产品名称: COP1 (phospho Ser387) Rabbit Polyclonal

Antibody

产品货号: APRab04487



产品概述 (Summary)

产品名称 (Production Name) COP1 (phospho Ser387) Rabbit Polyclonal Antibody

描述 (Description) Rabbit polyclonal Antibody

宿主 (Host)Rabbit应用 (Application)WB,ELISA种属反应性 (Reactivity)Human,Mouse

产品性能 (Performance)

偶联物 (Conjugation) Unconjugated 修饰 (Modification) Phosphorylated

同种型 (Isotype) IgG

克隆 (Clonality) Polyclonal 形式 (Form) Liquid

Store at 4°C short term. Aliquot and store at -20°C long term. Avoid 存放说明 (Storage)

freeze/thaw cycles.

Liquid in PBS containing 50% glycerol, 0.5% protective protein and 0.02% 储存溶液 (Buffer)

New type preservative N.

纯化方式 (Purification) Affinity purification

免疫原信息 (Immunogen)

基因名 (Gene Name) RFWD2

RFWD2; COP1; RNF200; E3 ubiquitin-protein ligase RFWD2; Constitutive

别名 (Alternative Names) photomorphogenesis protein 1 homolog; hCOP1; RING finger and WD repeat

domain protein 2; RING finger protein 200

基因 ID (Gene ID) 64326.0

Q8NHY2.The antiserum was produced against synthesized peptide derived

蛋白ID (SwissProt ID) from human RFWD2 around the phosphorylation site of Ser387. AA

range:353-402

产品应用 (Application)

Web:https://www.enkilife.cn E-mail:order@enkilife.cn (销售) tech@enkilife.cn (技支持) Tel:027-87002838

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稀释比 (Dilution Ratio) WB 1:500-1:2000,ELISA 1:5000-1:10000

蛋白分子量 (Molecular Weight) 100kDa

研究背景 (Background)

domain: The RING finger domain, in addition to its role in ubiquitination, functions as a structural scaffold to bring two clusters of positive-charged residues within spatial proximity to mimic a bipartite nuclear localization signal (NLS), function: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.,induction:By p53/TP53, pathway: Protein modification; protein ubiquitination, similarity: Belongs to the COP1 family, similarity: Contains 1 RING-type zinc finger., similarity: Contains 7 WD repeats., subcellular location: In the nucleus, it forms nuclear speckles., subunit: Homodimer. Homodimerization is mediated by the coiled coil domain. Component of the DCX DET1-COP1 ubiquitin ligase complex at least composed of RBX1, DET1, DDB1, CUL4A and COP1. Isoform 2 does not interact with CUL4A but still binds to RBX1, suggesting that the interaction may be mediated by another culllin protein. Isoform 1 and isoform 2 interact with CUL5 but not with CUL1, CUL2 not CUL3. Interacts with bZIP transcription factors JUN, JUNB and JUND but not with FOS, ATF2 nor XBP1. Interacts with p53 (TP53), tissue specificity: Ubiquitously expressed at low level. Expressed at higher level in testis, placenta, skeletal muscle and heart, domain: The RING finger domain, in addition to its role in ubiquitination, functions as a structural scaffold to bring two clusters of positive-charged residues within spatial proximity to mimic a bipartite nuclear localization signal (NLS), function: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.,induction:By p53/TP53.,pathway:Protein modification; protein ubiquitination..similarity:Belongs to the COP1 family..similarity:Contains 1 RING-type zinc finger..similarity:Contains 7 WD repeats., subcellular location: In the nucleus, it forms nuclear speckles., subunit: Homodimer. Homodimerization is mediated by the coiled coil domain. Component of the DCX DET1-COP1 ubiquitin ligase complex at least composed of RBX1, DET1, DDB1, CUL4A and COP1. Isoform 2 does not interact with CUL4A but still binds to RBX1, suggesting that the interaction may be mediated by another culllin protein. Isoform 1 and isoform 2 interact with CUL5 but not with CUL1, CUL2 not CUL3.

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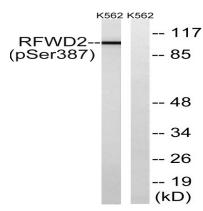


Interacts with bZIP transcription factors JUN, JUNB and JUND but not with FOS, ATF2 nor XBP1. Interacts with p53 (TP53).,tissue specificity:Ubiquitously expressed at low level. Expressed at higher level in testis, placenta, skeletal muscle and heart.,

研究领域 (Research Area)

p53;Ubiquitin mediated proteolysis;

图片 (Image Data)



Western blot analysis of lysates from K562 cells treated with UV 15 ', using RFWD2 (Phospho-Ser387) Antibody. The lane on the right is blocked with the phospho peptide.

注意事项 (Note)

For research use only.

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