

Terje Johansen

List of Publications by Year in descending order

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Version: 2024-02-01

156
papers

41,095
citations

9775

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h-index

7511

151
g-index

164
all docs

164
docs citations

164
times ranked

42598
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Guidelines for the use and interpretation of assays for monitoring autophagy (3rd edition). <i>Autophagy</i> , 2016, 12, 1-222. | 4.3 | 4,701 |
| 2 | p62/SQSTM1 Binds Directly to Atg8/LC3 to Facilitate Degradation of Ubiquitinated Protein Aggregates by Autophagy. <i>Journal of Biological Chemistry</i> , 2007, 282, 24131-24145. | 1.6 | 3,766 |
| 3 | Guidelines for the use and interpretation of assays for monitoring autophagy. <i>Autophagy</i> , 2012, 8, 445-544. | 4.3 | 3,122 |
| 4 | p62/SQSTM1 forms protein aggregates degraded by autophagy and has a protective effect on huntingtin-induced cell death. <i>Journal of Cell Biology</i> , 2005, 171, 603-614. | 2.3 | 2,854 |
| 5 | Guidelines for the use and interpretation of assays for monitoring autophagy in higher eukaryotes. <i>Autophagy</i> , 2008, 4, 151-175. | 4.3 | 2,064 |
| 6 | Selective autophagy mediated by autophagic adapter proteins. <i>Autophagy</i> , 2011, 7, 279-296. | 4.3 | 1,512 |
| 7 | Molecular definitions of autophagy and related processes. <i>EMBO Journal</i> , 2017, 36, 1811-1836. | 3.5 | 1,230 |
| 8 | p62/SQSTM1 Is a Target Gene for Transcription Factor NRF2 and Creates a Positive Feedback Loop by Inducing Antioxidant Response Element-driven Gene Transcription. <i>Journal of Biological Chemistry</i> , 2010, 285, 22576-22591. | 1.6 | 1,158 |
| 9 | A Role for NBR1 in Autophagosomal Degradation of Ubiquitinated Substrates. <i>Molecular Cell</i> , 2009, 33, 505-516. | 4.5 | 974 |
| 10 | Chapter 12 Monitoring Autophagic Degradation of p62/SQSTM1. <i>Methods in Enzymology</i> , 2009, 452, 181-197. | 0.4 | 936 |
| 11 | Interactions between Autophagy Receptors and Ubiquitin-like Proteins Form the Molecular Basis for Selective Autophagy. <i>Molecular Cell</i> , 2014, 53, 167-178. | 4.5 | 849 |
| 12 | The LIR motif " crucial for selective autophagy. <i>Journal of Cell Science</i> , 2013, 126, 3237-3247. | 1.2 | 718 |
| 13 | Autophagy in major human diseases. <i>EMBO Journal</i> , 2021, 40, e108863. | 3.5 | 615 |
| 14 | FYCO1 is a Rab7 effector that binds to LC3 and PI3P to mediate microtubule plus end-directed vesicle transport. <i>Journal of Cell Biology</i> , 2010, 188, 253-269. | 2.3 | 573 |
| 15 | TBK-1 Promotes Autophagy-Mediated Antimicrobial Defense by Controlling Autophagosome Maturation. <i>Immunity</i> , 2012, 37, 223-234. | 6.6 | 563 |
| 16 | Autophagy mediates degradation of nuclear lamina. <i>Nature</i> , 2015, 527, 105-109. | 13.7 | 510 |
| 17 | The Adaptor Protein p62/SQSTM1 Targets Invading Bacteria to the Autophagy Pathway. <i>Journal of Immunology</i> , 2009, 183, 5909-5916. | 0.4 | 501 |
| 18 | Regulation of selective autophagy: the p62/SQSTM1 paradigm. <i>Essays in Biochemistry</i> , 2017, 61, 609-624. | 2.1 | 490 |

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|----|---|------|-----------|
| 19 | Autophagy in healthy aging and disease. <i>Nature Aging</i> , 2021, 1, 634-650. | 5.3 | 467 |
| 20 | Selective Autophagy: ATG8 Family Proteins, LIR Motifs and Cargo Receptors. <i>Journal of Molecular Biology</i> , 2020, 432, 80-103. | 2.0 | 446 |
| 21 | NBR1 and p62 as cargo receptors for selective autophagy of ubiquitinated targets. <i>Cell Cycle</i> , 2009, 8, 1986-1990. | 1.3 | 399 |
| 22 | Microenvironmental autophagy promotes tumour growth. <i>Nature</i> , 2017, 541, 417-420. | 13.7 | 379 |
| 23 | Aggrephagy: Selective Disposal of Protein Aggregates by Macroautophagy. <i>International Journal of Cell Biology</i> , 2012, 2012, 1-21. | 1.0 | 363 |
| 24 | Repeated ER α endosome contacts promote endosome translocation and neurite outgrowth. <i>Nature</i> , 2015, 520, 234-238. | 13.7 | 343 |
| 25 | TRIMs and Galectins Globally Cooperate and TRIM16 and Galectin-3 Co-direct Autophagy in Endomembrane Damage Homeostasis. <i>Developmental Cell</i> , 2016, 39, 13-27. | 3.1 | 339 |
| 26 | Interaction Codes within the Family of Mammalian Phox and Bem1p Domain-containing Proteins. <i>Journal of Biological Chemistry</i> , 2003, 278, 34568-34581. | 1.6 | 332 |
| 27 | p62/SQSTM1 and ALFY interact to facilitate the formation of p62 bodies/ALIS and their degradation by autophagy. <i>Autophagy</i> , 2010, 6, 330-344. | 4.3 | 296 |
| 28 | FKBP8 recruits LC3A to mediate Parkin-independent mitophagy. <i>EMBO Reports</i> , 2017, 18, 947-961. | 2.0 | 295 |
| 29 | Plant NBR1 is a selective autophagy substrate and a functional hybrid of the mammalian autophagic adaptors NBR1 and p62/SQSTM1. <i>Autophagy</i> , 2011, 7, 993-1010. | 4.3 | 283 |
| 30 | Delivery of Cytosolic Components by Autophagic Adaptor Protein p62 Endows Autophagosomes with Unique Antimicrobial Properties. <i>Immunity</i> , 2010, 32, 329-341. | 6.6 | 276 |
| 31 | p62/SQSTM1: A Missing Link between Protein Aggregates and the Autophagy Machinery. <i>Autophagy</i> , 2006, 2, 138-139. | 4.3 | 274 |
| 32 | NBR1 acts as an autophagy receptor for peroxisomes. <i>Journal of Cell Science</i> , 2013, 126, 939-52. | 1.2 | 274 |
| 33 | TRIM Proteins Regulate Autophagy and Can Target Autophagic Substrates by Direct Recognition. <i>Developmental Cell</i> , 2014, 30, 394-409. | 3.1 | 269 |
| 34 | p62 and NDP52 Proteins Target Intracytosolic Shigella and Listeria to Different Autophagy Pathways. <i>Journal of Biological Chemistry</i> , 2011, 286, 26987-26995. | 1.6 | 257 |
| 35 | ATG8 Family Proteins Act as Scaffolds for Assembly of the ULK Complex. <i>Journal of Biological Chemistry</i> , 2012, 287, 39275-39290. | 1.6 | 257 |
| 36 | Expression pattern of zebrafish pax genes suggests a role in early brain regionalization. <i>Nature</i> , 1991, 353, 267-270. | 13.7 | 254 |

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|----|---|------|-----------|
| 37 | TRIM-mediated precision autophagy targets cytoplasmic regulators of innate immunity. <i>Journal of Cell Biology</i> , 2015, 210, 973-989. | 2.3 | 248 |
| 38 | Dedicated <sc>SNARE</sc> s and specialized <sc>TRIM</sc> cargo receptors mediate secretory autophagy. <i>EMBO Journal</i> , 2017, 36, 42-60. | 3.5 | 247 |
| 39 | SIRT1 is downregulated by autophagy in senescence and ageing. <i>Nature Cell Biology</i> , 2020, 22, 1170-1179. | 4.6 | 236 |
| 40 | Autophagic degradation of dBruce controls DNA fragmentation in nurse cells during late <i>Drosophila melanogaster</i> oogenesis. <i>Journal of Cell Biology</i> , 2010, 190, 523-531. | 2.3 | 224 |
| 41 | Starvation induces rapid degradation of selective autophagy receptors by endosomal microautophagy. <i>Journal of Cell Biology</i> , 2018, 217, 3640-3655. | 2.3 | 213 |
| 42 | Nucleocytoplasmic Shuttling of p62/SQSTM1 and Its Role in Recruitment of Nuclear Polyubiquitinated Proteins to Promyelocytic Leukemia Bodies. <i>Journal of Biological Chemistry</i> , 2010, 285, 5941-5953. | 1.6 | 200 |
| 43 | Selective Autophagy in Cancer Development and Therapy. <i>Cancer Research</i> , 2010, 70, 3431-3434. | 0.4 | 196 |
| 44 | Galectins Control mTOR in Response to Endomembrane Damage. <i>Molecular Cell</i> , 2018, 70, 120-135.e8. | 4.5 | 191 |
| 45 | The Selective Autophagy Receptor p62 Forms a Flexible Filamentous Helical Scaffold. <i>Cell Reports</i> , 2015, 11, 748-758. | 2.9 | 190 |
| 46 | iLIR. <i>Autophagy</i> , 2014, 10, 913-925. | 4.3 | 187 |
| 47 | Phospholipase C-mediated hydrolysis of phosphatidylcholine is an important step in PDGF-stimulated DNA synthesis. <i>Cell</i> , 1990, 61, 1113-1120. | 13.5 | 179 |
| 48 | NBR1 co-operates with p62 in selective autophagy of ubiquitinated targets. <i>Autophagy</i> , 2009, 5, 732-733. | 4.3 | 163 |
| 49 | Zebrafish contains two Pax6 genes involved in eye development1The sequence reported in this paper has been deposited in the GenBank data base (accession no. AF061252).1. <i>Mechanisms of Development</i> , 1998, 77, 185-196. | 1.7 | 159 |
| 50 | Following autophagy step by step. <i>BMC Biology</i> , 2011, 9, 39. | 1.7 | 155 |
| 51 | Organization of the mitochondrial genome of Atlantic cod, <i>Gadus morhua</i> . <i>Nucleic Acids Research</i> , 1990, 18, 411-419. | 6.5 | 144 |
| 52 | A reporter cell system to monitor autophagy based on p62/SQSTM1. <i>Autophagy</i> , 2010, 6, 784-793. | 4.3 | 138 |
| 53 | Mechanisms of Selective Autophagy. <i>Annual Review of Cell and Developmental Biology</i> , 2021, 37, 143-169. | 4.0 | 137 |
| 54 | Turnip Mosaic Virus Counteracts Selective Autophagy of the Viral Silencing Suppressor HCpro. <i>Plant Physiology</i> , 2018, 176, 649-662. | 2.3 | 136 |

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|----|---|-----|-----------|
| 55 | Cell death during <i>Drosophila melanogaster</i> early oogenesis is mediated through autophagy. <i>Autophagy</i> , 2009, 5, 298-302. | 4.3 | 124 |
| 56 | Defective recognition of LC3B by mutant SQSTM1/p62 implicates impairment of autophagy as a pathogenic mechanism in ALS-FTLD. <i>Autophagy</i> , 2016, 12, 1094-1104. | 4.3 | 123 |
| 57 | SQSTM1/p62 mediates crosstalk between autophagy and the UPS in DNA repair. <i>Autophagy</i> , 2016, 12, 1917-1930. | 4.3 | 120 |
| 58 | Molecular determinants regulating selective binding of autophagy adapters and receptors to ATG8 proteins. <i>Nature Communications</i> , 2019, 10, 2055. | 5.8 | 118 |
| 59 | Cloning and sequencing of the gene encoding the phosphatidylcholine-preferring phospholipase C of <i>Bacillus cereus</i> . <i>Gene</i> , 1988, 65, 293-304. | 1.0 | 116 |
| 60 | Autophagy: links with the proteasome. <i>Current Opinion in Cell Biology</i> , 2010, 22, 192-198. | 2.6 | 113 |
| 61 | Galectins control MTOR and AMPK in response to lysosomal damage to induce autophagy. <i>Autophagy</i> , 2019, 15, 169-171. | 4.3 | 112 |
| 62 | FYCO1 Contains a C-terminally Extended, LC3A/B-preferring LC3-interacting Region (LIR) Motif Required for Efficient Maturation of Autophagosomes during Basal Autophagy. <i>Journal of Biological Chemistry</i> , 2015, 290, 29361-29374. | 1.6 | 106 |
| 63 | Genome-wide siRNA screen reveals amino acid starvation-induced autophagy requires SCOC and WAC. <i>EMBO Journal</i> , 2012, 31, 1931-1946. | 3.5 | 105 |
| 64 | NIPSNAP1 and NIPSNAP2 Act as "Eat Me" Signals for Mitophagy. <i>Developmental Cell</i> , 2019, 49, 509-525.e12.3.1 | | 104 |
| 65 | Phosphorylation of Syntaxin 17 by TBK1 Controls Autophagy Initiation. <i>Developmental Cell</i> , 2019, 49, 130-144.e6. | 3.1 | 99 |
| 66 | Selective autophagy. <i>Essays in Biochemistry</i> , 2013, 55, 79-92. | 2.1 | 98 |
| 67 | Aurothiomalate Inhibits Transformed Growth by Targeting the PB1 Domain of Protein Kinase C δ . <i>Journal of Biological Chemistry</i> , 2006, 281, 28450-28459. | 1.6 | 92 |
| 68 | Noncoding control region of naturally occurring BK virus variants: Sequence comparison and functional analysis. <i>Virus Genes</i> , 1995, 10, 261-275. | 0.7 | 89 |
| 69 | Phosphorylation of the Transactivation Domain of Pax6 by Extracellular Signal-regulated Kinase and p38 Mitogen-activated Protein Kinase. <i>Journal of Biological Chemistry</i> , 1999, 274, 15115-15126. | 1.6 | 89 |
| 70 | TRIM-directed selective autophagy regulates immune activation. <i>Autophagy</i> , 2017, 13, 989-990. | 4.3 | 86 |
| 71 | Members of the autophagy class III phosphatidylinositol 3-kinase complex I interact with GABARAP and GABARAPL1 via LIR motifs. <i>Autophagy</i> , 2019, 15, 1333-1355. | 4.3 | 86 |
| 72 | CALCOCO 1 acts with VAMP-associated proteins to mediate ER-phagy. <i>EMBO Journal</i> , 2020, 39, e103649. | 3.5 | 86 |

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|----|---|-----|-----------|
| 73 | Identification of p62/SQSTM1 as a component of non-canonical Wnt VANGL2â€‘JNK signalling in breast cancer. <i>Nature Communications</i> , 2016, 7, 10318. | 5.8 | 85 |
| 74 | ATG4B contains a C-terminal LIR motif important for binding and efficient cleavage of mammalian orthologs of yeast Atg8. <i>Autophagy</i> , 2017, 13, 834-853. | 4.3 | 84 |
| 75 | Autophagy and endocytosis â€‘ interconnections and interdependencies. <i>Journal of Cell Science</i> , 2020, 133, . | 1.2 | 83 |
| 76 | Evidence for a Bifurcation of the Mitogenic Signaling Pathway Activated by Ras and Phosphatidylcholine-hydrolyzing Phospholipase C. <i>Journal of Biological Chemistry</i> , 1995, 270, 21299-21306. | 1.6 | 71 |
| 77 | Cellular and molecular mechanism for secretory autophagy. <i>Autophagy</i> , 2017, 13, 1084-1085. | 4.3 | 71 |
| 78 | Structural basis of p62/SQSTM1 helical filaments and their role in cellular cargo uptake. <i>Nature Communications</i> , 2020, 11, 440. | 5.8 | 71 |
| 79 | Superactivation of Pax6-mediated Transactivation from Paired Domain-binding Sites by DNA-independent Recruitment of Different Homeodomain Proteins. <i>Journal of Biological Chemistry</i> , 2001, 276, 4109-4118. | 1.6 | 70 |
| 80 | Structural and Functional Analysis of a Novel Interaction Motif within UFM1-activating Enzyme 5 (UBA5) Required for Binding to Ubiquitin-like Proteins and Ufmlylation. <i>Journal of Biological Chemistry</i> , 2016, 291, 9025-9041. | 1.6 | 69 |
| 81 | Reversion of Ras- and Phosphatidylcholine-hydrolyzing Phospholipase C-mediated Transformation of NIH 3T3 Cells by a Dominant Interfering Mutant of Protein Kinase C Î» Is Accompanied by the Loss of Constitutive Nuclear Mitogen-activated Protein Kinase/Extracellular Signal-regulated Kinase Activity. <i>Journal of Biological Chemistry</i> , 1997, 272, 11557-11565. | 1.6 | 68 |
| 82 | Zebrafish Pax9 Encodes Two Proteins with Distinct C-terminal Transactivating Domains of Different Potency Negatively Regulated by Adjacent N-terminal Sequences. <i>Journal of Biological Chemistry</i> , 1996, 271, 26914-26923. | 1.6 | 67 |
| 83 | The proteomic analysis of endogenous FAT10 substrates identifies p62/SQSTM1 as a substrate of FAT10ylation. <i>Journal of Cell Science</i> , 2012, 125, 4576-85. | 1.2 | 67 |
| 84 | FYCO1: Linking autophagosomes to microtubule plus end-directing molecular motors. <i>Autophagy</i> , 2010, 6, 550-552. | 4.3 | 65 |
| 85 | TRIM proteins regulate autophagy: TRIM5 is a selective autophagy receptor mediating HIV-1 restriction. <i>Autophagy</i> , 2014, 10, 2387-2388. | 4.3 | 64 |
| 86 | Nuclear Import and Export Signals Enable Rapid Nucleocytoplasmic Shuttling of the Atypical Protein Kinase C Î». <i>Journal of Biological Chemistry</i> , 2001, 276, 13015-13024. | 1.6 | 62 |
| 87 | Autophagy as a trigger for cell death: Autophagic degradation of inhibitor of apoptosis dBruce controls DNA fragmentation during late oogenesis in <i>Drosophila</i> . <i>Autophagy</i> , 2010, 6, 1214-1215. | 4.3 | 61 |
| 88 | p62/Sequestosome-1, Autophagy-related Gene 8, and Autophagy in <i>Drosophila</i> Are Regulated by Nuclear Factor Erythroid 2-related Factor 2 (NRF2), Independent of Transcription Factor TFEB. <i>Journal of Biological Chemistry</i> , 2015, 290, 14945-14962. | 1.6 | 61 |
| 89 | Nrf2 and SQSTM1/p62 jointly contribute to mesenchymal transition and invasion in glioblastoma. <i>Oncogene</i> , 2019, 38, 7473-7490. | 2.6 | 61 |
| 90 | Nucleotide sequence of the <i>Physarum polycephalum</i> small subunit ribosomal RNA as inferred from the gene sequence: secondary structure and evolutionary implications. <i>Current Genetics</i> , 1988, 14, 265-273. | 0.8 | 60 |

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|-----|---|-----|-----------|
| 91 | Conserved Atg8 recognition sites mediate Atg4 association with autophagosomal membranes and Atg8 deconjugation. <i>EMBO Reports</i> , 2017, 18, 765-780. | 2.0 | 59 |
| 92 | SQSTM1/p62 regulates the expression of junctional proteins through epithelial-mesenchymal transition factors. <i>Cell Cycle</i> , 2015, 14, 364-374. | 1.3 | 57 |
| 93 | Structural and functional analyses of DNA bending induced by Sp1 family transcription factors 1 Edited by T. Richmond. <i>Journal of Molecular Biology</i> , 1997, 267, 490-504. | 2.0 | 56 |
| 94 | SIRT1 â€“ a new mammalian substrate of nuclear autophagy. <i>Autophagy</i> , 2021, 17, 593-595. | 4.3 | 56 |
| 95 | p38MAPK-regulated induction of p62 and NBR1 after photodynamic therapy promotes autophagic clearance of ubiquitin aggregates and reduces reactive oxygen species levels by supporting Nrf2â€“antioxidant signaling. <i>Free Radical Biology and Medicine</i> , 2014, 67, 292-303. | 1.3 | 55 |
| 96 | Comparative Analyses of LTRs of the ERV-H Family of Primate-Specific Retrovirus-like Elements Isolated from Marmoset, African Green Monkey, and Man. <i>Virology</i> , 1997, 234, 14-30. | 1.1 | 54 |
| 97 | Structure and evolution of myxomycete nuclear group I introns: a model for horizontal transfer by intron homing. <i>Current Genetics</i> , 1992, 22, 297-304. | 0.8 | 53 |
| 98 | DOR/Tp53inp2 and Tp53inp1 Constitute a Metazoan Gene Family Encoding Dual Regulators of Autophagy and Transcription. <i>PLoS ONE</i> , 2012, 7, e34034. | 1.1 | 51 |
| 99 | Dynamic subcellular localization of the mono-ADP-ribosyltransferase ARTD10 and interaction with the ubiquitin receptor p62. <i>Cell Communication and Signaling</i> , 2012, 10, 28. | 2.7 | 50 |
| 100 | HIV-1 viral infectivity factor interacts with microtubule-associated protein light chain 3 and inhibits autophagy. <i>Aids</i> , 2015, 29, 275-286. | 1.0 | 50 |
| 101 | Kenny mediates selective autophagic degradation of the IKK complex to control innate immune responses. <i>Nature Communications</i> , 2017, 8, 1264. | 5.8 | 50 |
| 102 | The Nuclear Factor SPBP Contains Different Functional Domains and Stimulates the Activity of Various Transcriptional Activators. <i>Journal of Biological Chemistry</i> , 2000, 275, 40288-40300. | 1.6 | 49 |
| 103 | Rapid disappearance of one parental mitochondrial genotype after isogamous mating in the myxomycete <i>Physarum polycephalum</i> . <i>Current Genetics</i> , 1991, 19, 55-59. | 0.8 | 47 |
| 104 | A novel Bcr-Abl splice isoform is associated with the L248V mutation in CML patients with acquired resistance to imatinib. <i>Leukemia</i> , 2006, 20, 2057-2060. | 3.3 | 45 |
| 105 | The MH1 domain of Smad3 interacts with Pax6 and represses autoregulation of the Pax6 P1 promoter. <i>Nucleic Acids Research</i> , 2007, 35, 890-901. | 6.5 | 44 |
| 106 | ATG9A protects the plasma membrane from programmed and incidental permeabilization. <i>Nature Cell Biology</i> , 2021, 23, 846-858. | 4.6 | 43 |
| 107 | Members of the RTVL-H family of human endogenous retrovirus-like elements are expressed in placenta. <i>Gene</i> , 1989, 79, 259-267. | 1.0 | 42 |
| 108 | TRIM17 contributes to autophagy of midbodies while actively sparing other targets from degradation. <i>Journal of Cell Science</i> , 2016, 129, 3562-3573. | 1.2 | 40 |

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|-----|--|-----|-----------|
| 109 | Galectins and TRIMs directly interact and orchestrate autophagic response to endomembrane damage. <i>Autophagy</i> , 2017, 13, 1086-1087. | 4.3 | 40 |
| 110 | TRIM50 regulates Beclin 1 proautophagic activity. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2018, 1865, 908-919. | 1.9 | 39 |
| 111 | SAMM50 acts with p62 in piecemeal basal- and OXPHOS-induced mitophagy of SAM and MICOS components. <i>Journal of Cell Biology</i> , 2021, 220, . | 2.3 | 39 |
| 112 | Mammalian Atg8 proteins regulate lysosome and autolysosome biogenesis through <sc>SNARE</sc> s. <i>EMBO Journal</i> , 2019, 38, e101994. | 3.5 | 37 |
| 113 | NIPSNAP1 and NIPSNAP2 act as "eat me"-signals to allow sustained recruitment of autophagy receptors during mitophagy. <i>Autophagy</i> , 2019, 15, 1845-1847. | 4.3 | 35 |
| 114 | Regulation of Golgi turnover by CALCOCO1-mediated selective autophagy. <i>Journal of Cell Biology</i> , 2021, 220, . | 2.3 | 35 |
| 115 | SPBP Is a Sulforaphane Induced Transcriptional Coactivator of NRF2 Regulating Expression of the Autophagy Receptor p62/SQSTM1. <i>PLoS ONE</i> , 2014, 9, e85262. | 1.1 | 35 |
| 116 | Extrachromosomal ribosomal DNA of <i>Didymium iridis</i> : sequence analysis of the large subunit ribosomal RNA gene and sub-telomeric region. <i>Current Genetics</i> , 1992, 22, 305-312. | 0.8 | 32 |
| 117 | Regulator of Chromosome Condensation 2 Identifies High-Risk Patients within Both Major Phenotypes of Colorectal Cancer. <i>Clinical Cancer Research</i> , 2015, 21, 3759-3770. | 3.2 | 32 |
| 118 | Endosomal microautophagy is an integrated part of the autophagic response to amino acid starvation. <i>Autophagy</i> , 2019, 15, 182-183. | 4.3 | 32 |
| 119 | The third helix of the homeodomain of paired class homeodomain proteins acts as a recognition helix both for DNA and protein interactions. <i>Nucleic Acids Research</i> , 2005, 33, 2661-2675. | 6.5 | 29 |
| 120 | Zebrafish pou[c]: a divergent POU family gene ubiquitously expressed during embryogenesis. <i>Nucleic Acids Research</i> , 1993, 21, 475-483. | 6.5 | 28 |
| 121 | cDNA sequence of zebrafish (<i>Brachydanio rerio</i>) translation elongation factor-1 \pm : Molecular phylogeny of eukaryotes based on elongation factor-1 \pm protein sequences. <i>Biochimica Et Biophysica Acta Gene Regulatory Mechanisms</i> , 1994, 1219, 529-532. | 2.4 | 28 |
| 122 | Identification of two independent nucleosome-binding domains in the transcriptional co-activator SPBP. <i>Biochemical Journal</i> , 2012, 442, 65-75. | 1.7 | 28 |
| 123 | The ePHD protein SPBP interacts with TopBP1 and together they co-operate to stimulate Ets1-mediated transcription. <i>Nucleic Acids Research</i> , 2007, 35, 6648-6662. | 6.5 | 26 |
| 124 | TAK 1 converts Sequestosome 1/p62 from an autophagy receptor to a signaling platform. <i>EMBO Reports</i> , 2019, 20, e46238. | 2.0 | 24 |
| 125 | A Phylogenetic Study of SPBP and RAI1: Evolutionary Conservation of Chromatin Binding Modules. <i>PLoS ONE</i> , 2013, 8, e78907. | 1.1 | 22 |
| 126 | NIMA-related kinase 9 α -mediated phosphorylation of the microtubule-associated LC3B protein at Thr-50 suppresses selective autophagy of p62/sequestosome 1. <i>Journal of Biological Chemistry</i> , 2020, 295, 1240-1260. | 1.6 | 19 |

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| 127 | Regulation of Expression of Autophagy Genes by Atg8a-Interacting Partners Sequoia, YL-1, and Sir2 in <i>Drosophila</i> . <i>Cell Reports</i> , 2020, 31, 107695. | 2.9 | 19 |
| 128 | The FMRpolyGlycine Protein Mediates Aggregate Formation and Toxicity Independent of the CGG mRNA Hairpin in a Cellular Model for FXTAS. <i>Frontiers in Genetics</i> , 2019, 10, 249. | 1.1 | 18 |
| 129 | Zonda is a novel early component of the autophagy pathway in <i>Drosophila</i> . <i>Molecular Biology of the Cell</i> , 2017, 28, 3070-3081. | 0.9 | 17 |
| 130 | Bacillus cereus strain SE-1: nucleotide sequence of the sphingomyelinase C gene. <i>Nucleic Acids Research</i> , 1988, 16, 10370-10370. | 6.5 | 16 |
| 131 | SQSTM-1/p62 potentiates HTLV-1 Tax-mediated NF- κ B activation through its ubiquitin binding function. <i>Scientific Reports</i> , 2019, 9, 16014. | 1.6 | 15 |
| 132 | Pax6 Represses Androgen Receptor-Mediated Transactivation by Inhibiting Recruitment of the Coactivator SPBP. <i>PLoS ONE</i> , 2011, 6, e24659. | 1.1 | 14 |
| 133 | TRIM32 acts both as a substrate and a positive regulator of p62/SQSTM1 impaired in a muscular dystrophy disease. <i>Journal of Cell Science</i> , 2019, 132, . | 1.2 | 14 |
| 134 | Bicaudal D1 impairs autophagosome maturation in chronic obstructive pulmonary disease. <i>FASEB BioAdvances</i> , 2019, 1, 688-705. | 1.3 | 14 |
| 135 | Phosphorylation of the LIR Domain of SCOC Modulates ATG8 Binding Affinity and Specificity. <i>Journal of Molecular Biology</i> , 2021, 433, 166987. | 2.0 | 14 |
| 136 | NIMA-related kinase N^{E} -mediated phosphorylation of the microtubule-associated LC3B protein at Thr-50 suppresses selective autophagy of p62/sequestosome 1. <i>Journal of Biological Chemistry</i> , 2020, 295, 1240-1260. | 1.6 | 14 |
| 137 | Selective autophagy goes exclusive. <i>Nature Cell Biology</i> , 2014, 16, 395-397. | 4.6 | 11 |
| 138 | Sequence analysis of 12 structural genes and a novel non-coding region from mitochondrial DNA of Atlantic cod, <i>Gadus morhua</i> . <i>Biochimica Et Biophysica Acta Gene Regulatory Mechanisms</i> , 1994, 1218, 213-217. | 2.4 | 10 |
| 139 | Pax6 localizes to chromatin-rich territories and displays a slow nuclear mobility altered by disease mutations. <i>Cellular and Molecular Life Sciences</i> , 2010, 67, 4079-4094. | 2.4 | 9 |
| 140 | CALCOCO1 is a soluble reticulophagy receptor. <i>Autophagy</i> , 2020, 16, 1729-1731. | 4.3 | 9 |
| 141 | Use of Peptide Arrays for Identification and Characterization of LIR Motifs. <i>Methods in Molecular Biology</i> , 2019, 1880, 149-161. | 0.4 | 8 |
| 142 | The soluble reticulophagy receptor CALCOCO1 is also a Golgiphagy receptor. <i>Autophagy</i> , 2021, 17, 2051-2052. | 4.3 | 8 |
| 143 | The putative origin of heavy strand replication (oriH) in mitochondrial DNA is highly conserved among the teleost fishes. <i>DNA Sequence</i> , 1993, 3, 397-399. | 0.7 | 6 |
| 144 | Degradation of arouser by endosomal microautophagy is essential for adaptation to starvation in <i>Drosophila</i> . <i>Life Science Alliance</i> , 2021, 4, e202000965. | 1.3 | 6 |

| # | ARTICLE | IF | CITATIONS |
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