

Product datasheet for PH312386

C6orf150 (MB21D1) (NM_138441) Human Mass Spec Standard

Product data:

Product Type:	Mass Spec Standards
Description:	C6orf150 MS Standard C13 and N15-labeled recombinant protein (NP_612450)
Species:	Human
Expression Host:	HEK293
Expression cDNA Clone or AA Sequence:	RC212386
Predicted MW:	58.9 kDa
Protein Sequence:	>RC212386 protein sequence Red=Cloning site Green=Tags(s)

MQPWHGKAMQRASEAGATAPKASARNARGAPMDPNESPAAPAEALPKAGKFGPARKSGSRQKKSAPDTQE
RPPVVRATGARAKKAPQRAQDTQPSDATSAPGAEGLEPPAAREPALSRAGSCRQRGARCSTKPRPPPGPWD
VPSPGLPVSAPILVRRDAAPGASKLRVLEKLEKLSRDDISTAAGMVKGVDHLLRLKCDASAFRGVGLLN
TGSYYEHVKISAPNEFDVMFKLEVPRIQLEEYSNTRAYYFVKFKRNPKENHLSQFLEGEILSASKMLSKF
RKIIKKEEINDIKDQDVMKRKRGGSPAVTLLISEKISVDITLALLESKSSWPASTQEGLRIQNWLSAKVRK
QLRLKPFYLVPKHAKGNGFQEETWRLSFSHIEKEILNNHGKSKTCCENKEEKCCRKDKLKMKYLLEQL
KERFKDKKHLDFSSYHVKTAFHVCTQNPQDSQWDRKDLGLCFDNCVTFYFLQCLRTEKLENYFIPEFNL
FSSNLIDKRSKEFLTKQIEYERNNEFPVDFE

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_612450</u>
RefSeq Size:	1802
RefSeq ORF:	1566

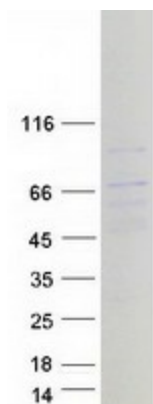


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Synonyms: C6orf150; h-cGAS; MB21D1
Locus ID: 115004
UniProt ID: [Q8N884](#)
Cytogenetics: 6q13

Summary:

Nucleotidyltransferase that catalyzes the formation of cyclic GMP-AMP (cGAMP) from ATP and GTP and plays a key role in innate immunity (PubMed:23258413, PubMed:23707061, PubMed:23722159, PubMed:24077100, PubMed:25131990, PubMed:29976794, PubMed:30799039). Catalysis involves both the formation of a 2',5' phosphodiester linkage at the GpA step and the formation of a 3',5' phosphodiester linkage at the ApG step, producing c[G(2',5')pA(3',5')p] (PubMed:28363908, PubMed:28214358). Acts as a key cytosolic DNA sensor, the presence of double-stranded DNA (dsDNA) in the cytoplasm being a danger signal that triggers the immune responses (PubMed:28363908). Binds cytosolic DNA directly, leading to activation and synthesis of cGAMP, a second messenger that binds to and activates TMEM173/STING, thereby triggering type-I interferon production (PubMed:28363908, PubMed:28314590). Preferentially recognizes and binds curved long DNAs (PubMed:30007416). In contrast to other mammals, human CGAS displays species-specific mechanisms of DNA recognition and produces less cyclic GMP-AMP (cGAMP), allowing a more fine-tuned response to pathogens (PubMed:30007416). Has antiviral activity by sensing the presence of dsDNA from DNA viruses in the cytoplasm (PubMed:28363908). Also acts as an innate immune sensor of infection by retroviruses, such as HIV-1, by detecting the presence of reverse-transcribed DNA in the cytosol (PubMed:23929945). Detection of retroviral reverse-transcribed DNA in the cytosol may be indirect and be mediated via interaction with PQBP1, which directly binds reverse-transcribed retroviral DNA (PubMed:26046437). Also detects the presence of DNA from bacteria, such as M.tuberculosis (PubMed:26048138). cGAMP can be transferred from producing cells to neighboring cells through gap junctions, leading to promote TMEM173/STING activation and convey immune response to connecting cells (PubMed:24077100). cGAMP can also be transferred between cells by virtue of packaging within viral particles contributing to IFN-induction in newly infected cells in a cGAS-independent but TMEM173/STING-dependent manner (PubMed:26229115). In addition to antiviral activity, also involved in the response to cellular stresses, such as senescence, DNA damage or genome instability (PubMed:28738408, PubMed:28759889). Acts as a regulator of cellular senescence by binding to cytosolic chromatin fragments that are present in senescent cells, leading to trigger type-I interferon production via TMEM173/STING and promote cellular senescence (By similarity). Also involved in the inflammatory response to genome instability and double-stranded DNA breaks: acts by localizing to micronuclei arising from genome instability (PubMed:28738408, PubMed:28759889). Micronuclei, which as frequently found in cancer cells, consist of chromatin surrounded by its own nuclear membrane: following breakdown of the micronuclear envelope, a process associated with chromothripsis, CGAS binds self-DNA exposed to the cytosol, leading to cGAMP synthesis and subsequent activation of TMEM173/STING and type-I interferon production (PubMed:28738408, PubMed:28759889). Acts as a suppressor of DNA repair in response to DNA damage: translocates to the nucleus following dephosphorylation at Tyr-215 and inhibits homologous recombination repair by interacting with PARP1, the CGAS-PARP1 interaction leading to impede the formation of the PARP1-TIMELESS complex (PubMed:30356214).[UniProtKB/Swiss-Prot Function]

Product images:

Coomassie blue staining of purified CGAS protein (Cat# [TP312386]). The protein was produced from HEK293T cells transfected with CGAS cDNA clone (Cat# [RC212386]) using MegaTran 2.0 (Cat# [TT210002]).