

Slc9a3 Cas9-CKO Strategy

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Project Overview

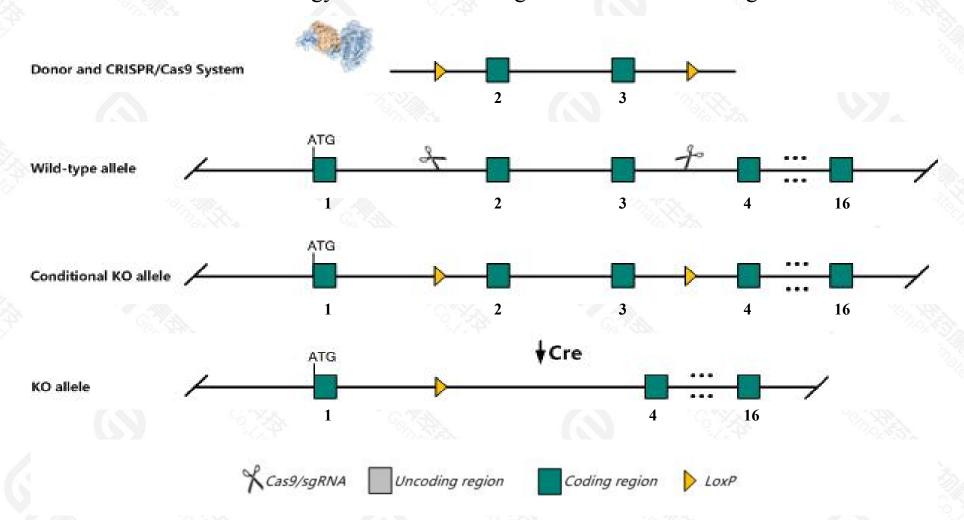


Project Name	Slc9a3			
Project type	Cas9-CKO			
Strain background	C57BL/6JGpt			

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Slc9a3* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Slc9a3* gene has 4 transcripts. According to the structure of *Slc9a3* gene, exon2-exon3 of *Slc9a3*-201(ENSMUST00000036208.7) transcript is recommended as the knockout region. The region contains 464bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Slc9a3* gene. The brief process is as follows:sgRNA was transcribed in vitro, donor 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.

Notice



- > According to the existing MGI data, homozygous mutant mice have diarrhea associated with defects of renal and intestinal absorption. Males are infertile.
- > The *Slc9a3* gene is located on the Chr13. 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)



Slc9a3 solute carrier family 9 (sodium/hydrogen exchanger), member 3 [Mus musculus (house mouse)]

Gene ID: 105243, updated on 2-Mar-2021

Summary



Official Symbol Slc9a3 provided by MGI

Official Full Name solute carrier family 9 (sodium/hydrogen exchanger), member 3 provided by MGI

Primary source MGI:MGI:105064

See related Ensembl:ENSMUSG00000036123

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 9030624013Rik, Al930210, NHE, NHE-, NHE-3, NHE3

Expression Biased expression in colon adult (RPKM 52.1), small intestine adult (RPKM 51.8) and 4 other tissuesSee more

Orthologs <u>human</u> all

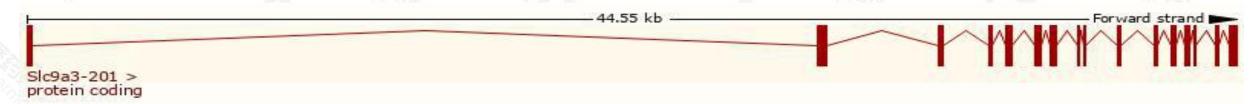
Transcript information (Ensembl)



The gene has 4 transcripts, all transcripts are shown below:

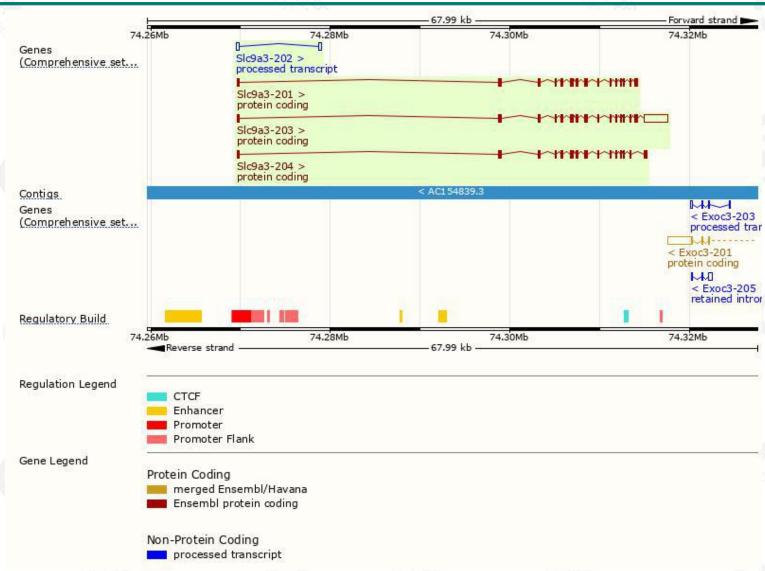
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Slc9a3-201	ENSMUST00000036208.7	2490	829aa	Protein coding	CCDS36731		TSL:2, GENCODE basic, APPRIS P2,
Slc9a3-203	ENSMUST00000221703.2	5118	829aa	Protein coding	-		TSL:5 , GENCODE basic , APPRIS ALT2 ,
Slc9a3-204	ENSMUST00000225423.2	2589	<u>862aa</u>	Protein coding	2		GENCODE basic , APPRIS ALT2 ,
Slc9a3-202	ENSMUST00000221410.2	578	No protein	Processed transcript	-		TSL:3,

The strategy is based on the design of *Slc9a3-201* transcript, the transcription is shown below:



Genomic location distribution





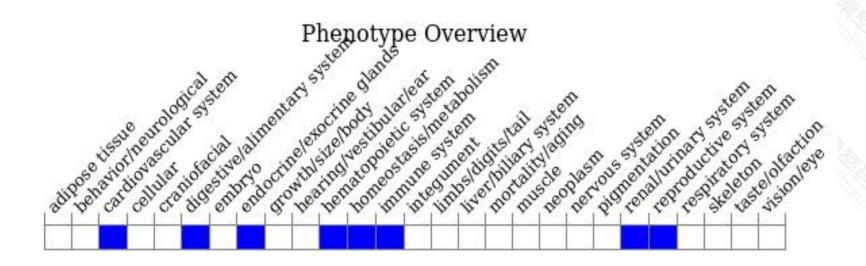
Protein domain





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, homozygous mutant mice have diarrhea associated with defects of renal and intestinal absorption. Males are infertile.



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

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