

# *Slc45a2* Cas9-KO Strategy

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**Reviewer:**

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# Project Overview

**Project Name**

*Slc45a2*

**Project type**

**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Slc45a2* gene. The schematic diagram is as follows:



- The *Slc45a2* gene has 1 transcript. According to the structure of *Slc45a2* gene, exon2-exon4 of *Slc45a2*-201 (ENSMUST00000117100.3) transcript is recommended as the knockout region. The region contains 647bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Slc45a2* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Homozygotes for spontaneous mutations exhibit varied degrees of hypopigmentation of the eyes, skin, and hair, especially the underfur. Eyes are very light at birth but darken with age.
- The *Slc45a2* gene is located on the Chr15. 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)

## Slc45a2 solute carrier family 45, member 2 [Mus musculus (house mouse)]

Gene ID: 22293, updated on 31-Jan-2019

### Summary



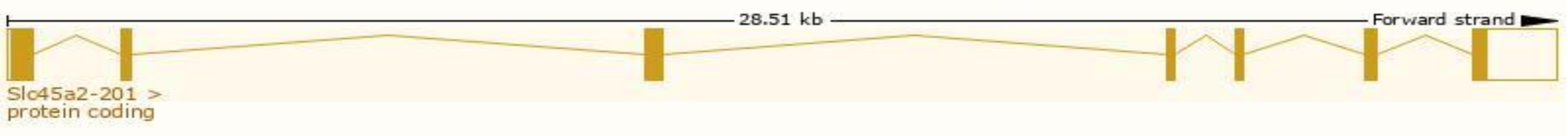
<b>Official Symbol</b>	Slc45a2 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	solute carrier family 45, member 2 provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:2153040</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000022243</a>
<b>Gene type</b>	protein coding
<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>	Aim-1, Aim1, Dbr, Matp, Oca4, blanc-sale, bls, uw
<b>Expression</b>	Low expression observed in reference dataset <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

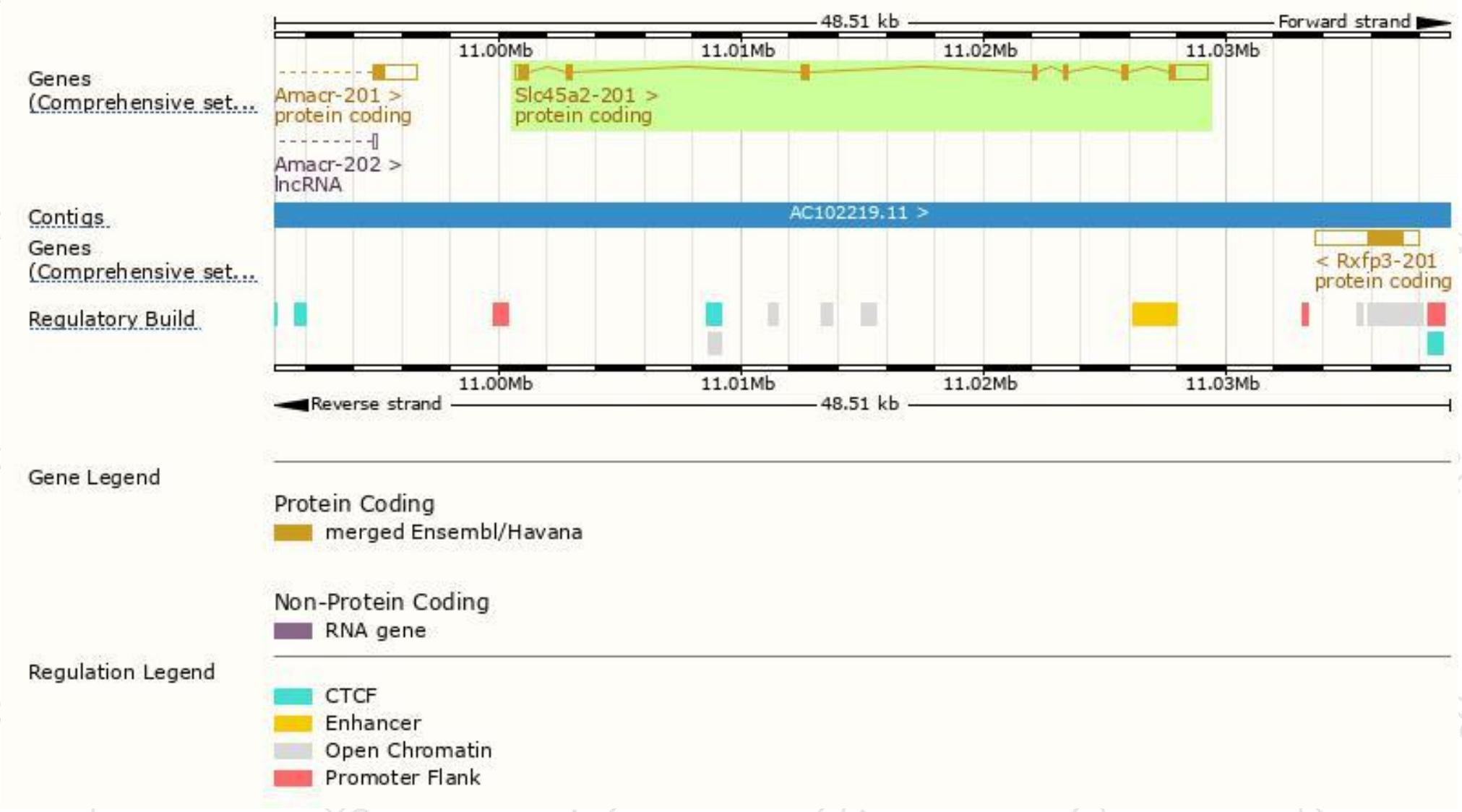
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Slc45a2-201	<a href="#">ENSMUST00000117100.3</a>	2983	<a href="#">530aa</a>	Protein coding	<a href="#">CCDS27382</a>	<a href="#">P58355 Q541S3</a>	TSL:1 GENCODE basic APPRIS P1

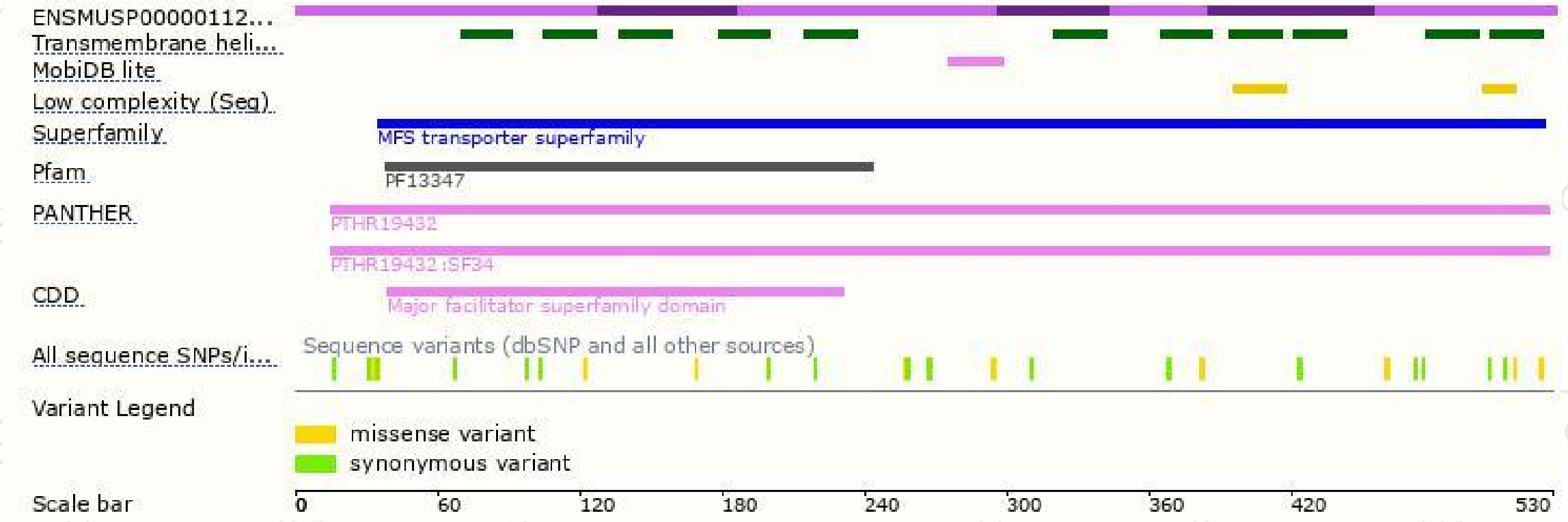
The strategy is based on the design of *Slc45a2-201* transcript, The transcription is shown below



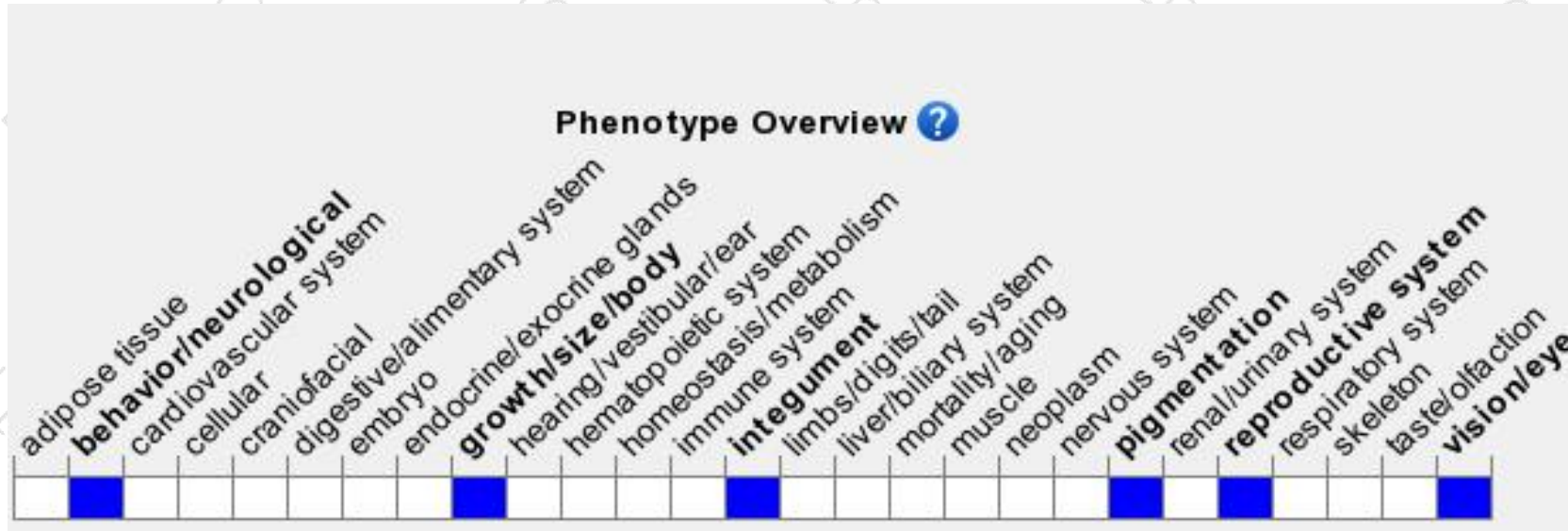
# Genomic location distribution



# 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, Homozygotes for spontaneous mutations exhibit varied degrees of hypopigmentation of the eyes, skin, and hair, especially the underfur. Eyes are very light at birth but darken with age.

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

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