

OriGene Technologies, Inc.

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Product datasheet for PH300224

Malignant T cell amplified sequence 1 (MCTS1) (NM_014060) Human Mass Spec Standard

Product data:

Product Type:	Mass Spec Standards
Description:	MCTS1 MS Standard C13 and N15-labeled recombinant protein (NP_054779)
Species:	Human
Expression Host:	HEK293
Expression cDNA Clone or AA Sequence:	RC200224
Predicted MW:	20.6 kDa
Protein Sequence:	<pre>>RC200224 protein sequence Red=Cloning site Green=Tags(s)</pre>
	MFKKFDEKENVSNCIQLKTSVIKGIKNQLIEQFPGIEPWLNQIMPKKDPVKIVRCHEHIEILTVNGELLF FRQREGPFYPTLRLLHKYPFILPHQQVDKGAIKFVLSGANIMCPGLTSPGAKLYPAAVDTIVAIMAEGKQ HALCVGVMKMSAEDIEKVNKGIGIENIHYLNDGLWHMKTYK
	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 054779</u>
RefSeq Size:	9805
RefSeq ORF:	543
Synonyms:	MCT-1; MCT1
Locus ID:	28985
UniProt ID:	Q9ULC4



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	PH300224
Cytogenetics:	Xq24
Summary:	Anti-oncogene that plays a role in cell cycle regulation; decreases cell doubling time and anchorage-dependent growth; shortens the duration of G1 transit time and G1/S transition. When constitutively expressed, increases CDK4 and CDK6 kinases activity and CCND1/cyclin D1 protein level, as well as G1 cyclin/CDK complex formation. Involved in translation initiation; promotes recruitment of aminoacetyled initiator tRNA to P site of 40S ribosomes. Can promote release of deacylated tRNA and mRNA from recycled 40S subunits following ABCE1-mediated dissociation of post-termination ribosomal complexes into subunits. Plays a role as translation enhancer; recruits the density-regulated protein/DENR and binds to the cap complex of the 5'-terminus of mRNAs, subsequently altering the mRNA translation profile; up-regulates protein levels of BCL2L2, TFDP1, MRE11, CCND1 and E2F1, while mRNA levels remains constant. Hyperactivates DNA damage signaling pathway; increased gamma-irradiation-induced phosphorylation of histone H2AX, and induces damage foci formation. Increases the overall number of chromosomal abnormalities such as larger chromosomes formation and multiples chromosomal fusions when overexpressed in gamma-irradiated cells. May play a role in promoting lymphoid tumor development: lymphoid cell lines overexpressing MCTS1 exhibit increased growth rates and display increased protection against apoptosis. Involved in the process of proteasome degradation to down-regulate Tumor suppressor p53/TP53 in breast cancer cell; Positively regulates phosphorylation of MAPK1 and MAPK3. Involved in translation initiation; promotes aminoacetyled initiator tRNA to P site of 40S ribosomes. Can promote release of deacylated tRNA and mRNA from recycled 40S subunits following ABCE1-mediated dissociation of post-termination ribosomal complexes into subunits.[UniProtKB/Swiss-Prot Function]
Protein Families:	Druggable Genome

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Product images:

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Coomassie blue staining of purified MCTS1 protein (Cat# [TP300224]). The protein was produced from HEK293T cells transfected with MCTS1 cDNA clone (Cat# [RC200224]) using MegaTran 2.0 (Cat# [TT210002]).

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