# B4galt3 Cas9-KO Strategy 

Designer: Yanhua Shen
Reviewer: Xueting Zhang
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## Project Overview

## Project Name <br> B4galt3

## Project type

Cas9－KO

Strain background

## C57BL／6JGpt

## Knockout strategy

This model will use CRISPR／Cas9 technology to edit the B4galt3 gene．The schematic diagram is as follows：

Wild－type allele


KO allele


## Technical routes

$>$ The B4galt3 gene has 11 transcripts．According to the structure of B4galt3 gene，exon1－exon7 of B4galt3－202
（ENSMUST00000111313．9）transcript is recommended as the knockout region．The region contains most of coding sequence． Knock out the region will result in disruption of protein function．

7 In this project we use CRISPR／Cas9 technology to modify B4galt3 gene．The brief process is as follows：CRISPR／Cas9 system

## Notice

Transcript 207 may be affected．The effect of transcripts 209，210，211 is unknown．
$>$ The $B 4 \mathrm{galt} 3$ gene is located on the Chrl．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）

## B4galt3 UDP－Gal：betaGIcNAc beta 1，4－galactosyltransferase，polypeptide 3 ［ Mus musculus（house mouse）］

Gene ID：57370，updated on 10－Sep－2019

## Summary

Official Symbol B4galt3 provided by MGI
Official Full Name UDP－Gal：betaGIcNAc beta 1，4－galactosyltransferase，polypeptide 3 provided by MGI
Primary source MGI：MGI：1928767
See related Ensembl：ENSMUSG00000052423
Gene type protein coding
RefSeq status VALIDATED
Organism Mus musculus
Lineage Eukaryota；Metazoa；Chordata；Craniata；Vertebrata；Euteleostomi；Mammalia；Eutheria；Euarchontoglires；Glires；Rodentia；Myomorpha；Muroidea；Muridae； Murinae；Mus；Mus
Also known as ESTM6；ESTM26；R74981；AA104562；AW125175；9530061M23Rik
Expression Ubiquitous expression in duodenum adult（RPKM 38．6），ovary adult（RPKM 29．1）and 28 other tissues See more
Orthologs human all

## Genomic context

Location： $1 \mathrm{H} 3 ; 179.29 \mathrm{~cm}$
See B4galt3 in Genome Data Viewer
Exon count： 8

## Transcript information（Ensembl）

The gene has 11 transcripts，all transcripts are shown below：

| Name | Transcript ID | bp | Protein | Biotype | CCDS | UniProt | Flags |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| B4galt3－202 | ENSMUST00000111313．9 | 2098 | 395aa | Protein coding | CCDS15486 | Q3U260 Q91YY2 | TSL：1 GENCODE basic APPRIS P1 |
| B4galt3－201 | ENSMUST00000064272．9 | 1998 | 395aa | Protein coding | CCDS15486 | Q3U260 Q91YY2 | TSL：1 GENCODE basic APPRIS P1 |
| B4galt3－205 | ENSMUST00000126699．3 | 1758 | 173aa | Protein coding | － | A0A0A6YXE7 | TSL： 5 GENCODE basic |
| B4galt3－211 | ENSMUST00000151863．7 | 869 | 13aa | Protein coding | － | AOAOA6YXS2 | CDS 3 ＇incomplete TSL：5 |
| B4galt 3－210 | ENSMUST00000141999．7 | 583 | 65 aa | Protein coding | － | D3YUU0 | CDS 3 ＇incomplete TSL： 3 |
| B4galt3－209 | ENSMUST00000141114．1 | 446 | 139aa | Protein coding | － | D3YVA5 | CDS $3^{\prime}$ incomplete TSL： 3 |
| B4galt3－207 | ENSMUST00000132890．1 | 3261 | No protein | Retained intron | － | － | TSL： 1 |
| B4galt3－206 | ENSMUST00000129985．2 | 2805 | No protein | Retained intron | － | － | TSL：5 |
| B4galt3－204 | ENSMUST00000125939．1 | 728 | No protein | Retained intron | － | － | TSL：3 |
| B4galt3－203 | ENSMUST00000123954．7 | 566 | No protein | Retained intron | － | － | TSL：2 |
| B4galt3－208 | ENSMUST00000138904．1 | 440 | No protein | IncRNA | － | － | TSL：2 |

The strategy is based on the design of B4galt3－202 transcript，The transcription is shown below


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## Genomic location distribution



## Protein domain

ENSMUSP00000106．．． Transmembrane heli．．．
MobiDB lite．
Low complexity（Seq）．
Superfamily．
Prints
Pfam

$$
\square
$$

Nucleotide－diphospho－sugar transferases


PANTHER
Beta－1，4－galactosy｜transferase
गTHR19300：SF33
Gene3D．
CDD
All seguence $\mathrm{SNPs} / \mathrm{i} . .$.


Variant Legendmissense variant
synonymous variant
Scale bar
40


Sequence variants（dbSNP and all other sources）
$\square$
 0
$0 \quad 40$

If you have any questions, you are welcome to inquire. Tel: 400-9660890



[^0]:    B4galt3－202 $>$
    protein coding

