

Product datasheet for **TP311483**

COP1 (RFWD2) (NM_001001740) Human Recombinant Protein

Product data:

Product Type: Recombinant Proteins
Description: Recombinant protein of human ring finger and WD repeat domain 2 (RFWD2), transcript variant 2, 20 µg
Species: Human
Expression Host: HEK293T
Expression cDNA Clone or AA Sequence: >RC211483 representing NM_001001740
Red=Cloning site **Green**=Tags(s)

MSGSRQAGSGSAGTSPGSSAASSVTSASSSLSSSPSPPSVAVSAAALVSGGVAQAAGSGGLGGPVRPVLV
APAVSGSGGGAVSTGLSRHSCAARPSAGVGGSSSLGSGSRKRPLLAPLCNGLINSYEDKSNDFVCPICF
DMIEEAYMTKCGHSFCYKCIHQSLDNNRCPKCNVVDNIDHLYPNFLVNELILKQKQRFEEKRFLDHS
NGHRWQIFQDWLGTQDNLDLANVNLMLLELLVQKKKQLEAESHAALQILMEFLKVARRNKREEMSGLYS
PVSEDSTVPQFEAPSPSHSSIIDSTEYSQPPGFSGSSQTKKQPWYNSTLASRRKRLTAHFEDLEQCYFST
RMSRISDDSRASQLDEFQECLSKFTRYNSVRPLATLSYASDLYNGSSIVSSIEFDRDCDYFAIAGVTKK
IKVYEYDTVIQDAVDIHYPENEMTCNSKISCISWSSYHKNLLASSDYEGTVILWDGFTGQRSKVYQEHEK
RCWSVDFNLMDPKLLASGSDDAKVKLWSTNLDNSVASIEAKANVCCVKFSPSSRYHLAFGCADHCVHYD
LRNTKQPIMVFKGHRKAVSYAKFVSGEEIVSASTDSQLKLWNVGKPYCLRSFKGHINEKNFVGLASNGDY
IACGSENNSLYLYKGLSKTLLTFKFDTVKSVLDKDRKEDDTNEFVSAVCWRALPDGESNVLIAANSQGT
IKVLELV

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

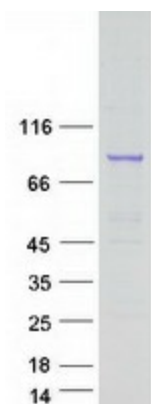
Tag: C-Myc/DDK
Predicted MW: 77.5 kDa
Concentration: >0.05 µg/µL as determined by microplate BCA method
Purity: > 80% as determined by SDS-PAGE and Coomassie blue staining
Buffer: 25 mM Tris-HCl, 100 mM glycine, pH 7.3, 10% glycerol
Preparation: Recombinant protein was captured through anti-DDK affinity column followed by conventional chromatography steps.
Note: For testing in cell culture applications, please filter before use. Note that you may experience some loss of protein during the filtration process.



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Storage:	Store at -80°C.
Stability:	Stable for 12 months from the date of receipt of the product under proper storage and handling conditions. Avoid repeated freeze-thaw cycles.
RefSeq:	NP_001001740
Locus ID:	64326
UniProt ID:	Q8NHY2
RefSeq Size:	2729
Cytogenetics:	1q25.1-q25.2
RefSeq ORF:	2121
Synonyms:	CFAP78; FAP78; RFWD2; RNF200
Summary:	<p>E3 ubiquitin-protein ligase that mediates ubiquitination and subsequent proteasomal degradation of target proteins. E3 ubiquitin ligases accept ubiquitin from an E2 ubiquitin-conjugating enzyme in the form of a thioester and then directly transfers the ubiquitin to targeted substrates. Involved in JUN ubiquitination and degradation. Directly involved in p53 (TP53) ubiquitination and degradation, thereby abolishing p53-dependent transcription and apoptosis. Ubiquitinates p53 independently of MDM2 or RCHY1. Probably mediates E3 ubiquitin ligase activity by functioning as the essential RING domain subunit of larger E3 complexes. In contrast, it does not constitute the catalytic RING subunit in the DCX DET1-COP1 complex that negatively regulates JUN, the ubiquitin ligase activity being mediated by RBX1. Involved in 14-3-3 protein sigma/SFN ubiquitination and proteasomal degradation, leading to AKT activation and promotion of cell survival. Ubiquitinates MTA1 leading to its proteasomal degradation. Upon binding to TRIB1, ubiquitinates CEBPA, which lacks a canonical COP1-binding motif (Probable). [UniProtKB/Swiss-Prot Function]</p>
Protein Pathways:	p53 signaling pathway, Ubiquitin mediated proteolysis

Product images:



Coomassie blue staining of purified COP1 protein (Cat# TP311483). The protein was produced from HEK293T cells transfected with COP1 cDNA clone (Cat# [RC211483]) using MegaTran 2.0 (Cat# [TT210002]).