

## Product datasheet for **SC202086**

### PHGDH (NM\_006623) Human 3' UTR Clone

#### Product data:

Product Type:	3' UTR Clones
Product Name:	PHGDH (NM_006623) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	PHGDH
Synonyms:	3-PGDH; 3PGDH; HEL-S-113; NLS; NLS1; PDG; PGAD; PGD; PGDH; PHGDHD; SERA
ACCN:	NM_006623
Insert Size:	205 bp
Insert Sequence:	>SC202086 3'UTR clone of NM_006623 The sequence shown below is from the reference sequence of NM_006623. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site  GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC GTGACTGAAGCCTTCCAGTTCACCTTCTAACCTTGGAGCTCACTGGTCCCTGCCTCTGGGGCTTTTCTG AAGAAACCCACCCACTGTGATCAATAGGGAGAGAAAAATCCACATTCTTGGGCTGAACGCGGCCTCTGA CACTGCTTACACTGCACTCTGACCCTGTAGTACAGCAATAACCGTCTAATAAAGAGCCTACCCCAA ACGCGTAAGCGGCCGCGCATCTAGATTGAAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA CGAGATTCGATTCCACCGCCCTTCTATGAAAGG
Restriction Sites:	Sgfl-MluI
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.
RefSeq:	<a href="#">NM_006623.4</a>



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**Summary:** This gene encodes the enzyme which is involved in the early steps of L-serine synthesis in animal cells. L-serine is required for D-serine and other amino acid synthesis. The enzyme requires NAD/NADH as a cofactor and forms homotetramers for activity. Mutations in this gene have been found in a family with congenital microcephaly, psychomotor retardation and other symptoms. Multiple alternatively spliced transcript variants have been found, however the full-length nature of most are not known. [provided by RefSeq, Aug 2011]

**Locus ID:** 26227

**MW:** 7.6