

# ***Ints1* Cas9-CKO Strategy**

**Designer: Xueting Zhang**

**Reviewer: Daohua Xu**

**Design Date: 2020-7-28**

# Project Overview

---

**Project Name**

***Ints1***

---

**Project type**

**Cas9-CKO**

---

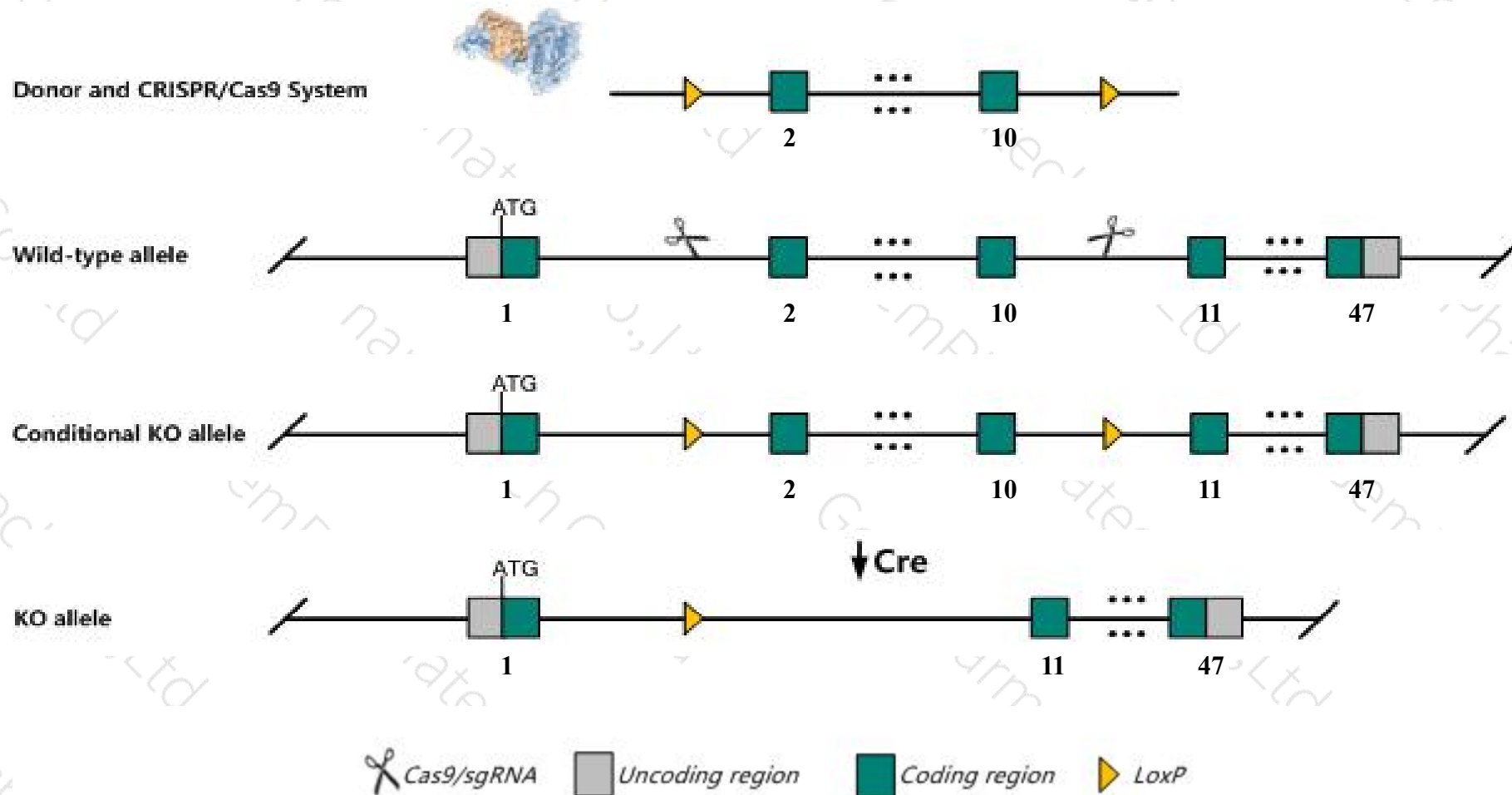
**Strain background**

**C57BL/6JGpt**

---

# Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Ints1* gene. The schematic diagram is as follows:



- The *Ints1* gene has 10 transcripts. According to the structure of *Ints1* gene, exon2-exon10 of *Ints1-210*(ENSMUST00000200393.4) transcript is recommended as the knockout region. The region contains 1550bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Ints1* gene. The brief process is as follows: sgRNA was transcribed in vitro, donor vector was constructed. Cas9, sgRNA 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 was 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.

- According to the existing MGI data, homozygous mutation of this gene results in embryonic lethality at the blastocyst stage.
- Transcript *Ints1*-202&207&208&209 may not be affected.
- The effect on transcript *Ints1*-206 is unknown.
- The floxed region is near to the N-terminal of *Mir7037* gene, this strategy may influence the regulatory function of the N-terminal of *Mir7037* gene.
- The *Ints1* gene is located on the Chr5. 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)

## Ints1 integrator complex subunit 1 [Mus musculus (house mouse)]

Gene ID: 68510, updated on 13-Mar-2020

### Summary



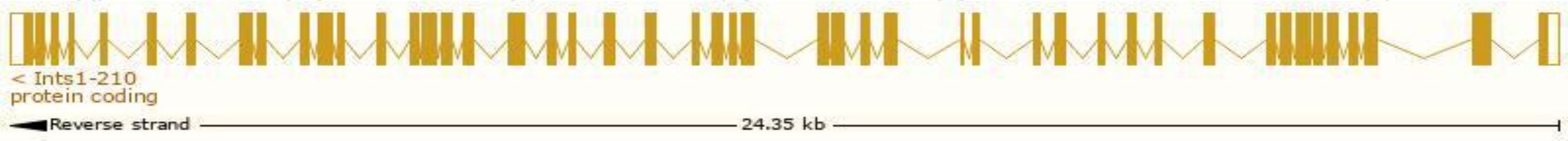
<b>Official Symbol</b>	Ints1 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	integrator complex subunit 1 provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:1915760</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000029547</a>
<b>Gene type</b>	protein coding
<b>RefSeq 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>	1110015K06Rik
<b>Expression</b>	Ubiquitous expression in thymus adult (RPKM 35.7), adrenal adult (RPKM 20.2) and 28 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

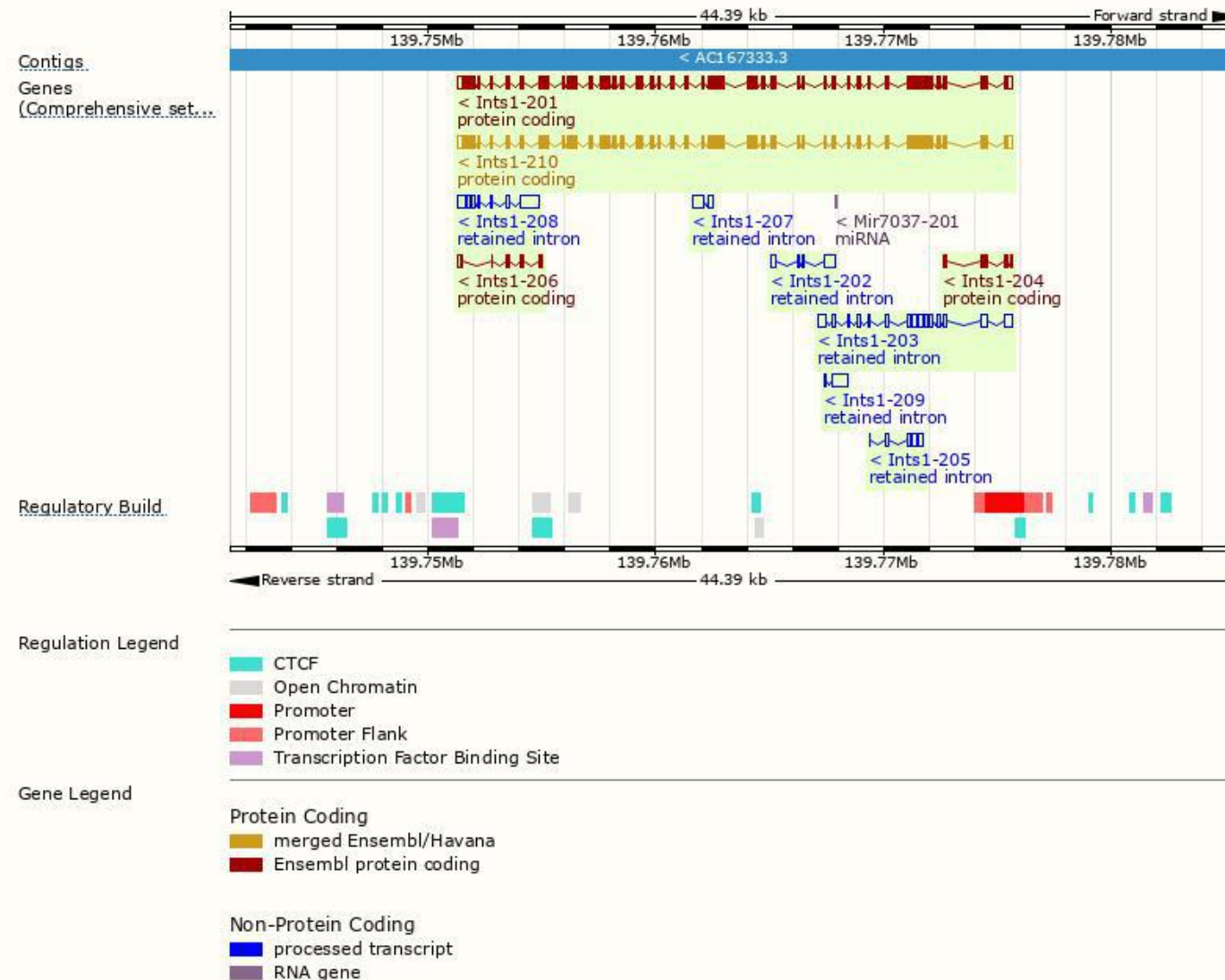
The gene has 10 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ints1-210	<a href="#">ENSMUST00000200393.4</a>	7070	<a href="#">2222aa</a>	Protein coding	<a href="#">CCDS57399</a>	<a href="#">A0A0G2JH17</a>	TSL:1 GENCODE basic APPRIS P2
Ints1-201	<a href="#">ENSMUST00000072607.8</a>	7102	<a href="#">2220aa</a>	Protein coding	-	<a href="#">K3W4P2</a>	TSL:5 GENCODE basic APPRIS ALT2
Ints1-204	<a href="#">ENSMUST00000196864.1</a>	621	<a href="#">175aa</a>	Protein coding	-	<a href="#">A0A0G2JDG8</a>	CDS 3' incomplete TSL:3
Ints1-206	<a href="#">ENSMUST00000197187.1</a>	563	<a href="#">137aa</a>	Protein coding	-	<a href="#">A0A0G2JFR3</a>	CDS 5' incomplete TSL:5
Ints1-203	<a href="#">ENSMUST00000196379.4</a>	2761	No protein	Retained intron	-	-	TSL:1
Ints1-208	<a href="#">ENSMUST00000198615.1</a>	1634	No protein	Retained intron	-	-	TSL:2
Ints1-209	<a href="#">ENSMUST00000200339.1</a>	810	No protein	Retained intron	-	-	TSL:2
Ints1-202	<a href="#">ENSMUST00000196319.1</a>	807	No protein	Retained intron	-	-	TSL:2
Ints1-207	<a href="#">ENSMUST00000197727.1</a>	716	No protein	Retained intron	-	-	TSL:3
Ints1-205	<a href="#">ENSMUST00000197156.1</a>	694	No protein	Retained intron	-	-	TSL:3

The strategy is based on the design of *Ints1-210* transcript,the transcription is shown below:

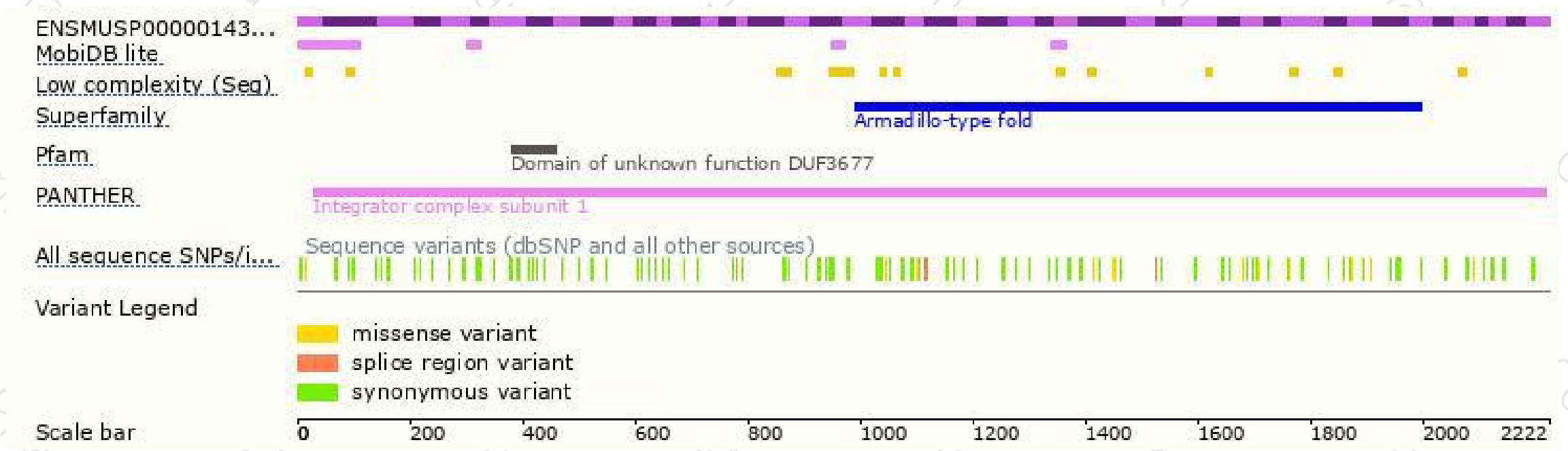


# Genomic location distribution

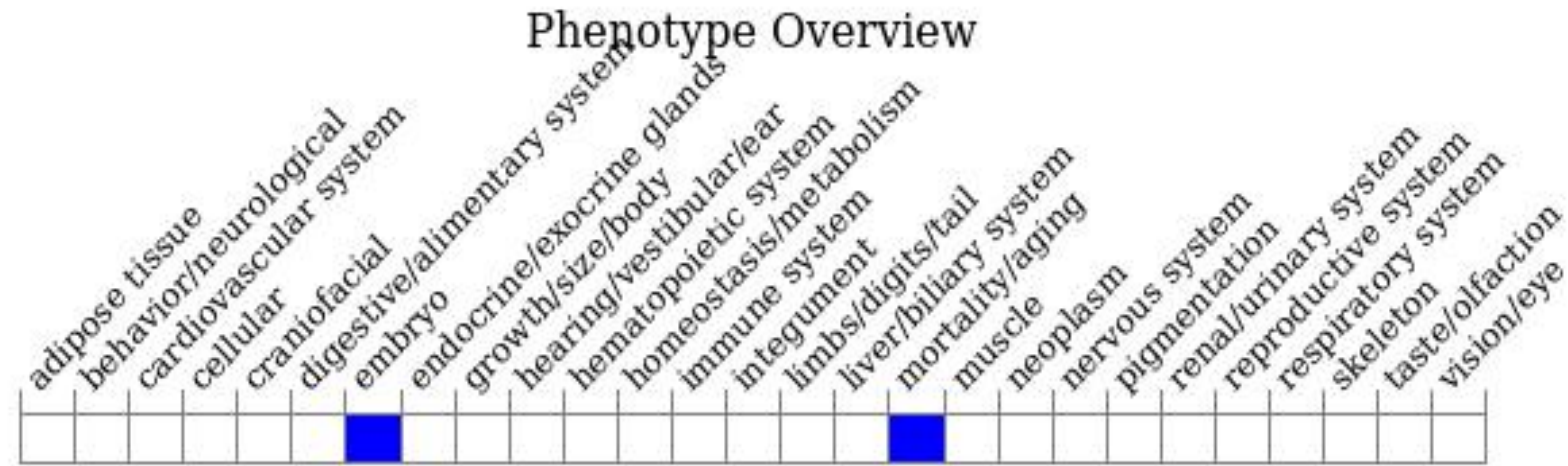




# Protein domain



# 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, homozygous mutation of this gene results in embryonic lethality at the blastocyst stage.

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

Tel: 025-5864 1534

