

# Anti-p62 Antibody

Rabbit polyclonal antibody to p62 Catalog # AP61633

### **Product Information**

Application	WB, IHC-P
Primary Accession	<u>Q13501</u>
Other Accession	<u>Q64337</u>
Reactivity	Human, Mouse, Rat
Host	Rabbit
Clonality	Polyclonal
Calculated MW	47687

## **Additional Information**

Gene ID	8878
Other Names	ORCA; OSIL; Sequestosome-1; EBI3-associated protein of 60 kDa; EBIAP; p60; Phosphotyrosine-independent ligand for the Lck SH2 domain of 62 kDa; Ubiquitin-binding protein p62
Target/Specificity	Recombinant full length protein of human p62
Dilution	WB~~1:1000 IHC-P~~N/A
Format	Liquid in 0.42% Potassium phosphate, 0.87% Sodium chloride, pH 7.3, 30% glycerol, and 0.09% (W/V) sodium azide.
Storage	Store at -20 °C.Stable for 12 months from date of receipt

#### **Protein Information**

Name	SQSTM1 {ECO:0000303 PubMed:16286508,
	ECO:0000312 HGNC:HGNC:11280}
Function	Molecular adapter required for selective macroautophagy (aggrephagy) by acting as a bridge between polyubiquitinated proteins and autophagosomes
	(PubMed: <u>15340068</u> , PubMed: <u>15953362</u> , PubMed: <u>16286508</u> ,
	PubMed: <u>17580304</u> , PubMed: <u>20168092</u> , PubMed: <u>22017874</u> ,
	PubMed: <u>22622177</u> , PubMed: <u>24128730</u> , PubMed: <u>28404643</u> ,
	PubMed: <u>29343546</u> , PubMed: <u>29507397</u> , PubMed: <u>31857589</u> ,
	PubMed: <u>33509017</u> , PubMed: <u>34471133</u> , PubMed: <u>34893540</u> ,
	PubMed: <u>35831301</u> , PubMed: <u>37306101</u> , PubMed: <u>37802024</u> ). Promotes the
	recruitment of ubiquitinated cargo proteins to autophagosomes via multiple
	domains that bridge proteins and organelles in different steps
	(PubMed: <u>16286508</u> , PubMed: <u>20168092</u> , PubMed: <u>22622177</u> ,
	PubMed: <u>24128730</u> , PubMed: <u>28404643</u> , PubMed: <u>29343546</u> ,

PubMed:29507397, PubMed:34893540, PubMed:37802024). SQSTM1 first mediates the assembly and removal of ubiquitinated proteins by undergoing liquid-liquid phase separation upon binding to ubiquitinated proteins via its UBA domain, leading to the formation of insoluble cytoplasmic inclusions, known as p62 bodies (PubMed:15911346, PubMed:20168092, PubMed:22017874, PubMed:24128730, PubMed:29343546, PubMed:29507397, PubMed:31857589, PubMed:37802024). SQSTM1 then interacts with ATG8 family proteins on autophagosomes via its LIR motif, leading to p62 body recruitment to autophagosomes, followed by autophagic clearance of ubiquitinated proteins (PubMed: 16286508, PubMed: 17580304, PubMed:20168092, PubMed:22622177, PubMed:24128730, PubMed:<u>28404643</u>, PubMed:<u>37802024</u>). SQSTM1 is itself degraded along with its ubiquitinated cargos (PubMed: 16286508, PubMed: 17580304, PubMed: 37802024). Also required to recruit ubiquitinated proteins to PML bodies in the nucleus (PubMed:20168092). Also involved in autophagy of peroxisomes (pexophagy) in response to reactive oxygen species (ROS) by acting as a bridge between ubiquitinated PEX5 receptor and autophagosomes (PubMed:26344566). Acts as an activator of the NFE2L2/NRF2 pathway via interaction with KEAP1: interaction inactivates the BCR(KEAP1) complex by sequestering the complex in inclusion bodies, promoting nuclear accumulation of NFE2L2/NRF2 and subsequent expression of cytoprotective genes (PubMed:20452972, PubMed:28380357, PubMed:33393215, PubMed:37306101). Promotes relocalization of 'Lys-63'-linked ubiquitinated STING1 to autophagosomes (PubMed:<u>29496741</u>). Involved in endosome organization by retaining vesicles in the perinuclear cloud: following ubiquitination by RNF26, attracts specific vesicle-associated adapters, forming a molecular bridge that restrains cognate vesicles in the perinuclear region and organizes the endosomal pathway for efficient cargo transport (PubMed:27368102, PubMed:33472082). Sequesters tensin TNS2 into cytoplasmic puncta, promoting TNS2 ubiquitination and proteasomal degradation (PubMed:<u>25101860</u>). May regulate the activation of NFKB1 by TNF-alpha, nerve growth factor (NGF) and interleukin-1 (PubMed: 10356400, PubMed:10747026, PubMed:11244088, PubMed:12471037, PubMed:16079148, PubMed:19931284). May play a role in titin/TTN downstream signaling in muscle cells (PubMed: 15802564). Adapter that mediates the interaction between TRAF6 and CYLD (By similarity). Cytoplasmic vesicle, autophagosome. Preautophagosomal structure. Cytoplasm, cytosol. Nucleus, PML body. Late endosome. Lysosome. Nucleus Endoplasmic reticulum. Cytoplasm, myofibril, sarcomere

{ECO:0000250|UniProtKB:008623}. Note=In cardiac muscle, localizes to the sarcomeric band (By similarity). Localizes to cytoplasmic membraneless inclusion bodies, known as p62 bodies, containing polyubiquitinated protein aggregates (PubMed:11786419, PubMed:20357094, PubMed:22017874, PubMed:29343546, PubMed:29507397, PubMed:31857589, PubMed:37306101, PubMed:37802024). In neurodegenerative diseases, detected in Lewy bodies in Parkinson disease, neurofibrillary tangles in Alzheimer disease, and HTT aggregates in Huntington disease (PubMed:15158159). In protein aggregate diseases of the liver, found in large amounts in Mallory bodies of alcoholic and nonalcoholic steatohepatitis, hyaline bodies in hepatocellular carcinoma, and in SERPINA1 aggregates (PubMed:11981755) Enriched in Rosenthal fibers of pilocytic astrocytoma (PubMed:11786419). In the cytoplasm, observed in both membrane-free ubiquitin-containing protein aggregates (sequestosomes) and membranesurrounded autophagosomes (PubMed:15953362, PubMed:17580304) Colocalizes with TRIM13 in the perinuclear endoplasmic reticulum (PubMed:22178386). Co-localizes with TRIM5 in cytoplasmic bodies (PubMed:20357094). When nuclear export is blocked by treatment with leptomycin B, accumulates in PML bodies (PubMed:20168092) {ECO:0000250|UniProtKB:008623, ECO:0000269|PubMed:11786419,

**Cellular Location** 

ECO:000269 | PubMed:11981755, ECO:000269 | PubMed:15158159, ECO:000269 | PubMed:15953362, ECO:0000269 | PubMed:17580304, ECO:0000269 | PubMed:20168092, ECO:0000269 | PubMed:20357094, ECO:0000269 | PubMed:22017874, ECO:0000269 | PubMed:22178386, ECO:0000269 | PubMed:29343546, ECO:0000269 | PubMed:29507397, ECO:0000269 | PubMed:31857589, ECO:0000269 | PubMed:37306101, ECO:0000269 | PubMed:37802024}

**Tissue Location**Ubiquitously expressed.

## Background

Recombinant full length protein of human p62

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