

Product datasheet for **RG238538**

COP1 (RFWD2) (NM_001286644) Human Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	COP1 (RFWD2) (NM_001286644) Human Tagged ORF Clone
Tag:	TurboGFP
Symbol:	COP1
Synonyms:	CFAP78; FAP78; RFWD2; RNF200
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-AC-GFP (PS100010)
E. coli Selection:	Ampicillin (100 ug/mL)



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ORF Nucleotide Sequence:

>RG238538 representing NM_001286644.
 Blue=ORF Red=Cloning site Green=Tag(s)

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GCTCGTTAGTGAACCGTCAGAATTTTGTAAACGACTCACTATAGGGCGGCCGGGAATTCGTGACTG
GATCCGGTACCGAGGAGATCTGCCGCCCGGATCGCC
ATGTTGTGGACAATATTGACCATCTGTATCCTAATTTCTTGGGAATCACATGCAGCCCACTACAGATT
CTTATGGAAATTCCTCAAGTTGCAAGAAGAAATAGAGAGAGCAACTGGAACAGATCCAGAAGGAGCTA
AGTGTGTTTGAAGAGGATATTAAGAGAGTGGAAGAAATGAGTGGCTTATACTCTCCTGTCAGTGAGGAT
AGCACAGTGCCTCAATTTGAAGCTCCTTCTCCATCACACAGTAGTATTATTGATTCCACAGAATACAGC
CAACCTCCAGTTTCAGTGGCAGTTCTCAGACAAAGAAACAGCCTTGGTATAATAGCACGTTAGCATCA
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TATAATTCAGTACGACCTTAGCCACATTGTCATATGCTAGTGTCTCTATAATGGTTCCAGTATAGTC
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ATAGCTTGTGGAAGTAAAATAACTCTCTCTACCTGTACTATAAAGGACTTTCTAAGACTTTGCTAACT
TTTAAGTTTGATACAGTCAAAGTGTCTCGACAAAGACCGAAAAGAAGATGATACAAATGAATTTGTT
AGTGCTGTGTGCTGGAGGGCACTACCAGATGGGGAGTCCAATGTGCTGATTGCTGCTAACAGTCAGGGT
ACAATTAAGGTGCTAGAATTGGTA
ACGCGTACGCGGCCGCTCGAG - GFP Tag - GTTTAAAC
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Protein Sequence:

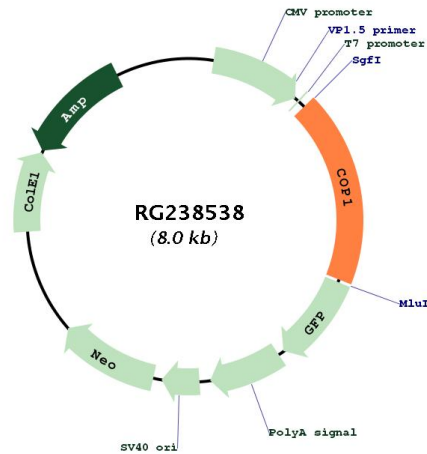
>Peptide sequence encoded by RG238538
 Blue=ORF Red=Cloning site Green=Tag(s)

```
MLWTIL TICILISWESHAAQLQILMEFLKVARRNKREQLQIQKELSVLEEDIKRVEEMSGLYSPVSED
STVPQFEAPSPSHSSIIDSTEYSQPPGFSGSSQTKKQPWYNSTLASRRKRLTAHFEDLEQCYFSTRMSR
ISDDSRASQLDEFQECLSKFTRYNSVRPLATLSYASDL YNGSSIVSSIEFDRDCDYFAIAGVTKKIKV
YEYDVTIQDAVDIHYPENEMTCNSKISCSWSSYHKNL LASSDYEGTVILWDGFTGQRSKVYQEHEKRC
WSVDFNLM DPKLLASGSDDAKVKLWSTNLDNSVASIEAKANVCCVKFSPSSRYHLAFGCADHCVHYDDL
RNTKQPI MVFKGHRKAVSYAKFVSGEEIVSASTDSQLKLWNVGKPYCLRSFKGHINEKNFVGLASNGDY
IACGSENNSLYL YKGLSKTLLTFKFDTVKSVL DKDRKEDDTNEFVSAVCWRALPDGESNVLIAANSQG
TIKVLELV
TRTRPLEMESDESGLPAMEIECRITGTLNGVEFELVGGGEGTPEQGRMTNKMSTKGALTFSPYLLSHV
MGYGFYHFGTYPSGYENPFLHAINNGGYNTRIEKYEDGGVLHVSFSYRYEAGRVIGDFKVMGTGFPED
SVIFTDKIIRSNATVEHLHPMGDNDLDGSFTRTFSLRDGGYSSVVD SHMHFKSAIHPSILQNGGPMFA
FRRVEEDHSNTELGIVEYQHAFKTPDADAGEERV
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Restriction Sites:

Sgfl-Mlul

Plasmid Map:



ACCN: NM_001286644

ORF Size: 1473 bp

OTI Disclaimer: The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. [More info](#)

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

RefSeq: [NM_001286644.1](#), [NP_001273573.1](#)

RefSeq Size:	2609 bp
RefSeq ORF:	1476 bp
Locus ID:	64326
UniProt ID:	Q8NHY2
Cytogenetics:	1q25.1-q25.2
Protein Pathways:	p53 signaling pathway, Ubiquitin mediated proteolysis
MW:	56.1 kDa
Gene 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>