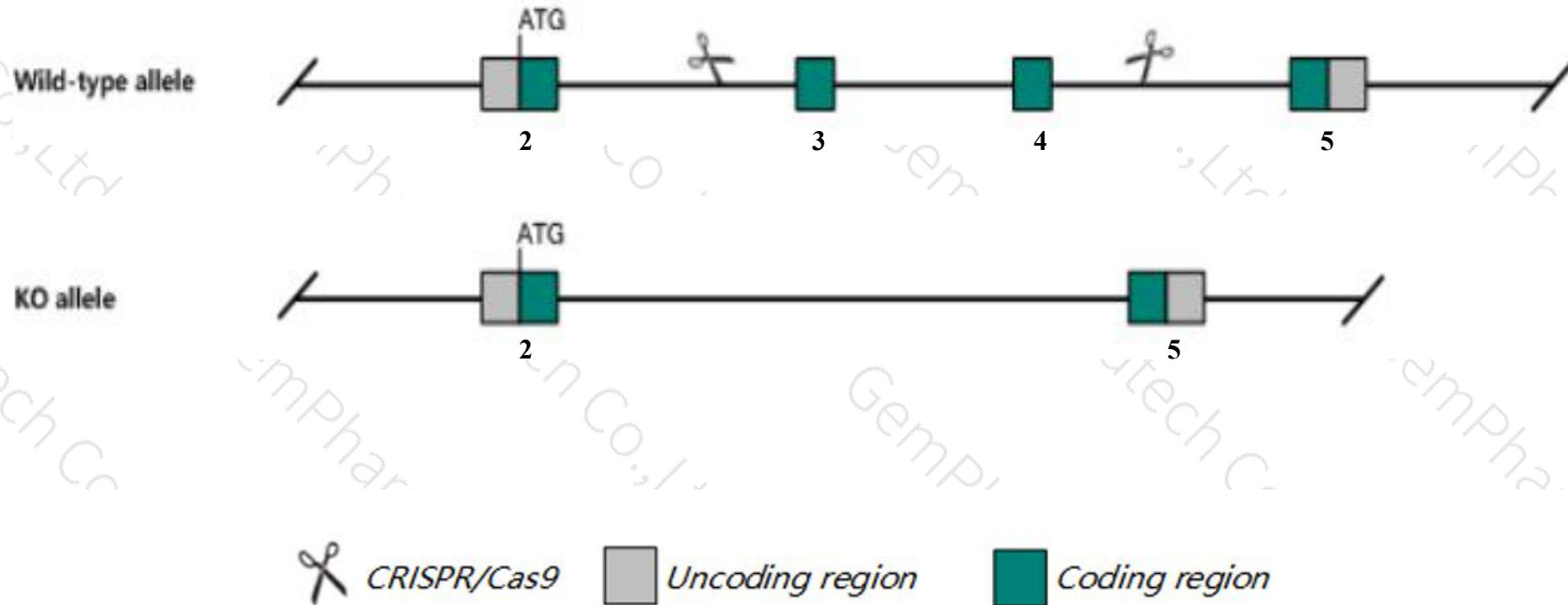
**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Lyzl6* gene. The schematic diagram is as follows:



- The *Lyzl6* gene has 3 transcripts. According to the structure of *Lyzl6* gene, exon3-exon4 of *Lyzl6-201*(ENSMUST00000021328.7) transcript is recommended as the knockout region. The region contains 238bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Lyzl6* gene. The brief process is as follows: CRISPR/Cas9 system 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.

- According to the existing MGI data, male mice homozygous for a mutation are viable and show normal fertility.
- The *Lyz16* 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.



# Gene information (NCBI)

Lyzl6 lysozyme-like 6 [Mus musculus (house mouse)]

Gene ID: 69444, updated on 13-Mar-2020

## Summary



Official Symbol [Lyzl6](#) provided by [MGI](#)

Official Full Name [lysozyme-like 6](#) provided by [MGI](#)

Primary source [MGI:MGI:1916694](#)

See related [Ensembl:ENSMUSG00000020945](#)

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 1700023H08Rik, Lyc1, Lyzb

Expression Restricted expression toward testis adult (RPKM 87.6) [See more](#)

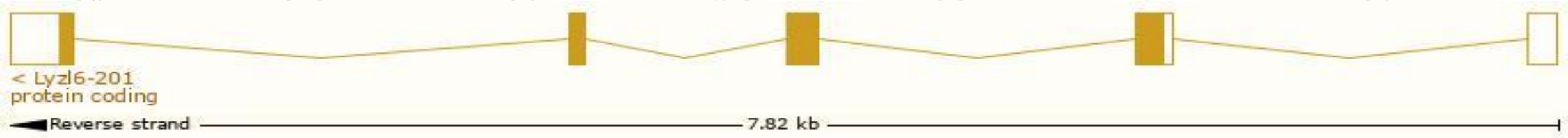
Orthologs [human all](#)

# Transcript information (Ensembl)

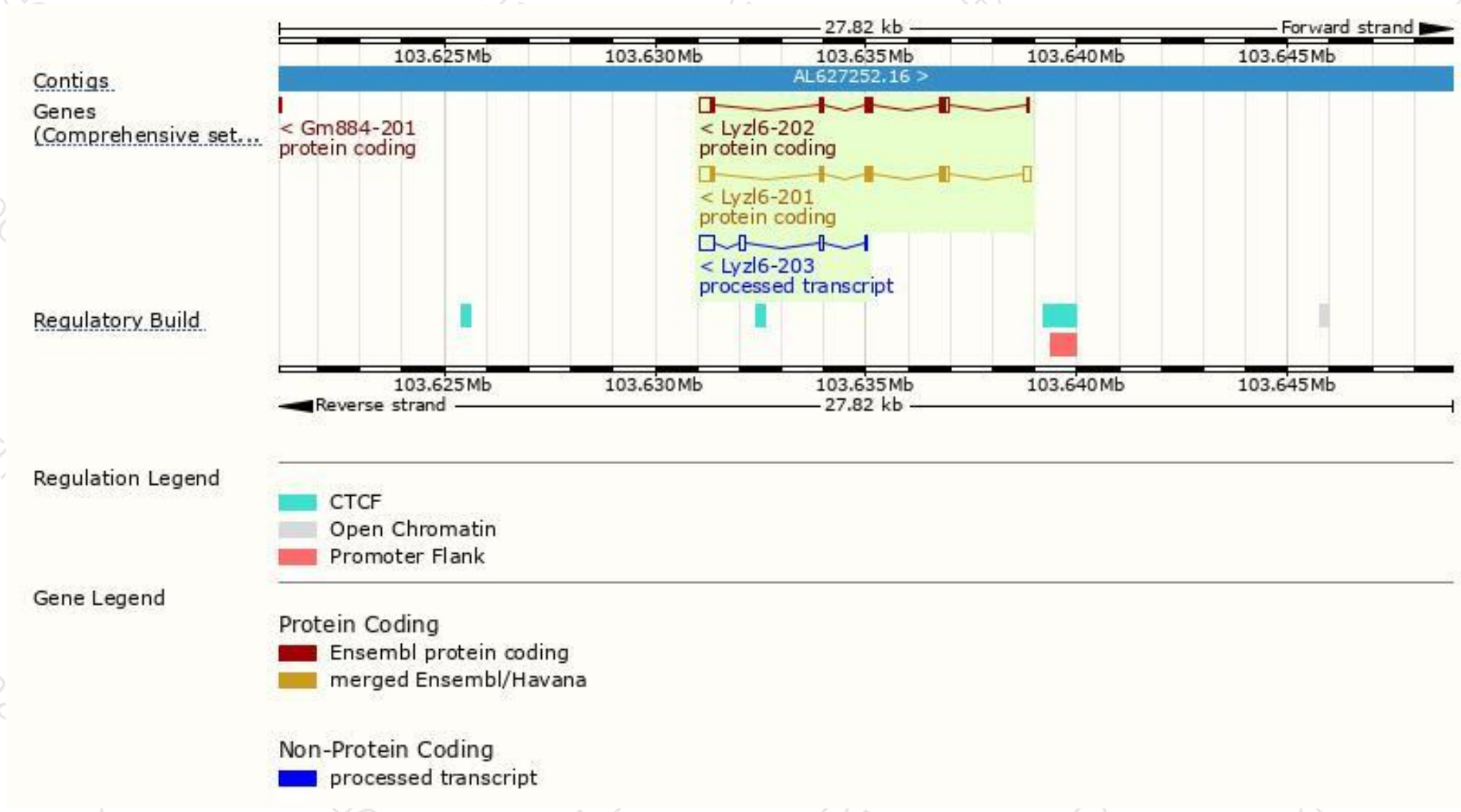
The gene has 3 transcripts,all transcripts are shown below:

| Name      | Transcript ID                        | bp  | Protein               | Biotype              | CCDS                      | UniProt                           | Flags                         |
|-----------|--------------------------------------|-----|-----------------------|----------------------|---------------------------|-----------------------------------|-------------------------------|
| Lyzl6-201 | <a href="#">ENSMUST00000021328.7</a> | 900 | <a href="#">148aa</a> | Protein coding       | <a href="#">CCDS25519</a> | <a href="#">A0A077S2U3 Q9DA11</a> | TSL:1 GENCODE basic APPRIS P1 |
| Lyzl6-202 | <a href="#">ENSMUST00000107014.7</a> | 802 | <a href="#">148aa</a> | Protein coding       | <a href="#">CCDS25519</a> | <a href="#">A0A077S2U3 Q9DA11</a> | TSL:1 GENCODE basic APPRIS P1 |
| Lyzl6-203 | <a href="#">ENSMUST00000140430.1</a> | 512 | No protein            | Processed transcript | -                         | -                                 | TSL:5                         |

The strategy is based on the design of *Lyzl6-201* transcript,the transcription is shown below:

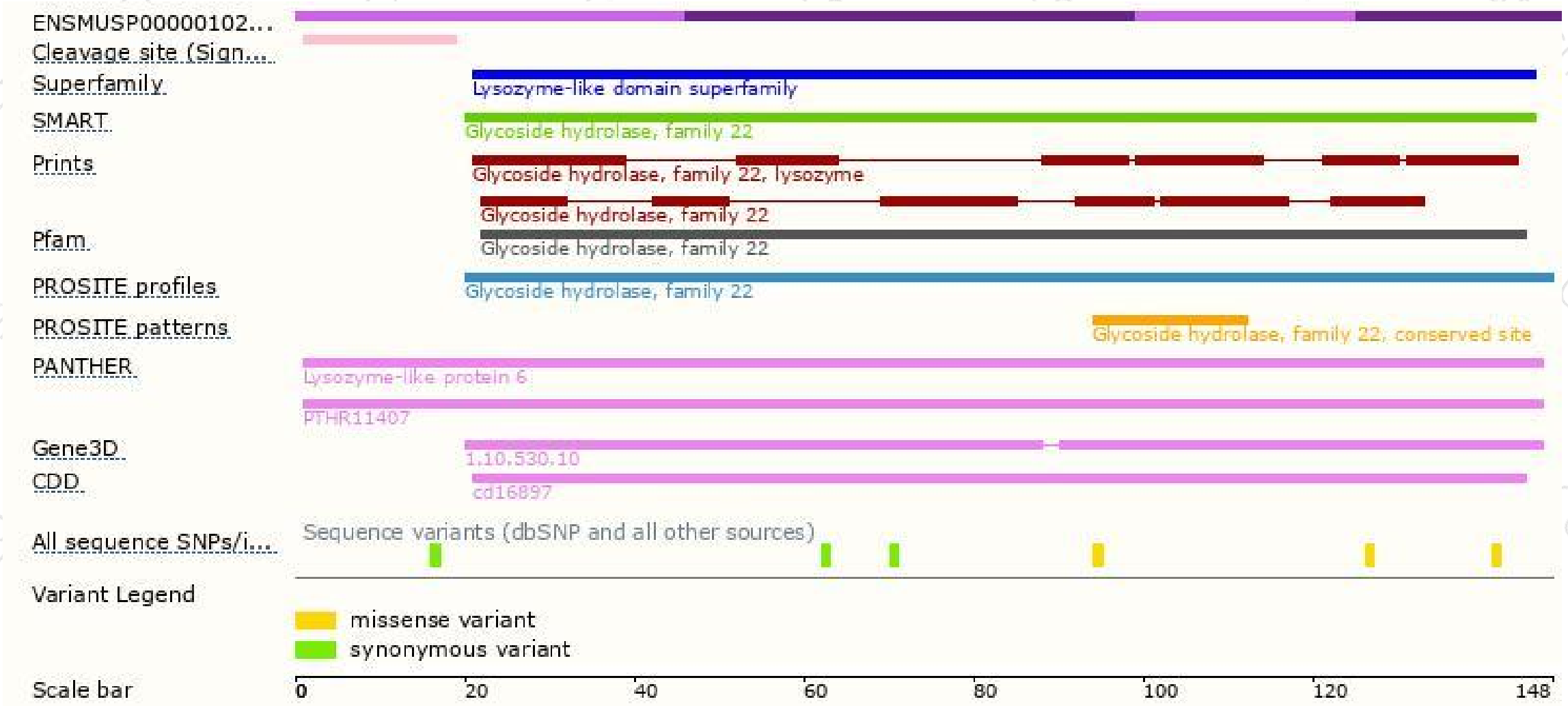


# Genomic location distribution





# Protein domain



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

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