

# ***Slc12a2-T197A&T201A&T206A* Mouse Model Strategy**

## **-CRISPR/Cas9 technology**

**Designer: Daohua Xu**

**Reviewer: Jia Yu**

**Design Date: 2020-12-8**

# Project Overview

**Project Name**

**Slc12a2-T197A&T201A&T206A**

**Project type**

**cas9-ki(PM)**

**Strain background**

**C57BL/6JGpt**

# Technical Description

- The mouse *Slc12a2* gene has 2 transcripts.
- This project produced *Slc12a2-T197A&T201A&T206A* point mutation on exon1 of the transcript of *Slc12a2-201*(ENSMUST00000115366.2). The 197th,201th and 206th amino acid will be mutated from T to A.
- In this project, *Slc12a2* gene will be modified by CRISPR/Cas9 technology. The brief process is as follows: In vitro, sgRNA and donor vectors were constructed. Cas9, sgRNA and donor were injected into the fertilized eggs of C57BL/6JGpt mice for homologous recombination, and obtained positive F0 mice identified by PCR and sequencing analysis. The stable inheritable positive F1 mice model was obtained by mating F0 mice with C57BL/6JGpt mice.

# Analysis of Homology

	204	210	220	230	240	250	260	270	280	290
NKCC1-mouse-201	197	THTNTYYLRTFGHNTMDAVPRIDHYRHTAAQLGEKLLRPSLAELHDELEKEPFEDGFANGEEESTPTRDAVVAYTAESKGVVKFGWIKGVLVF								
NKCC1-human-201	203	THTNTYYLRTFGHNTMDAVPRIDHYRHTAAQLGEKLLRPSLAELHDELEKEPFEDGFANGEEESTPTRDAVVYTAESKGVVKFGWIKGVLVF								
Consensus	204	THTNTYYLRTFGHNTMDAVPRIDHYRHTAAQLGEKLLRPSLAELHDELEKEPFEDGFANGEEESTPTRDAVV YTAESKGVVKFGWIKGVLVF								
Ready		consensus positions: 95.5% identity positions: 93.9% aln: 265								

Identity positions: 93.9%



# Mutation Site

Before mutation

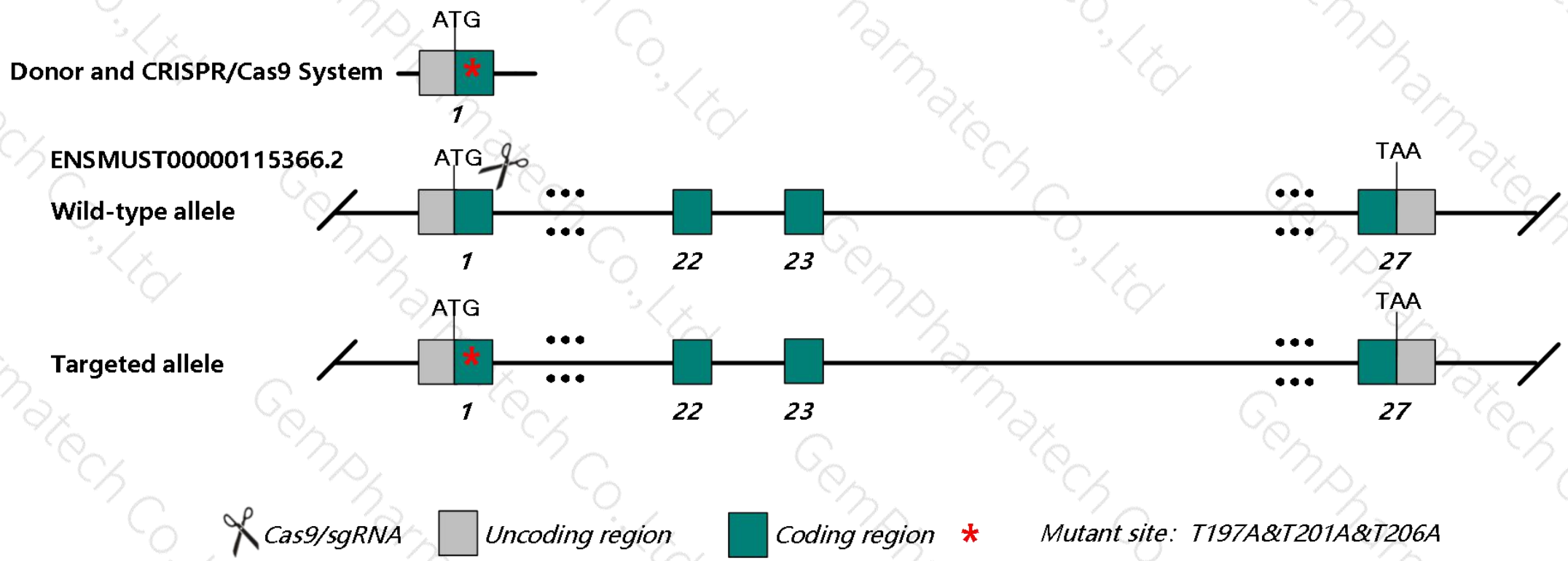
	+3	V	G	G	D	G	P	N	V	S	F	Q	N	G	G	D	T	V	L	S	E	G	S	S	L	H	S	G	G	G	S	G	H	H	Q			
20601		G	T	C	G	G	T	G	G	C	A	A	C	G	G	C	A	A	C	G	G	T	G	C	T	T	C	A	G	A	C	G	G	T	G	G	T	
		C	A	G	C	C	A	C	C	G	C	T	G	C	A	A	G	G	T	G	C	C	G	G	T	T	G	C	A	A	G	G	T	T	G	G	T	
	+3	?Q	Q	Y	Y	Y	D	T	H	T	N	T	Y	Y	L	R	T	F	G	H	N	T	M	D	A	V	P	R	I	D	H	Y	R	H	T			
20701		A	G	C	A	G	T	A	C	T	A	C	T	A	C	T	A	C	T	A	C	T	A	C	T	A	C	T	A	C	T	A	C	T	A	C	T	
		T	C	G	T	C	A	T	G	A	T	G	A	T	G	A	T	G	A	T	G	A	T	G	A	T	G	A	T	G	A	T	G	A	T	G	A	T

After mutation

	+3	V	G	G	D	G	P	N	V	S	F	Q	N	G	G	D	T	V	L	S	E	G	S	S	L	H	S	G	G	G	S	G	H	H	Q	
20601		G	T	C	G	G	T	G	G	C	G	A	C	G	G	C	G	G	T	G	C	T	T	C	A	G	A	C	G	G	T	G	G	T	G	G
		C	A	G	C	C	A	C	C	G	C	T	G	C	A	A	G	G	T	G	C	C	G	G	T	T	G	C	A	A	G	G	T	T	G	G
	+3	?Q	Q	Y	Y	Y	D	A	H	T	N	A	Y	Y	L	R	A	F	G	H	N	T	M	D	A	V	P	R	I	D	H	Y	R	H	T	
20701		A	G	C	A	G	T	A	C	T	A	C	T	A	C	T	A	C	T	A	C	T	A	C	T	A	C	T	A	C	T	A	C	T	A	C
		T	C	G	T	C	A	T	G	A	T	G	A	T	G	A	T	G	A	T	G	A	T	G	A	T	G	A	T	G	A	T	G	A	T	G

The green region is exon1 of *Slc12a2-201*, and the yellow region represents the p.T197A&T201A&T206A mutation site.

This model uses CRISPR/Cas9 technology to edit the *Slc12a2* gene and the schematic diagram is as follow:



- According to the data of MGI, homozygous mutants show variably severe deafness, head-shaking, circling, reduced endolymph secretion, male sterility, growth retardation, hypotension, reduced salivation, delayed ductal outgrowth of mammary epithelium and increased periweaning mortality.
- Two to four synonymous mutations of amino acids would be introduced on exon1 of *Slc12a2*.
- Mouse *Slc12a2* gene is located on Chr18. Please take the loci in consideration when breeding this mutation mice with other gene modified strains, if the other gene is also on Chr18, it may be extremely hard to get double gene positive homozygotes.
- The scheme is designed according to the genetic information in the existing database. Due to the complex process of gene transcription and translation, it cannot be predicted completely at the present technology level.

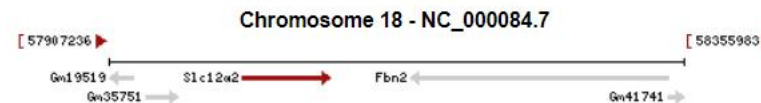
# Gene name and location (NCBI)

## Slc12a2 solute carrier family 12, member 2 [ *Mus musculus* (house mouse) ]

Gene ID: 20496, updated on 22-Nov-2020

### Summary

**Official Symbol** Slc12a2 provided by [MGI](#)  
**Official Full Name** solute carrier family 12, member 2 provided by [MGI](#)  
**Primary source** [MGI:MGI:101924](#)  
**See related** [Ensembl:ENSMUSG00000024597](#)  
**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** sy; Nkcc; mBSC; Nkcc1; mBSC2; sy-ns; 9330166H04R; 9330166H04Rik  
**Expression** Broad expression in mammary gland adult (RPKM 43.3), colon adult (RPKM 19.3) and 20 other tissues [See more](#)  
**Orthologs** [human](#) [all](#)





# Transcript information (Ensembl)

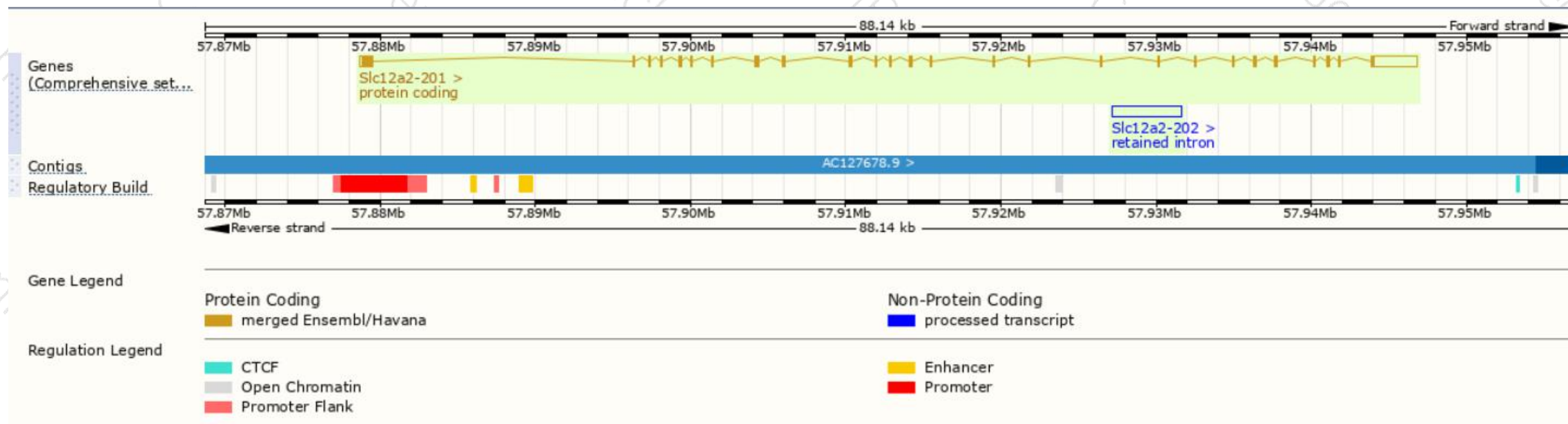
The gene has 2 transcripts, and all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt Match	Flags
Slc12a2-201	<a href="#">ENSMUST00000115366.2</a>	6520	<a href="#">1206aa</a>	Protein coding	<a href="#">CCDS37826</a>	<a href="#">E9QM38</a>	TSL:1 GENCODE basic APPRIS P1
Slc12a2-202	<a href="#">ENSMUST00000236673.1</a>	4480	No protein	Retained intron	-	-	-

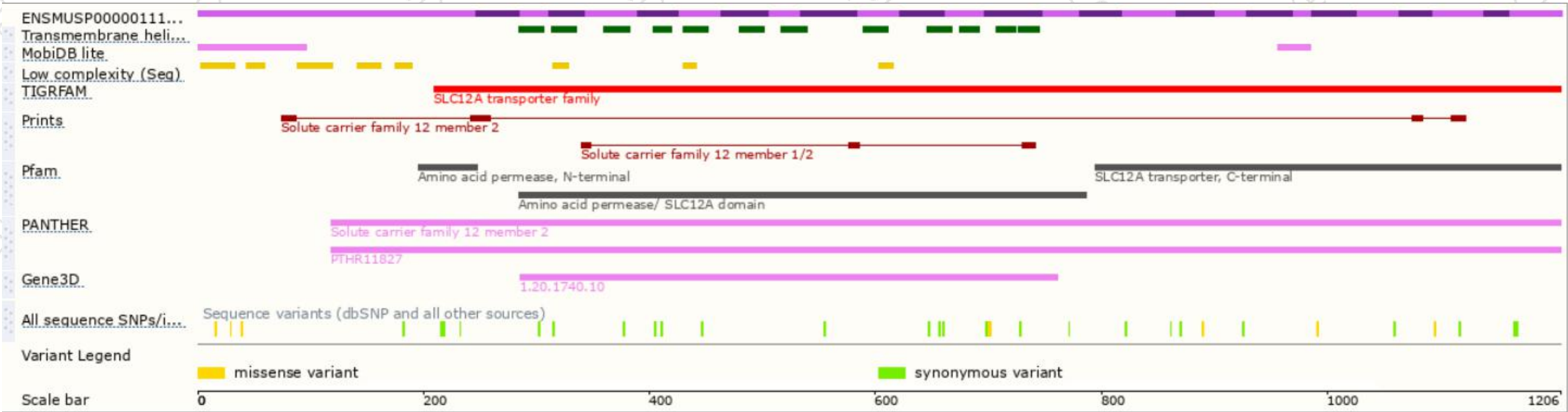
The strategy is based on the design of *Slc12a2-201* transcript, the transcription is shown below:



# Genomic location distribution



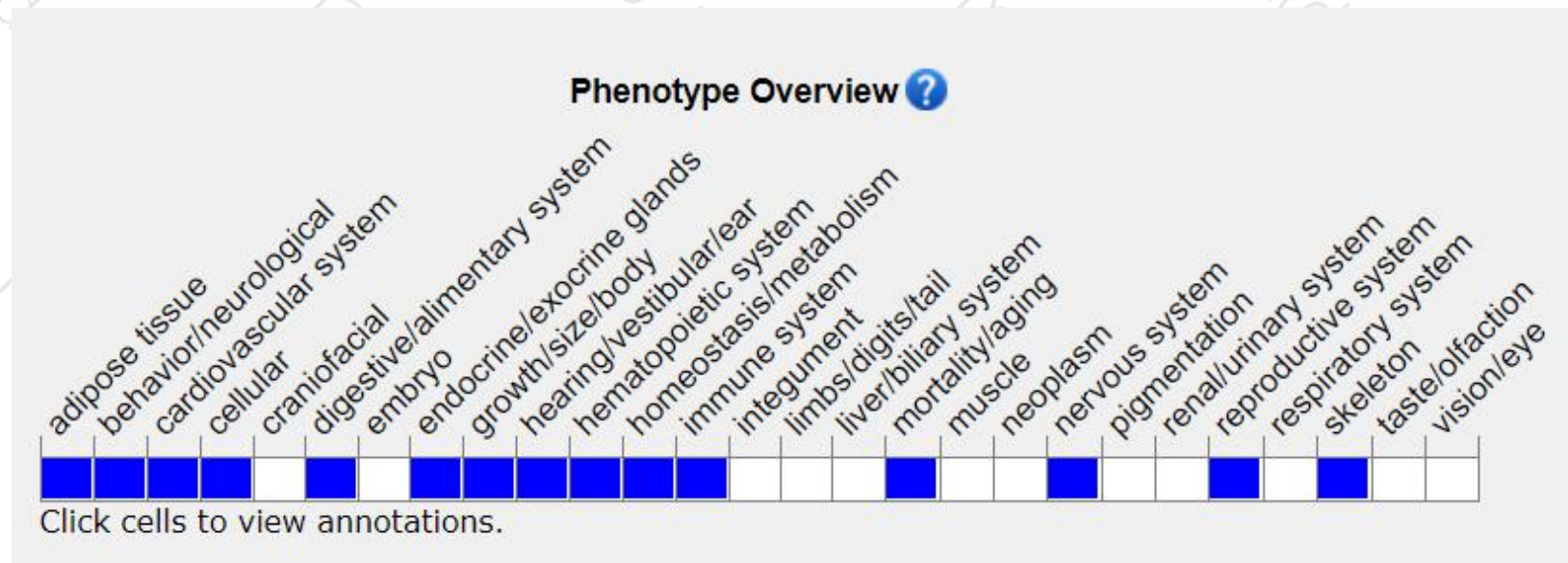
# Protein domain



# Mouse phenotype description(MGI )

URL link is as follows:

<http://www.informatics.jax.org/marker/MGI:101924>



Homozygous mutants show variably severe deafness, head-shaking, circling, reduced endolymph secretion, male sterility, growth retardation, hypotension, reduced salivation, delayed ductal outgrowth of mammary epithelium and increased periweaning mortality.



If you have any questions, please feel free to contact us.

Tel: 025-5864 1534



集萃药康生物科技  
GemPharmatech Co.,Ltd

