

#6895 Store at -20°C

COPS5 Antibody



Orders ■ 877-616-CELL (2355)
orders@cellsignal.com
Support ■ 877-678-TECH (8324)
info@cellsignal.com
Web ■ www.cellsignal.com

rev. 02/16/16

For Research Use Only. Not For Use In Diagnostic Procedures.

Applications W, IP, IF-IC Endogenous	Species Cross-Reactivity* H, M, R, Mk, (X, Z, B, Dg, Pg, Hr)	Molecular Wt. 37 kDa	Source Rabbit**
--	---	-------------------------	--------------------

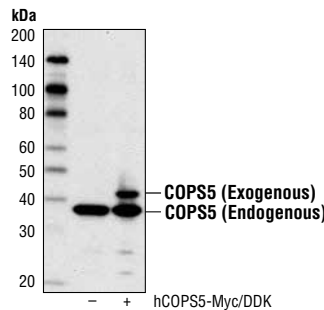
Background: The COP9 Signalosome (CSN) is a ubiquitously expressed multiprotein complex that is involved in a vast array of cellular and developmental processes, which is thought to be attributed to its control over the ubiquitin-proteasome pathway. Typically, the CSN is composed of eight highly conserved subunits (CSN1-CSN8), each of which is homologous to one of the eight subunits that form the lid of the 26S proteasome particle, suggesting that these complexes have a common evolutionary ancestor (1). CSN was first identified in *Arabidopsis thaliana* mutants with a light-grown seedling phenotype when grown in the dark (2-4). The subsequent cloning of the constitutive morphogenesis 9 (cop9) mutant from *Arabidopsis thaliana* was soon followed by the biochemical purification of the COP9-containing multiprotein complex (4). It is now widely accepted that the CSN directly interacts with cullin-RING ligase (CRL) families of ubiquitin E3 complexes, and that CSN is required for their proper function (5). In addition, CSN may also regulate protein homeostasis through its association with protein kinases and deubiquitylating enzymes. Collectively, these activities position the CSN as a pivotal regulator of the DNA-damage response, cell-cycle control, and gene expression (1).

COPS5/CSN5/Jab1 (c-Jun activation domain-binding protein-1) was originally identified as a transcriptional coactivator of c-Jun and subsequently discovered to be a fifth component and integral part of the CSN (6). As the catalytic center of the CSN, COPS5 is able to integrate multiple functions of the CSN complex such as cell-cycle control, transcription, and DNA-damage response by regulating the activity of CRLs through deneddylation of cullins (7). Indeed, COPS5 harbors a Mpr1-Pad1-N-terminal (MPN) domain with an embedded Jab1/CSN5 MPN domain metalloenzyme (JAMM) motif that is essential for the CSN isopeptidase activity responsible for deneddylation of CRLs. COPS5 is an evolutionarily conserved 38 kDa protein in humans, mice, fission yeast, and plants, which suggests that it is critical to cell survival and proliferation. A role for COPS5 as a positive regulator of cellular proliferation is supported by evidence that it functionally inactivates several key tumor suppressors such as p53, RUNX3, Smad4, and p27^{Kip1} through altered subcellular localization, degradation, and deneddylation (8-12). These findings are underscored by the observation that COPS5 overexpression has been identified in a number of different tumor types and has been

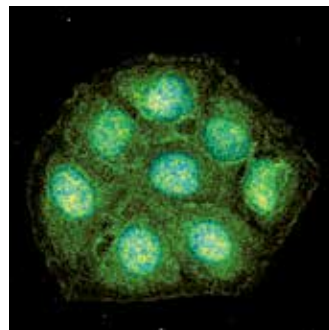
implicated in the initiation and progression of several types of cancer (13). Moreover, COPS5-deficient mice display an embryonically lethal phenotype highlighted by elevated expression of COPS5 targets such as p53 and p27 (14,15).

Specificity/Sensitivity: COPS5 Antibody recognizes endogenous levels of total COPS5 protein. Based upon sequence homology, this antibody is not predicted to cross-react with PSMD14/POH1.

Source/Purification: Polyclonal antibodies are produced by immunizing animals with a synthetic peptide corresponding to residues near the amino-terminus of human COPS5 protein. Antibodies are purified by protein A and peptide affinity chromatography.



Western blot analysis of extracts from 293T cells, either mock-transfected (-) or transfected with a Myc/DDK-tagged human COPS5 cDNA expression construct (+), using COPS5 Antibody.



Confocal immunofluorescent analysis of U-2 OS cells using COPS5 Antibody (green). Blue pseudocolor = DRAQ5[®] #4084 (fluorescent DNA dye).

Entrez-Gene ID #10987
Swiss-Prot Acc. #Q92905

Storage: Supplied in 10 mM sodium HEPES (pH 7.5), 150 mM NaCl, 100 µg/ml BSA and 50% glycerol. Store at -20°C. Do not aliquot the antibody.

*Species cross-reactivity is determined by western blot.

**Anti-rabbit secondary antibodies must be used to detect this antibody.

Recommended Antibody Dilutions:

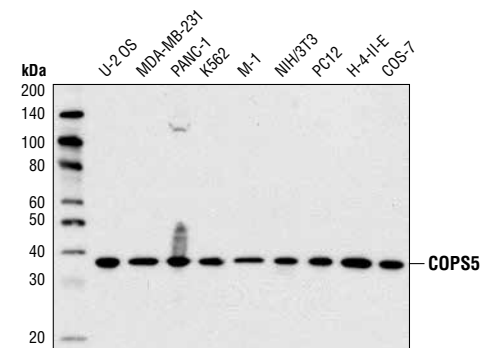
Western blotting	1:1000
Immunoprecipitation	1:50
Immunofluorescence (IF-IC)	1:50

For application specific protocols please see the web page for this product at www.cellsignal.com.

Please visit www.cellsignal.com for a complete listing of recommended companion products.

DRAQ5[®] is a registered trademark of Biostatus Limited.

IMPORTANT: For western blots, incubate membrane with diluted antibody in 5% w/v BSA, 1X TBS, 0.1% Tween-20 at 4°C with gentle shaking, overnight.



Western blot analysis of extracts from various cell lines using COPS5 Antibody.

© 2011 Cell Signaling Technology, Inc. Cell Signaling Technology[®] is a trademark of Cell Signaling Technology, Inc.

Applications Key: W—Western IP—Immunoprecipitation IHC—Immunohistochemistry ChIP—Chromatin Immunoprecipitation IF—Immunofluorescence F—Flow cytometry E-P—ELISA-Peptide
Species Cross-Reactivity Key: H—human M—mouse R—rat Hm—hamster Mk—monkey Mi—mink C—chicken Dm—D. melanogaster X—Xenopus Z—zebrafish B—bovine
Dg—dog Pg—pig Sc—S. cerevisiae Ce—C. elegans Hr—horse AI—all species expected Species enclosed in parentheses are predicted to react based on 100% homology.

Background References:

- (1) Wei, N. and Deng, X.W. (2003) *Annu Rev Cell Dev Biol* 19, 261-86.
- (2) Kwok, S.F. et al. (1996) *Plant Physiol* 110, 731-42.
- (3) Wei, N. et al. (1994) *Cell* 78, 117-24.
- (4) Chamovitz, D.A. et al. (1996) *Cell* 86, 115-21.
- (5) Cope, G.A. and Deshaies, R.J. (2003) *Cell* 114, 663-71.
- (6) Claret, F.X. et al. (1996) *Nature* 383, 453-7.
- (7) Wei, N. et al. (2008) *Trends Biochem Sci* 33, 592-600.
- (8) Bech-Otschir, D. et al. (2001) *EMBO J* 20, 1630-9.
- (9) Oh, W. et al. (2006) *J Biol Chem* 281, 17457-65.
- (10) Wan, M. et al. (2002) *EMBO Rep* 3, 171-6.
- (11) Tomoda, K. et al. (2002) *J Biol Chem* 277, 2302-10.
- (12) Kim, J.H. et al. (2009) *J Cell Biochem* 107, 557-65.
- (13) Shackleford, T.J. and Claret, F.X. (2010) *Cell Div* 5, 26.
- (14) Tian, L. et al. (2010) *Oncogene* 29, 6125-37.
- (15) Tomoda, K. et al. (2004) *J Biol Chem* 279, 43013-8.