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## Product datasheet for PH304322

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## MALT1 (NM_173844) Human Mass Spec Standard

## Product data:

Product Type:
Description:
Species:
Expression Host:
Expression cDNA Clone
or AA Sequence:
Predicted MW:
Protein Sequence:

## Mass Spec Standards

MALT1 MS Standard C13 and N15-labeled recombinant protein (NP_776216)
Human
HEK293
RC204322
90.9 kDa
>RC204322 representing NM_173844
Red=Cloning site Green=Tags(s)
MSLLGDPLQALPPSAAPTGPLLAPPAGATLNRLREPLLRRLSELLDQAPEGRGWRRLAELAGSRGRLRLS CLDLEQCSLKVLEPEGSPSLCLLKLMGEKGCTVTELSDFLQAMEHTEVLQLLSPPGIKITVNPESKAVLA GQFVKLCCRATGHPFVQYQWFKMNKEIPNGNTSELIFNAVHVKDAGFYVCRVNNNFTFEFSQWSQLDVCD IPESFQRSVDGVSESKLQICVEPTSQKLMPGSTLVLQCVAVGSPIPHYQWFKNELPLTHETKKLYMVPYV DLEHQGTYWCHVYNDRDSQDSKKVEIIIDELNNLGHPDNKEQTTDQPLAKDKVALLIGNMNYREHPKLKA PLVDVYELTNLLRQLDFKVVSLLDLTEYEMRNAVDEFLLLLDKGVYGLLYYAGHGYENFGNSFMVPVDAP NPYRSENCLCVQNILKLMQEKETGLNVFLLDMCRKRNDYDDTIPILDALKVTANIVFGYATCQGAEAFEI QHSGLANGIFMKFLKDRLLEDKKITVLLDEVAEDMGKCHLTKGKQALEIRSSLSEKRALTDPIQGTEYSA ESLVRNLQWAKAHELPESMCLKFDCGVQIQLGFAAEFSNVMIIYTSIVYKPPEIIMCDAYVTDFPLDLDI DPKDANKGTPEETGSYLVSKDLPKHCLYTRLSSLQKLKEHLVFTVCLSYQYSGLEDTVEDKQEVNVGKPL IAKLDMHRGLGRKTCFQTCLMSNGPYQSSAATSGGAGHYHSLQDPFHGVYHSHPGNPSNVTPADSCHCSR TPDAFISSFAHHASCHFSRSNVPVETTDEIPFSFSDRLRISEK

## TRTRPLEQKLISEEDLAANDILDYKDDDDKV

## Tag: C-Myc/DDK

## Purity:

## Concentration:

Labeling Method:
Buffer:
Storage:
Stability:
RefSeq:
$>80 \%$ as determined by SDS-PAGE and Coomassie blue staining
$>0.05 \mu \mathrm{~g} / \mu \mathrm{L}$ as determined by microplate BCA method
Labeled with [U- 13C6, 15N4]-L-Arginine and [U- 13C6, 15N2]-L-Lysine 25 mM Tris-HCl, 100 mM glycine, pH 7.3

Store at $-80^{\circ} \mathrm{C}$. Avoid repeated freeze-thaw cycles.
Stable for 3 months from receipt of products under proper storage and handling conditions.
NP 776216

| RefSeq Size: | 4996 |
| :---: | :---: |
| RefSeq ORF: | 2439 |
| Synonyms: | IMD12; MLT; MLT1; PCASP1 |
| Locus ID: | 10892 |
| UniProt ID: | Q9UDY8, A8K5S1 |
| Cytogenetics: | 18q21.32 |
| Summary: | This gene encodes a caspase-like protease that plays a role in BCL10-induced activation of NF-kappaB. The protein is a component of the CARMA1-BCL10-MALT1 (CBM) signalosome that triggers NF-kappaB signaling and lymphoctye activation following antigen-receptor stimulation. Mutations in this gene result in immunodeficiency 12 (IMD12). This gene has been found to be recurrently rearranged in chromosomal translocations with other genes in mucosa-associated lymphoid tissue lymphomas, including at $(11 ; 18)(q 21 ; q 21)$ translocation with the baculoviral IAP repeat-containing protein 3 (also known as apoptosis inhibitor 2) locus [BIRC3(API2)-MALT1], and a $\mathrm{t}(14 ; 18)(\mathrm{q} 32 ; \mathrm{q} 21)$ translocation with the immunoglobulin heavy chain locus (IGH-MALT1). Alternatively spliced transcript variants have been described for this gene. [provided by RefSeq, May 2018] |
| Protein Families: | Druggable Genome, Protease |
| Protein Pathways: | B cell receptor signaling pathway, $T$ cell receptor signaling pathway |

## Product images:



Coomassie blue staining of purified MALT1 protein (Cat\# [TP304322]). The protein was produced from HEK293T cells transfected with MALT1 cDNA clone (Cat\# [RC204322]) using MegaTran 2.0 (Cat\# [TT210002]).

