

Product datasheet for SC208051

OriGene Technologies, Inc.

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Glucose 6 Phosphate Dehydrogenase (G6PD) (NM 000402) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: Glucose 6 Phosphate Dehydrogenase (G6PD) (NM_000402) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: G6PD Synonyms: G6PD1

ACCN: NM_000402

Insert Size: 650 bp

Insert Sequence: >SC208051 3'UTR clone of NM_000402

The sequence shown below is from the reference sequence of NM_000402. 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).

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.



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MW:

RefSeq: NM 000402.4

Summary: This gene encodes glucose-6-phosphate dehydrogenase. This protein is a cytosolic enzyme

encoded by a housekeeping X-linked gene whose main function is to produce NADPH, a key

electron donor in the defense against oxidizing agents and in reductive biosynthetic reactions. G6PD is remarkable for its genetic diversity. Many variants of G6PD, mostly

produced from missense mutations, have been described with wide ranging levels of enzyme activity and associated clinical symptoms. G6PD deficiency may cause neonatal jaundice, acute hemolysis, or severe chronic non-spherocytic hemolytic anemia. Two transcript variants

encoding different isoforms have been found for this gene. [provided by RefSeq, Jul 2008]

Locus ID: 2539

22.8