

# *Bcl3* Cas9-KO Strategy

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**Design Date:** 2019-8-3

# Project Overview

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**Project Name**

*Bcl3*

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**Project type**

**Cas9-KO**

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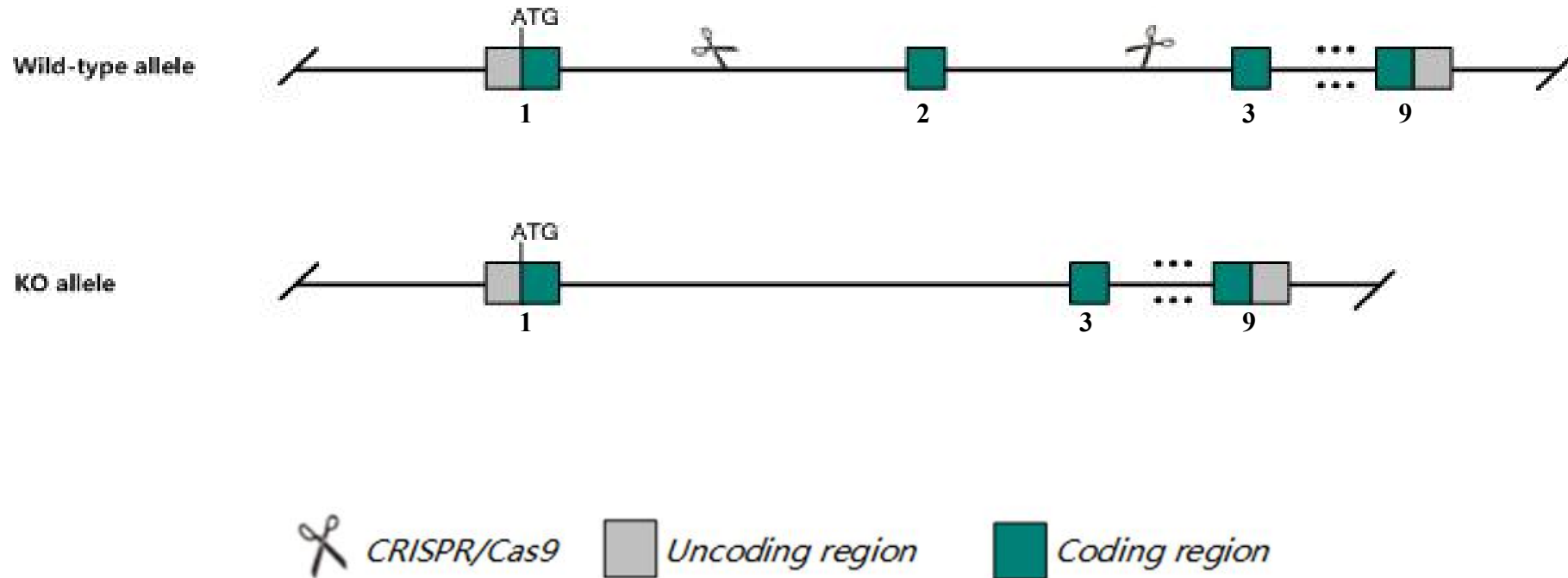
**Strain background**

**C57BL/6JGpt**

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# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Bcl3* gene. The schematic diagram is as follows:



The *Bcl3* gene has 6 transcripts. According to the structure of *Bcl3* gene, exon2 of *Bcl3-201* (ENSMUST00000120537.7) transcript is recommended as the knockout region. The region contains 154bp coding sequence. Knock out the region will result in disruption of protein function.

In this project we use CRISPR/Cas9 technology to modify *Bcl3* gene. The brief process is as follows: CRISPR/Cas9 system w

According to the existing MGI data, Mice lacking functional copies of this gene exhibit defects of the immune system including disruption of the humoral immune response and abnormal spleen and Peyer's patch organogenesis. Mutant mice show increased susceptibility to pathogens.

The *Bcl3* gene is located on the Chr7. 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.

## Bcl3 B cell leukemia/lymphoma 3 [Mus musculus (house mouse)]

Gene ID: 12051, updated on 25-Mar-2019

### Summary

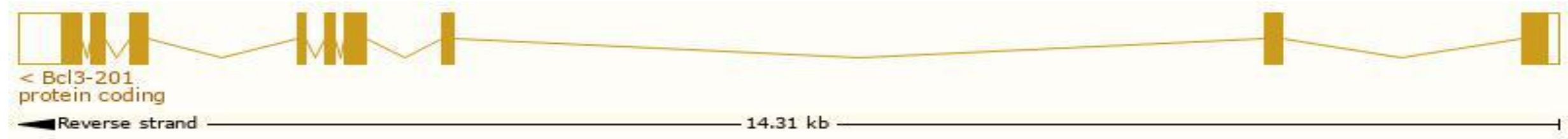
|                           |                                                                                                                                                                           |
|---------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <b>Official Symbol</b>    | Bcl3 provided by <a href="#">MGI</a>                                                                                                                                      |
| <b>Official Full Name</b> | B cell leukemia/lymphoma 3 provided by <a href="#">MGI</a>                                                                                                                |
| <b>Primary source</b>     | <a href="#">MGI:MGI:88140</a>                                                                                                                                             |
| <b>See related</b>        | <a href="#">Ensembl:ENSMUSG00000053175</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>      | AI528691, Bcl-3                                                                                                                                                           |
| <b>Expression</b>         | Biased expression in duodenum adult (RPKM 114.5), small intestine adult (RPKM 82.2) and 11 other tissues <a href="#">See more</a>                                         |
| <b>Orthologs</b>          | <a href="#">human</a> <a href="#">all</a>                                                                                                                                 |

# Transcript information      Ensembl

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

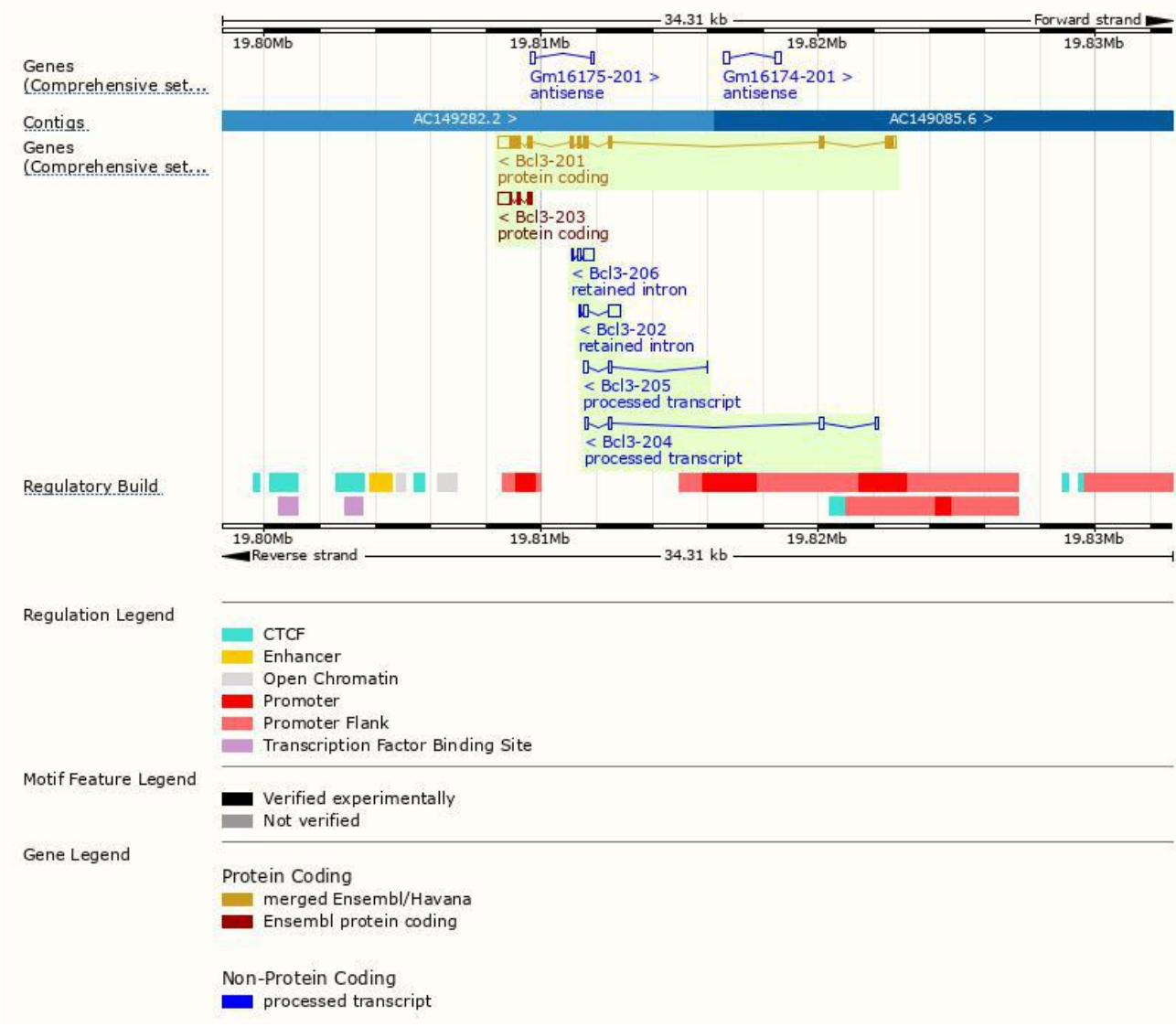
| Name     | Transcript ID                        | bp   | Protein               | Biotype         | CCDS                      | UniProt                | Flags                         |
|----------|--------------------------------------|------|-----------------------|-----------------|---------------------------|------------------------|-------------------------------|
| Bcl3-206 | <a href="#">ENSMUST00000152768.1</a> | 502  | No protein            | Retained intron | -                         | -                      | TSL:3                         |
| Bcl3-205 | <a href="#">ENSMUST00000141996.1</a> | 346  | No protein            | lncRNA          | -                         | -                      | TSL:3                         |
| Bcl3-204 | <a href="#">ENSMUST00000139680.1</a> | 460  | No protein            | lncRNA          | -                         | -                      | TSL:3                         |
| Bcl3-203 | <a href="#">ENSMUST00000135609.7</a> | 755  | <a href="#">114aa</a> | Protein coding  | -                         | <a href="#">F6YNH8</a> | CDS 5' incomplete TSL:3       |
| Bcl3-202 | <a href="#">ENSMUST00000123375.7</a> | 694  | No protein            | Retained intron | -                         | -                      | TSL:3                         |
| Bcl3-201 | <a href="#">ENSMUST00000120537.7</a> | 1850 | <a href="#">448aa</a> | Protein coding  | <a href="#">CCDS20914</a> | <a href="#">Q9Z2F6</a> | TSL:1 GENCODE basic APPRIS P1 |

The strategy is based on the design of *Bcl3-201* transcript,The transcription is shown below





# Genomic location distribution

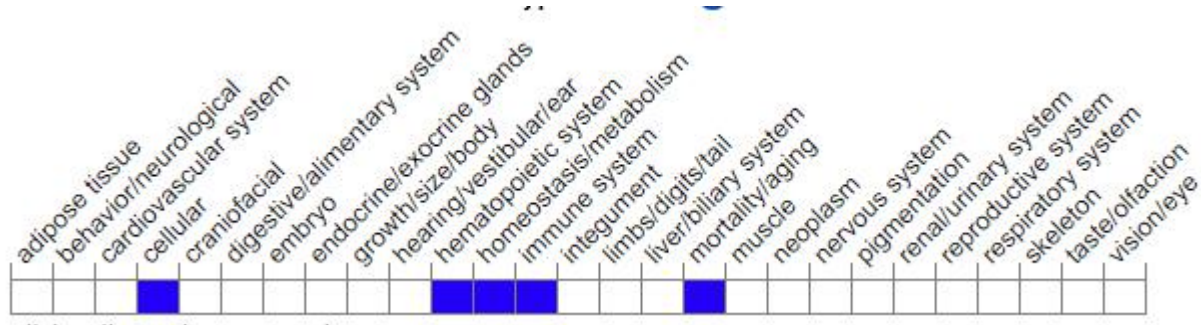




# Protein domain



# 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 lacking functional copies of this gene exhibit defects of the immune system including disruption of the humoral immune response and abnormal spleen and Peyer's patch organogenesis. Mutant mice show increased susceptibility to pathogens.

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