

Product datasheet for SC208654

GLUT8 (SLC2A8) (NM 014580) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: GLUT8 (SLC2A8) (NM_014580) Human 3' UTR Clone

Symbol: GLUT8

Synonyms: GLUT8; GLUTX1

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_014580

Insert Size: 679 bp

Insert Sequence: >SC208654 3'UTR clone of NM_014580

The sequence shown below is from the reference sequence of NM_014580. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).



OriGene Technologies, Inc. 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com



GLUT8 (SLC2A8) (NM_014580) Human 3' UTR Clone - SC208654

Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

RefSeq: <u>NM 014580.5</u>

Summary: This gene belongs to the solute carrier 2A family, which includes intracellular glucose

transporters. Based on sequence comparison, the glucose transporters are grouped into three classes and this gene is a member of class II. The encoded protein, like other members of the family, contains several conserved residues and motifs and 12 transmembrane

domains with both amino and carboxyl ends being on the cytosolic side of the membrane. Alternatively spliced transcript variants have been described for this gene. [provided by

RefSeq, Nov 2012]

Locus ID: 29988

MW: 25.2