

# *Clic6* Cas9-KO Strategy

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**Reviewer:**

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# Project Overview

**Project Name**

*Clic6*

**Project type**

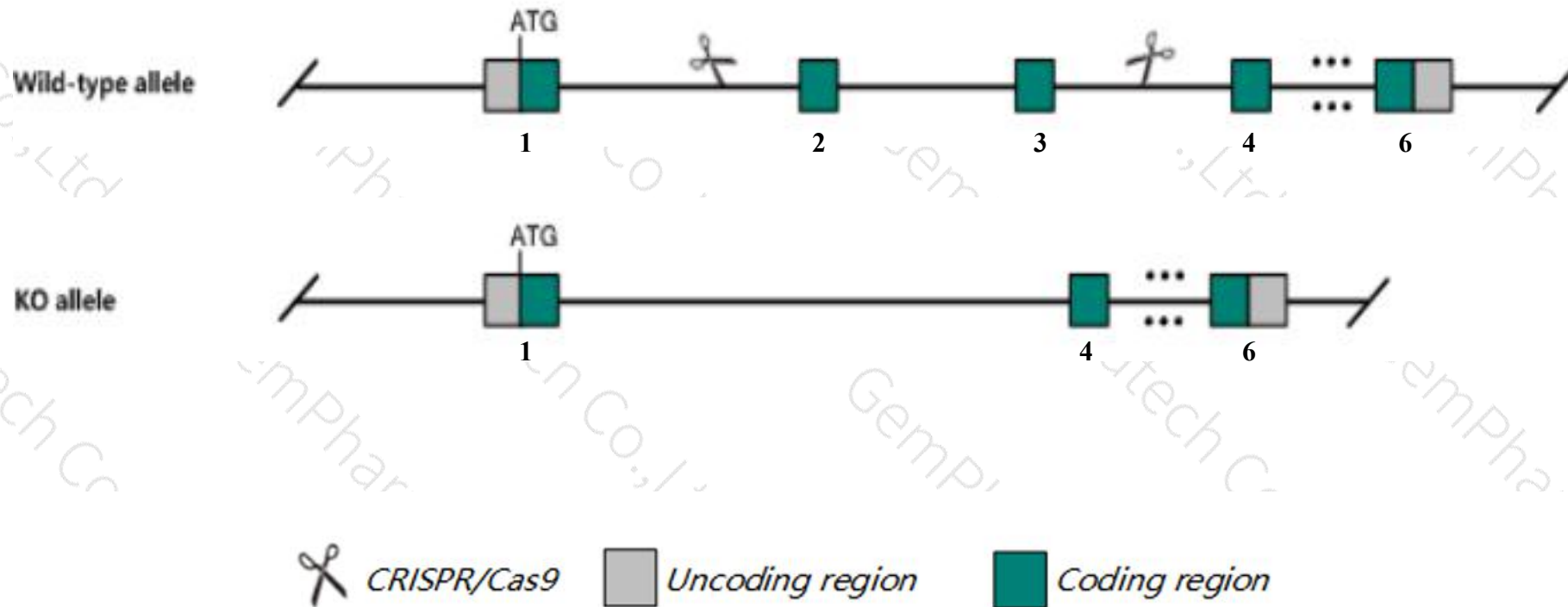
**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Clic6* gene. The schematic diagram is as follows:



- The *Clic6* gene has 2 transcripts. According to the structure of *Clic6* gene, exon2-exon3 of *Clic6-201* (ENSMUST00000023670.3) transcript is recommended as the knockout region. The region contains 236bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Clic6* gene. The brief process is as follows: CRISPR/Cas9 system

- Some amino acids will remain at the N-terminus and some functions may be retained.
- The effect of transcript 202 is unknown.
- The *Clic6* gene is located on the Chr16. 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)

## Clc6 chloride intracellular channel 6 [Mus musculus (house mouse)]

Gene ID: 209195, updated on 13-Mar-2020

### Summary



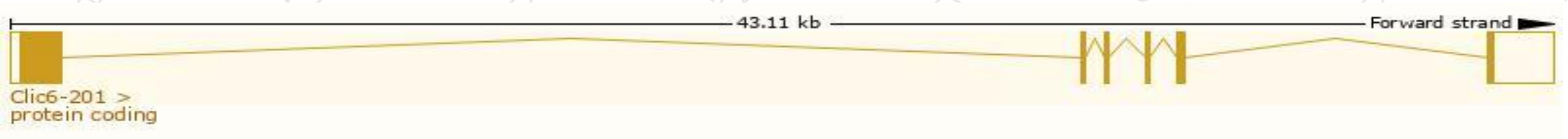
<b>Official Symbol</b>	Clc6 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	chloride intracellular channel 6 provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:2146607</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000022949</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	PROVISIONAL
<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>	5730466J16Rik, AL022908, AW045520, CLIC1L
<b>Expression</b>	Biased expression in stomach adult (RPKM 15.5), placenta adult (RPKM 13.6) and 14 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

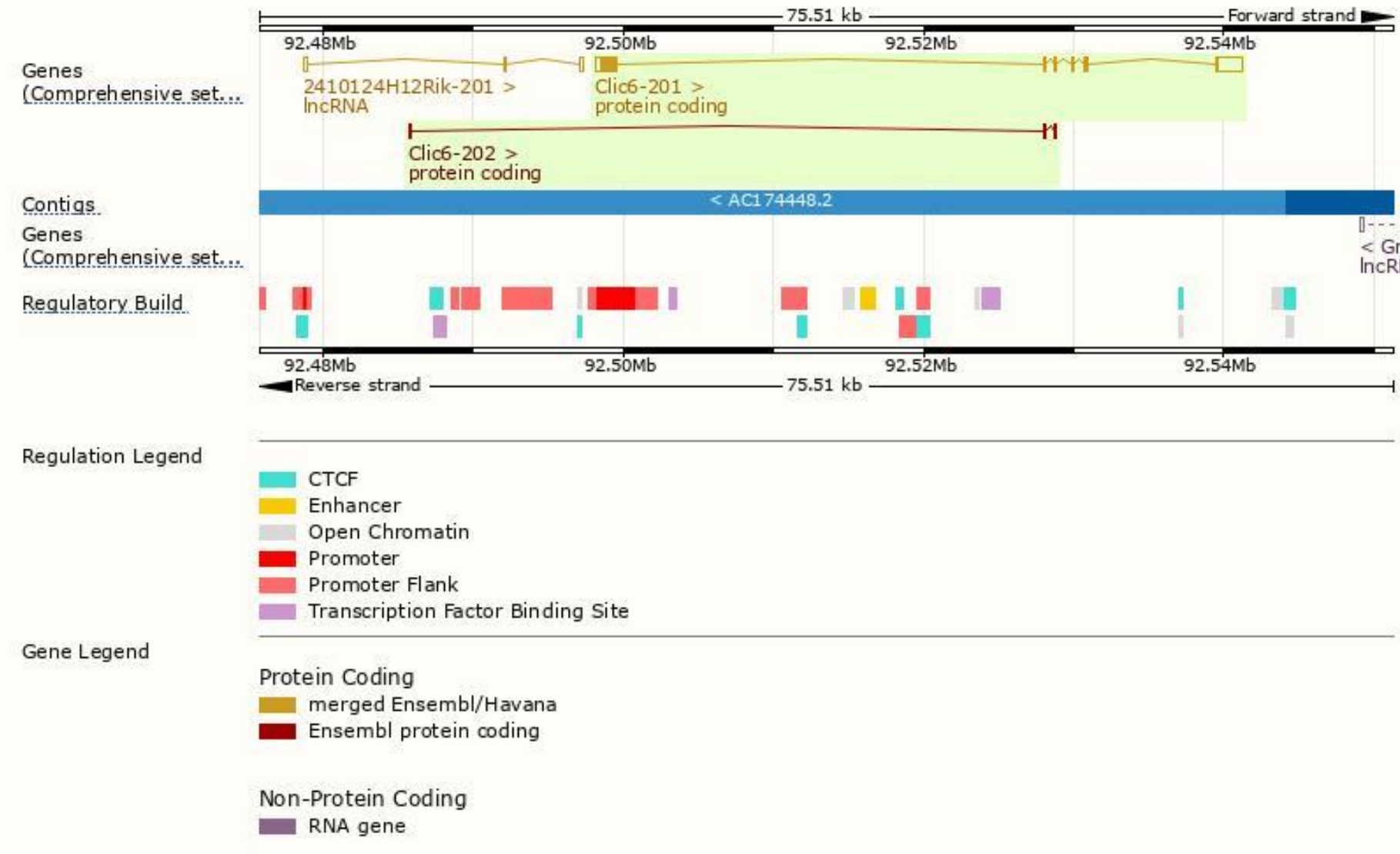
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Clic6-201	<a href="#">ENSMUST00000023670.3</a>	3758	<a href="#">596aa</a>	Protein coding	<a href="#">CCDS28338</a>	<a href="#">Q8BHB9</a>	TSL:1 GENCODE basic APPRIS is a system to annotate alternatively spliced transcripts based on a range of computational methods to identify the most functionally important transcript(s) of a gene. APPRIS P1
Clic6-202	<a href="#">ENSMUST00000162181.7</a>	376	<a href="#">100aa</a>	Protein coding	-	<a href="#">E0CY42</a>	CDS 3' incomplete TSL:2

The strategy is based on the design of *Clic6-201* 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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