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Product datasheet for PH304814

Thymidylate Synthase (TYMS) (NM_001071) Human Mass Spec Standard

Product data:

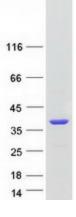
Product Type:	Mass Spec Standards
Description:	TYMS MS Standard C13 and N15-labeled recombinant protein (NP_001062)
Species:	Human
Expression Host:	HEK293
Expression cDNA Clone or AA Sequence:	RC204814
Predicted MW:	35.5 kDa
Protein Sequence:	<pre>>RC204814 representing NM_001071 Red=Cloning site Green=Tags(s)</pre>
	MPVAGSELPRRPLPPAAQERDAEPRPPHGELQYLGQIQHILRCGVRKDDRTGTGTLSVFGMQARYSLRDE FPLLTTKRVFWKGVLEELLWFIKGSTNAKELSSKGVKIWDANGSRDFLDSLGFSTREEGDLGPVYGFQWR HFGAEYRDMESDYSGQGVDQLQRVIDTIKTNPDDRRIIMCAWNPRDLPLMALPPCHALCQFYVVNSELSC QLYQRSGDMGLGVPFNIASYALLTYMIAHITGLKPGDFIHTLGDAHIYLNHIEPLKIQLQREPRPFPKLR ILRKVEKIDDFKAEDFQIEGYNPHPTIKMEMAV
	TRTRPLEQKLISEEDLAANDILDYKDDDDKV
Tag:	C-Myc/DDK
Purity:	> 80% as determined by SDS-PAGE and Coomassie blue staining
Concentration:	>0.05 µg/µL as determined by microplate BCA method
Labeling Method:	Labeled with [U- 13C6, 15N4]-L-Arginine and [U- 13C6, 15N2]-L-Lysine
Buffer:	25 mM Tris-HCl, 100 mM glycine, pH 7.3
Storage:	Store at -80°C. Avoid repeated freeze-thaw cycles.
Stability:	Stable for 3 months from receipt of products under proper storage and handling conditions.
RefSeq:	<u>NP 001062</u>
RefSeq Size:	1536
RefSeq ORF:	939
Synonyms:	HST422; TMS; TS
Locus ID:	7298



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	Thymidylate Synthase (TYMS) (NM_001071) Human Mass Spec Standard – PH304814
UniProt ID:	<u>P04818, Q53Y97</u>
Cytogenetics:	18p11.32
Summary:	Thymidylate synthase catalyzes the methylation of deoxyuridylate to deoxythymidylate using, 10-methylenetetrahydrofolate (methylene-THF) as a cofactor. This function maintains the dTMP (thymidine-5-prime monophosphate) pool critical for DNA replication and repair. The enzyme has been of interest as a target for cancer chemotherapeutic agents. It is considered to be the primary site of action for 5-fluorouracil, 5-fluoro-2-prime-deoxyuridine, and some folate analogs. Expression of this gene and that of a naturally occurring antisense transcript, mitochondrial enolase superfamily member 1 (GeneID:55556), vary inversely when cell- growth progresses from late-log to plateau phase. Polymorphisms in this gene may be associated with etiology of neoplasia, including breast cancer, and response to chemotherapy. [provided by RefSeq, Aug 2017]
Protein Families:	Druggable Genome
Protein Pathways	: Metabolic pathways, One carbon pool by folate, Pyrimidine metabolism
Product imag	es:



Coomassie blue staining of purified TYMS protein (Cat# [TP304814]). The protein was produced from HEK293T cells transfected with TYMS cDNA clone (Cat# [RC204814]) using MegaTran 2.0 (Cat# [TT210002]).

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