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SLC22A8 Knockout 293T Cell Line, Homozygous

Catalog No.: RM02122

Basic Information

Catalog No.
RM02122

Category
Cell Line

Parental Cell line
293T

Genotype
Knockout

Background

This gene encodes a protein involved in the sodium-independent transport and excretion of organic anions, some of which are potentially toxic. The encoded protein is an integral membrane protein and appears to be localized to the basolateral membrane of the kidney. Multiple alternatively spliced transcript variants that encode different protein isoforms have been described for this gene. [provided by RefSeq, May 2010]

Gene Information

Gene Symbol
SLC22A8

Species
Human

Gene ID
9376

Swiss Prot
Q8TCC7

Synonyms
OAT3

Product Information

Description
SLC22A8 Knockout 293T Cell Line is engineered from 293T cell line with Gene-Editing Technology.
Allele-1:35bp deletion in exon2
Allele-2:35bp deletion in exon2
Mammalian cells such as human, rat and mouse cells are normally diploid with two alleles. Homozygote: both alleles were knocked out, mRNA has no signal, no expression of proteins. Heterozygote: only one allele was knocked out, the mRNA transcript levels was decreased compared to wild type, and the protein expression levels was also lower than that of the wild type.

Packaging
1 vial parental cell line and 1 vial knockout cell line

Shipping Conditions
Dry ice

Amount
1~5x10^6 cells/vial

Storage
Store at -20℃

Protocol
Upon arrival, it should be maintained in DMEM medium with 10%(v/v) fetal bovine serum and 100U penicillin-streptomycin, at 37℃ with 5% CO_2 condition.
1. Thaw the vial in 37℃ water bath ,and shake it to melt as soon as possible.
2. Transfer the cell suspension to a 15mL conical tube with pre-warmed 5mL complete medium and centrifuge 1000rpm for approximately 5 minutes at room temperature.
3. Remove and discard the supernatant.
4. Resuspend the cell pellet with 1mL pre-warmed complete medium and seed in 10cm dish.
5. Add 8-10mL of complete medium.
6. Incubate the culture at 37℃ incubator with 5% CO_2.
7. A subcultivation ratio of 1:2-1:4 is recommended.

Contact

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Sequencing data

WT GCTGGCAGCCAGCG**********TACATGGTCTTCCG
Mut GCTGGCAGCCAGCG***Deletion***TACATGGTCTTCCG
Allele-1: 35bp deletion in exon2

WT GCTGGCAGCCAGCG**********TACATGGTCTTCCG
Mut GCTGGCAGCCAGCG***Deletion***TACATGGTCTTCCG
Allele-2: 35bp deletion in exon2

Genome sequence analysis of PCR products from parental (WT) and SLC22A8 knockout (KO) 293T cells, using sanger sequencing.