

Product datasheet for PH302638

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

SDOS (NUDT16L1) (NM_032349) Human Mass Spec Standard

Product data:

Product Type: Mass Spec Standards

Description: NUDT16L1 MS Standard C13 and N15-labeled recombinant protein (NP_115725)

Species:HumanExpression Host:HEK293

Expression cDNA Clone

RC202638

or AA Sequence:

Predicted MW:

23.3 kDa

Protein Sequence: >RC202638 protein sequence

Red=Cloning site Green=Tags(s)

MSTAAVPELKQISRVEAMRLGPGWSHSCHAMLYAANPGQLFGRIPMRFSVLMQMRFDGLLGFPGGFVDRR FWSLEDGLNRVLGLGCLRLTEADYLSSHLTEGPHRVVAHLYARQLTLEQLHAVEISAVHSRDHGLEVL GLVRVPLYTQKDRVGGFPNFLSNAFVSTAKCQLLFALKVLNMMPEEKLVEALAAATEKQKKALEKLLPAS

S

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: NP 115725

RefSeq Size: 1367 RefSeq ORF: 633

Synonyms: SDOS; TIRR

Locus ID: 84309

UniProt ID: Q9BRJ7, B2RD96





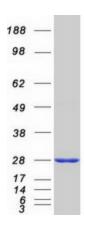
Cytogenetics:

16p13.3

Summary:

Key regulator of TP53BP1 required to stabilize TP53BP1 and regulate its recruitment to chromatin (PubMed:28241136). In absence of DNA damage, interacts with the tandem Tudor-like domain of TP53BP1, masking the region that binds histone H4 dimethylated at 'Lys-20' (H4K20me2), thereby preventing TP53BP1 recruitment to chromatin and maintaining TP53BP1 localization to the nucleus (PubMed:28241136). Following DNA damage, ATM-induced phosphorylation of TP53BP1 and subsequent recruitment of RIF1 leads to dissociate NUDT16L1/TIRR from TP53BP1, unmasking the tandem Tudor-like domain and allowing recruitment of TP53BP1 to DNA double strand breaks (DSBs) (PubMed:28241136). Binds U8 snoRNA (PubMed:18820299).[UniProtKB/Swiss-Prot Function]

Product images:



Coomassie blue staining of purified NUDT16L1 protein (Cat# [TP302638]). The protein was produced from HEK293T cells transfected with NUDT16L1 cDNA clone (Cat# [RC202638]) using MegaTran 2.0 (Cat# [TT210002]).