Phospho-SQSTM1/p62 (Ser349) Antibody



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For Research Use Only. Not for Use in Diagnostic Procedures.

Applications: WB	Reactivity: H M	Sensitivity: Endogenous	MW (kDa): 62	Source: Rabbit	UniProt ID: #Q13501	Entrez-Gene Id 8878	
Product Usage Information	Ар	Application			Dilution		
	We	Western Blotting			1:1000		
Storage		pplied in 10 mM sodi C. Do not aliquot the	\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	5), 150 mM NaCl, 10	00 μg/ml BSA and 50% ç	ylycerol. Store at –	
Specificity / Sens		Phospho-SQSTM1/p62 (Ser349) Antibody recognizes endogenous levels of SQSTM1/p62 protein only when phosphorylated at Ser349.					
Species predicted to react based on 100%							

Source / Purification

sequence homology:

Polyclonal antibodies are produced by immunizing animals with a synthetic phosphopeptide corresponding to residues surrounding Ser349 of human SQSTM1/p62 protein. Antibodies are purified by protein A and peptide affinity chromatography.

Background

Sequestosome 1 (SQSTM1, p62) is a ubiquitin binding protein involved in cell signaling, oxidative stress, and autophagy (1-4). It was first identified as a protein that binds to the SH2 domain of p56Lck (5) and independently found to interact with PKCζ (6,7). SQSTM1 was subsequently found to interact with ubiquitin, providing a scaffold for several signaling proteins and triggering degradation of proteins through the proteasome or lysosome (8). Interaction between SQSTM1 and TRAF6 leads to the K63-linked polyubiquitination of TRAF6 and subsequent activation of the NF-κB pathway (9). Protein aggregates formed by SQSTM1 can be degraded by the autophagosome (4,10,11). SQSTM1 binds autophagosomal membrane protein LC3/Atg8, bringing SQSTM1-containing protein aggregates to the autophagosome (12). Lysosomal degradation of autophagosomes leads to a decrease in SQSTM1 levels during autophagy; conversely, autophagy inhibitors stabilize SQSTM1 levels. Studies have demonstrated a link between SQSTM1 and oxidative stress. SQSTM1 interacts with KEAP1, which is a cytoplasmic inhibitor of NRF2, a key transcription factor involved in cellular responses to oxidative stress (3). Thus, accumulation of SQSTM1 can lead to an increase in NRF2 activity.

Phosphorylation of SQSTM1 at Ser349 (Ser351 in mouse) during oxidative stress increases its binding to KEAP1, thereby increasing NRF2 activity (13).

Background References

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- 5. Joung, I. et al. (1996) Proc Natl Acad Sci USA 93, 5991-5.
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- 7. Puls, A. et al. (1997) Proc Natl Acad Sci USA 94, 6191-6.
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- 13. Ichimura, Y. et al. (2013) Mol Cell 51, 618-31.

Species Reactivity

Species reactivity is determined by testing in at least one approved application (e.g., western blot).

3/23/24. 11:41 AM Phospho-SOSTM1/p62 (Ser349) Antibody (#95697) Datasheet Without Images Cell Signaling Technology

Western Blot Buffer

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

Applications Key

Cross-Reactivity Key

WB: Western Blotting

H: human M: mouse R: rat Hm: hamster Mk: monkey Vir: virus Mi: mink C: chicken Dm: D. melanogaster

X: Xenopus Z: zebrafish B: bovine Dq: dog Pq: pig Sc: S. cerevisiae Ce: C. elegans Hr: horse

GP: Guinea Pig Rab: rabbit All: all species expected

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