

Product datasheet for **SC206369**

GSTM4 (NM_147148) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	GSTM4 (NM_147148) Human 3' UTR Clone
Symbol:	GSTM4
Synonyms:	GSTM4-4; GTM4
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_147148
Insert Size:	496 bp
Insert Sequence:	>SC206369 3'UTR clone of NM_147148 The sequence shown below is from the reference sequence of NM_147148. The complete sequence of this clone may contain minor differences, such as SNPs. Blue =Stop Codon Red =Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
CGCTTTGAGGTTTCTGTGGCATAATGTGATGGTCAATTTTCTGCATCAACTTGACTGGGCTAAGGGAT
GCTCAGATGGCAGGTAATAATCATTGTGCTTGTGAGGGTGTTCAGAAAGAGATTTGCCTTTGAATCAGA
AGACAGCAAAGATTTCTTCAGCAATGAAGGAGGCATCCACCAAAGTGCAGGGCCAGAGAGAAGAAA
AAGACAGGAAGGGTGAATTTGACCTCTGACTGGGACATCCATCTCTGCCTATCTGGGACCTCCACA
CTCCTGGTTCTCTGGCCTTCAGACTTGATCAGGGACTAACACCATCGCCTCCCACCCCACTTTGTTC
TGAGGCCTTTAGCCTCTGAATGATACCACTGGCTTTCTGCTTCTATCCTGCAGTCGGCAGATCATG
GGACTTCTTCACTCCAAAATTGTGTGAGCCAATCCATAACAGATAGATAAATTTATAAATAAACACA
CAAATTTCTTACA
ACGCGTAAGCGGCCGCGGCATCTAGATTCTGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
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Restriction Sites:	SgfI-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).



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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_147148.3</u>
Summary:	Cytosolic and membrane-bound forms of glutathione S-transferase are encoded by two distinct supergene families. At present, eight distinct classes of the soluble cytoplasmic mammalian glutathione S-transferases have been identified: alpha, kappa, mu, omega, pi, sigma, theta and zeta. This gene encodes a glutathione S-transferase that belongs to the mu class. The mu class of enzymes functions in the detoxification of electrophilic compounds, including carcinogens, therapeutic drugs, environmental toxins and products of oxidative stress, by conjugation with glutathione. The genes encoding the mu class of enzymes are organized in a gene cluster on chromosome 1p13.3 and are known to be highly polymorphic. These genetic variations can change an individual's susceptibility to carcinogens and toxins as well as affect the toxicity and efficacy of certain drugs. Diversification of these genes has occurred in regions encoding substrate-binding domains, as well as in tissue expression patterns, to accommodate an increasing number of foreign compounds. Multiple transcript variants, each encoding a distinct protein isoform, have been identified. [provided by RefSeq, Jul 2008]
Locus ID:	2948
MW:	18.4