Store at -20C

SQSTM1/p62 (D10E10) Rabbit



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Applications: Reactivity: Sensitivity: MW (kDa): Source/Isotype: **UniProt ID:** Entrez-Gene Id: IP, IF-IC Н Endogenous 62 Rabbit IgG #Q13501 8878

Product Usage Application Dilution Information 1:50 Immunoprecipitation

Immunofluorescence (Immunocytochemistry) 1:100 - 1:400

Supplied in 10 mM sodium HEPES (pH 7.5), 150 mM NaCl, 100 µg/ml BSA, 50% glycerol and less than **Storage**

0.02% sodium azide. Store at -20°C. Do not aliquot the antibody.

SOSTM1/p62 (D10E10) Rabbit mAb is recommended to detect endogenous levels of total SOSTM1/p62 Specificity / Sensitivity

protein by immunofluorescence. Products SQSTM1/p62 (D5E2) Rabbit mAb #8025 and SQSTM1/p62 Antibody #5114 are preferred for western blot.

Source / Purification Monoclonal antibody is produced by immunizing animals with a synthetic peptide corresponding to

residues near the carboxy terminus of human SQSTM1/p62 protein.

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 PKC (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-kB pathway (9). Protein aggregates formed by SOSTM1 can be degraded by the autophagosome (4,10,11). SOSTM1 binds autophagosomal membrane protein LC3/Atg8, bringing SQSTM1-containing protein aggregates to the autophagosome (12). Lysosomal degradation of autophagosomes leads to a decrease in SOSTM1 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.

1. Kirkin, V. et al. (2009) Mol Cell 34, 259-69. **Background References**

2. Seibenhener, M.L. et al. (2007) FEBS Lett 581, 175-9.

3. Komatsu, M. et al. (2010) Nat Cell Biol 12, 213-23.

4. Bjørkøy, G. et al. (2006) Autophagy 2, 138-9.

5. Joung, I. et al. (1996) Proc Natl Acad Sci USA 93, 5991-5.

6. Sanchez, P. et al. (1998) Mol Cell Biol 18, 3069-80.

7. Puls. A. et al. (1997) Proc Natl Acad Sci USA 94, 6191-6.

8. Vadlamudi, R.K. et al. (1996) J Biol Chem 271, 20235-7.

9. Wooten, M.W. et al. (2005) J Biol Chem 280, 35625-9.

10. Bjørkøy, G. et al. (2005) J Cell Biol 171, 603-14.

11. Komatsu, M. et al. (2007) Cell 131, 1149-63.

12. Pankiv, S. et al. (2007) J Biol Chem 282, 24131-45.

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

IP: Immunoprecipitation IF-IC: Immunofluorescence (Immunocytochemistry) **Applications Key**

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

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

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

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Limited Uses

SQSTM1/p62 (D10E10) Rabbit mAb (#7695) Datasheet Without Images Cell Signaling Technology

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