

# ***B4galt1 Cas9-KO Strategy***

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# Project Overview



**Project Name**

***B4galt1***

**Project type**

**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *B4galt1* gene. The schematic diagram is as follows:



- The *B4galt1* gene has 2 transcripts. According to the structure of *B4galt1* gene, exon2 of *B4galt1-201* (ENSMUST00000030121.12) 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 *B4galt1* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Homozygotes for targeted null mutations exhibit growth retardation, low viability, excessive epithelial cell proliferation of skin and small intestine, sperm with reduced fertilizing capacity, birthing difficulty, and mammary gland defects.
- The *B4galt1* gene is located on the Chr4. 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)

## B4galt1 UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 1 [ *Mus musculus* (house mouse) ]

Gene ID: 14595, updated on 27-Aug-2019

### Summary

<b>Official Symbol</b>	B4galt1 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 1 provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:95705</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG00000028413</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	REVIEWED
<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>	GalT; Ggtb; Ggtb2; b4Gal-T1; B-1,4-GalT; B-1,4-GalT1; beta4Gal-T1; beta-1,4-GalT; beta-1,4-GalT1
<b>Summary</b>	This gene encodes two distinct enzyme isoforms, a long membrane-bound form and a short soluble form. These alternate isoforms are thought to be produced through alternative nested transcription initiation and different in-frame start codon usage. These enzymes catalyze the transfer of galactose to acceptor sugars, such as N-acetylglucosamine and glucose. The long form of this enzyme is localized to the trans-Golgi membrane and is involved in glycoconjugate biosynthesis. The short form functions in lactose biosynthesis through formation of a heterodimer with alpha-lactalbumin. [provided by RefSeq, Nov 2012]
<b>Expression</b>	Ubiquitous expression in adrenal adult (RPKM 28.6), ovary adult (RPKM 25.6) and 25 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

### Genomic context

**Location:** 4 A5; 4 20.46 cM

**Exon count:** 7

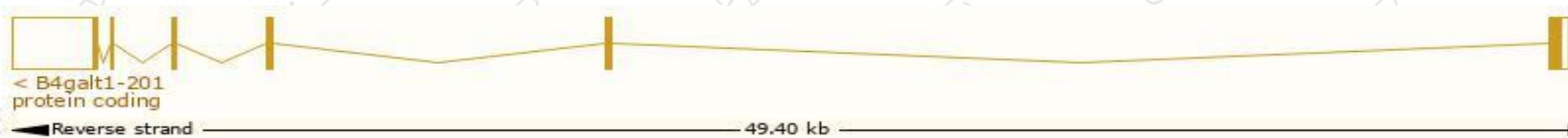
See B4galt1 in [Genome Data Viewer](#)

# Transcript information (Ensembl)

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

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
<b>B4galt1-201</b>	<a href="#">ENSMUST00000030121.12</a>	3983	<a href="#">399aa</a>	Protein coding	<a href="#">CCDS18051</a>	<a href="#">P15535 Q3U478</a>	TSL:1 GENCODE basic APPRIS P1
<b>B4galt1-202</b>	<a href="#">ENSMUST00000108096.2</a>	1413	<a href="#">346aa</a>	Protein coding	-	<a href="#">B1AXY5</a>	TSL:1 GENCODE basic

The strategy is based on the design of *B4galt1-201* transcript, The transcription is shown below



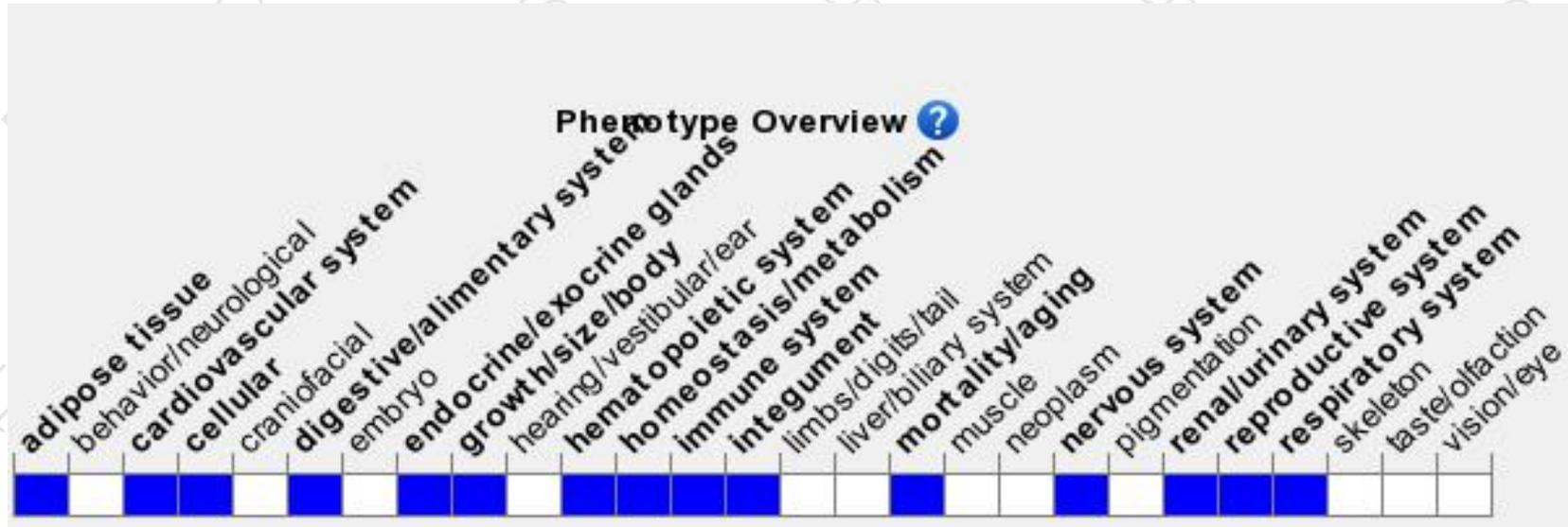
# Genomic location distribution



# 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 targeted null mutations exhibit growth retardation, low viability, excessive epithelial cell proliferation of skin and small intestine, sperm with reduced fertilizing capacity, birthing difficulty, and mammary gland defects.

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

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