

Gsto1 Cas9-CKO Strategy

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Project Overview



Project Name Gsto1

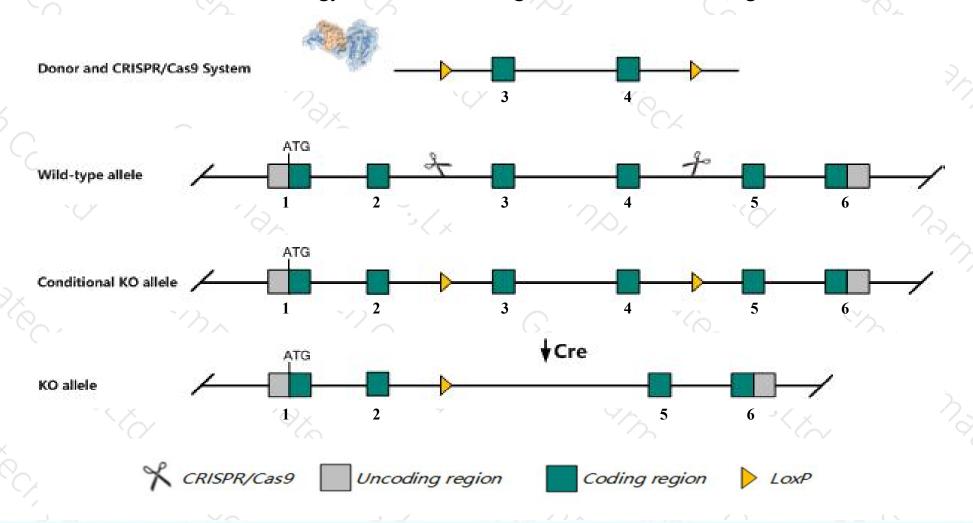
Project type Cas9-CKO

Strain background C57BL/6JGpt

Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Gsto1* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Gsto1* gene has 6 transcripts. According to the structure of *Gsto1* gene, exon3-exon4 of *Gsto1-201* (ENSMUST00000026050.7) transcript is recommended as the knockout region. The region contains 322bp coding sequence. Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Gsto1* gene. The brief process is as follows:CRISPR/Cas9 system 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 will be 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, Mice homozygous for a mutant allele do not display abnormal arsenate metabolism following acute arsenate exposure.
- The floxed region is near to the N-terminal of *Gsto2* gene, this strategy may influence the regulatory function of the N-terminal of *Gsto2* gene.
- ➤ Transcript *Gsto1*-202&203 may not be affected.
- > The partial sequence of intron of *Cfap43* gene will be deleted.
- The *Gsto1* gene is located on the Chr19. 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)



Gsto1 glutathione S-transferase omega 1 [Mus musculus (house mouse)]

Gene ID: 14873, updated on 10-Oct-2019

Summary

☆ ?

Official Symbol Gsto1 provided by MGI

Official Full Name glutathione S-transferase omega 1 provided by MGI

Primary source MGI:MGI:1342273

See related Ensembl: ENSMUSG00000025068

Gene type protein coding
RefSeq status REVIEWED
Organism Mus musculus

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;

Muroidea; Muridae; Murinae; Mus; Mus

Also known as p28; Gstx; Spg-r; Gsto-1; AA407097; AI194287; AU018802

Summary This gene encodes a member of the omega class of glutathione S-transferase (GST) proteins. GSTs are involved in the metabolism of xenobiotics

and carcinogens. There is evidence that the encoded protein is involved in the biotransformation of arsenic. [provided by RefSeq, Dec 2015]

Expression Broad expression in stomach adult (RPKM 297.2), bladder adult (RPKM 158.9) and 20 other tissues See more

Orthologs human all

Genomic context



Location: 19; 19 D1 See Gsto1 in Genome Data Viewer

Exon count: 6

Annotation release	Status	Assembly		Location		
108	current	GRCm38.p6 (GCF_000001635.26)	19	NC_000085.6 (4785497347864790)		
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	19	NC_000085.5 (4792947947939280)		

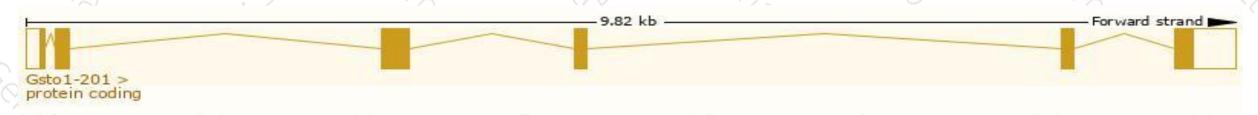
Transcript information (Ensembl)



The gene has 6 transcripts, all transcripts are shown below:

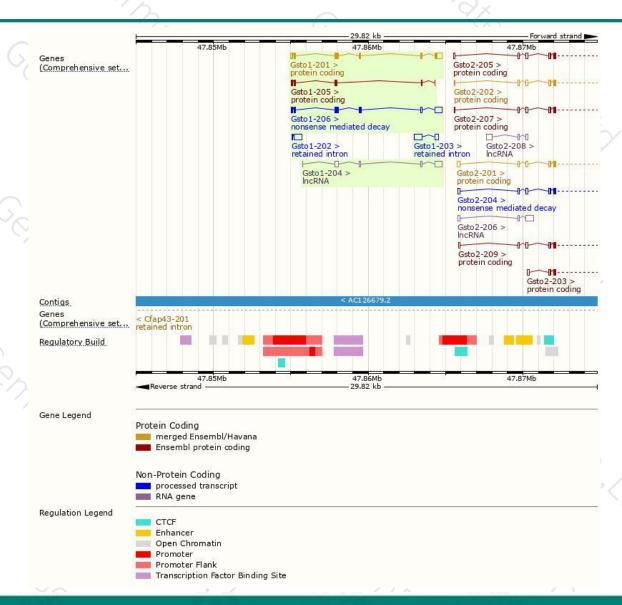
					122			
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags	
Gsto1-201	ENSMUST00000026050.7	1182	240aa	Protein coding	CCDS29893	009131	TSL:1 GENCODE basic APPRIS P1	
Gsto1-205	ENSMUST00000237121.1	384	<u>108aa</u>	Protein coding		*	CDS 3' incomplete	
Gsto1-206	ENSMUST00000237472.1	1138	<u>166aa</u>	Nonsense mediated decay	4-	22		
Gsto1-203	ENSMUST00000143694.1	759	No protein	Retained intron	92	20	TSL:2	
Gsto1-202	ENSMUST00000130157.1	551	No protein	Retained intron	-	-	TSL:1	
Gsto1-204	ENSMUST00000236902.1	950	No protein	IncRNA	-	-5		

The strategy is based on the design of *Gsto1-201* transcript, The transcription is shown below



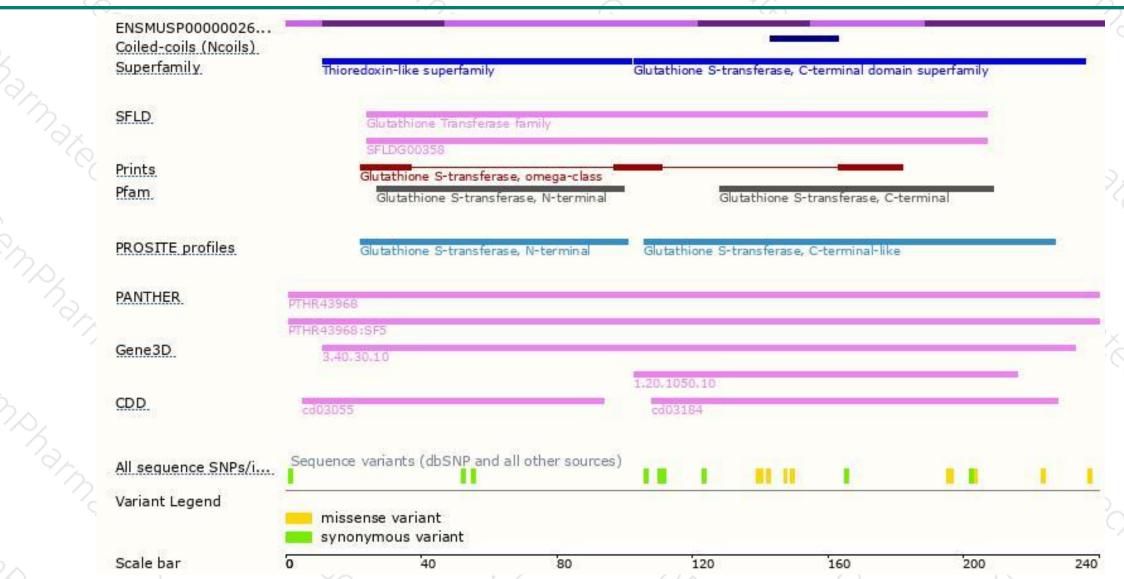
Genomic location distribution





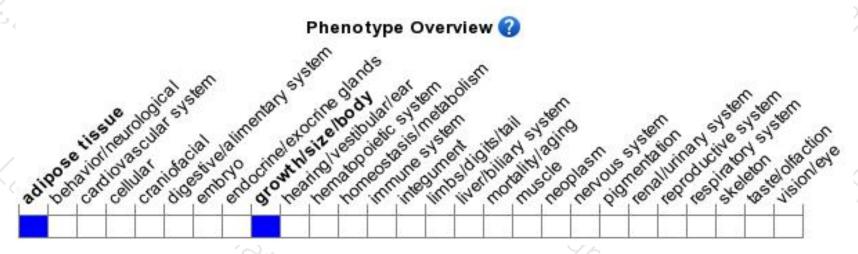
Protein domain





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, Mice homozygous for a mutant allele do not display abnormal arsenate metabolism following acute arsenate exposure.



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





