

Masahide Oku

List of Publications by Year in descending order

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papers

2,248
citations

304602

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docs citations

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times ranked

4810
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#	ARTICLE	IF	CITATIONS
1	Regulation of Peroxisome Homeostasis by Post-Translational Modification in the Methylophilic Yeast <i>Komagataella phaffii</i> . <i>Frontiers in Cell and Developmental Biology</i> , 2022, 10, 887806.	1.8	4
2	The methanol sensor Wsc1 and MAPK Mpk1 suppress degradation of methanol-induced peroxisomes in methylophilic yeast. <i>Journal of Cell Science</i> , 2021, 134, .	1.2	6
3	A peroxisome deficiency-induced reductive cytosol state up-regulates the brain-derived neurotrophic factor pathway. <i>Journal of Biological Chemistry</i> , 2020, 295, 5321-5334.	1.6	12
4	Peroxisomal Fba2p and Tal2p complementally function in the rearrangement pathway for xylulose 5-phosphate in the methylophilic yeast <i>Pichia pastoris</i> . <i>Journal of Bioscience and Bioengineering</i> , 2019, 128, 33-38.	1.1	9
5	Evolution from covalent conjugation to non-covalent interaction in the ubiquitin-like ATG12 system. <i>Nature Structural and Molecular Biology</i> , 2019, 26, 289-296.	3.6	39
6	Peroxisome Degradation and Its Molecular Machinery. , 2019, , 43-58.		0
7	Three Distinct Types of Microautophagy Based on Membrane Dynamics and Molecular Machineries. <i>BioEssays</i> , 2018, 40, e1800008.	1.2	180
8	Ethanol represses the expression of methanol-inducible genes via acetyl-CoA synthesis in the yeast <i>Komagataella phaffii</i> . <i>Scientific Reports</i> , 2018, 8, 18051.	1.6	10
9	Experimental Systems to Study Yeast Pexophagy. <i>Methods in Molecular Biology</i> , 2017, 1595, 249-255.	0.4	3
10	Evidence for ESCRT- and clathrin-dependent microautophagy. <i>Journal of Cell Biology</i> , 2017, 216, 3263-3274.	2.3	127
11	Role of Acyl Chain Composition of Phosphatidylcholine in Tafazzin-Mediated Remodeling of Cardiolipin in Liposomes. <i>Biochemistry</i> , 2017, 56, 6268-6280.	1.2	17
12	Autophagy-independent function of Atg8 in lipid droplet dynamics in yeast. <i>Journal of Biochemistry</i> , 2016, 161, mvw078.	0.9	10
13	Mechanism for Remodeling of the Acyl Chain Composition of Cardiolipin Catalyzed by <i>Saccharomyces cerevisiae</i> Tafazzin. <i>Journal of Biological Chemistry</i> , 2016, 291, 15491-15502.	1.6	24
14	Pexophagy in yeasts. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2016, 1863, 992-998.	1.9	48
15	A defect of the vacuolar putative lipase Atg15 accelerates degradation of lipid droplets through lipolysis. <i>Autophagy</i> , 2015, 11, 1247-1258.	4.3	32
16	Yeast nitrogen utilization in the phyllosphere during plant lifespan under regulation of autophagy. <i>Scientific Reports</i> , 2015, 5, 9719.	1.6	17
17	Regulation of nitrate and methylamine metabolism by multiple nitrogen sources in the methylophilic yeast <i>Candida boidinii</i> . <i>FEMS Yeast Research</i> , 2015, 15, fov084.	1.1	7
18	The emerging role of autophagy in peroxisome dynamics and lipid metabolism of phyllosphere microorganisms. <i>Frontiers in Plant Science</i> , 2014, 5, 81.	1.7	9

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19	Atg21 regulates pexophagy via its PI(3)P-binding activity in <i>Pichia pastoris</i> . <i>FEMS Yeast Research</i> , 2014, 14, 435-444.	1.1	6
20	Functional link between R-ab GTPase-mediated membrane trafficking and PI4,5P ₂ signaling. <i>Genes To Cells</i> , 2014, 19, 177-197.	0.5	2
21	The Tor and Sin3-Rpd3 complex regulate expression of the mitophagy receptor protein Atg32. <i>Journal of Cell Science</i> , 2014, 127, 3184-96.	1.2	40
22	Mitochondrial impairment triggers cytosolic oxidative stress and cell death following proteasome inhibition. <i>Scientific Reports</i> , 2014, 4, 5896.	1.6	168
23	Hyper-Activation of the Target of Rapamycin (Tor) Kinase 1 Decreases Intracellular Glutathione Content in <i>Saccharomyces cerevisiae</i> as Revealed by LC-MS/MS Analysis. <i>Bioscience, Biotechnology and Biochemistry</i> , 2013, 77, 1608-1611.	0.6	3
24	A fluorescence resonance energy transfer (FRET)-based redox sensor reveals physiological role of thioredoxin in the yeast <i>Saccharomyces cerevisiae</i> . <i>FEBS Letters</i> , 2013, 587, 793-798.	1.3	14
25	Atg18 phosphoregulation controls organellar dynamics by modulating its phosphoinositide-binding activity. <i>Journal of Cell Biology</i> , 2013, 202, 685-698.	2.3	45
26	Atg18 lifts up from and lands on the vacuolar membrane mediated by phosphorylation of its propellers. <i>Autophagy</i> , 2013, 9, 2161-2162.	4.3	1
27	Assessment of Physiological Redox State with Novel FRET Protein Probes. <i>Antioxidants and Redox Signaling</i> , 2012, 16, 698-704.	2.5	15
28	Yeast Methylo-trophy and Autophagy in a Methanol-Oscillating Environment on Growing <i>Arabidopsis thaliana</i> Leaves. <i>PLoS ONE</i> , 2011, 6, e25257.	1.1	51
29	Inhibition of surgical trauma-enhanced peritoneal dissemination of tumor cells by human catalase derivatives in mice. <i>Free Radical Biology and Medicine</i> , 2011, 51, 773-779.	1.3	6
30	Yeast Methylo-trophy: Metabolism, Gene Regulation and Peroxisome Homeostasis. <i>International Journal of Microbiology</i> , 2011, 2011, 1-8.	0.9	113
31	Peroxisomes as dynamic organelles: autophagic degradation. <i>FEBS Journal</i> , 2010, 277, 3289-3294.	2.2	72
32	A Novel Fluorescent Sensor Protein for Visualization of Redox States in the Cytoplasm and in Peroxisomes. <i>Molecular and Cellular Biology</i> , 2010, 30, 3758-3766.	1.1	100
33	Atg8 regulates vacuolar membrane dynamics in a lipidation-independent manner in <i>Pichia pastoris</i> . <i>Journal of Cell Science</i> , 2010, 123, 4107-4116.	1.2	52
34	Atg8 regulates vacuolar membrane dynamics in a lipidation-independent manner in <i>Pichia pastoris</i> . <i>Development (Cambridge)</i> , 2010, 137, e2406-e2406.	1.2	0
35	Atg26-Mediated Pexophagy Is Required for Host Invasion by the Plant Pathogenic Fungus <i>Colletotrichum orbiculare</i> . <i>Plant Cell</i> , 2009, 21, 1291-1304.	3.1	138
36	Control of hypoxia-induced tumor cell adhesion by cytophilic human catalase. <i>Free Radical Biology and Medicine</i> , 2009, 47, 1772-1778.	1.3	4

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37	Lagâ€­phase autophagy in the methylotrophic yeast <i>Pichia pastoris</i> . <i>Genes To Cells</i> , 2009, 14, 861-870.	0.5	18
38	Chapter 15 Pexophagy in <i>Pichia pastoris</i> . <i>Methods in Enzymology</i> , 2008, 451, 217-228.	0.4	8
39	Functions of PI4P and Sterol Glucoside Necessary for the Synthesis of a Nascent Membrane Structure During Pexophagy. <i>Autophagy</i> , 2007, 3, 35-37.	4.3	16
40	Pexophagy: Autophagic degradation of peroxisomes. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2006, 1763, 1767-1775.	1.9	193
41	Role of Vac8 in Formation of the Vacuolar Sequestering Membrane during Micropexophagy. <i>Autophagy</i> , 2006, 2, 272-279.	4.3	28
42	PI4P-signaling pathway for the synthesis of a nascent membrane structure in selective autophagy. <i>Journal of Cell Biology</i> , 2006, 173, 709-717.	2.3	77
43	A Sorting Nexin PpAtg24 Regulates Vacuolar Membrane Dynamics during Pexophagy via Binding to Phosphatidylinositol-3-Phosphate. <i>Molecular Biology of the Cell</i> , 2005, 16, 446-457.	0.9	69
44	Pexophagy: The Selective Autophagy of Peroxisomes. <i>Autophagy</i> , 2005, 1, 75-83.	4.3	250
45	Peroxisome degradation requires catalytically active sterol glucosyltransferase with a GRAM domain. <i>EMBO Journal</i> , 2003, 22, 3231-3241.	3.5	96
46	Paz2 and 13 otherPAZgene products regulate vacuolar engulfment of peroxisomes during micropexophagy. <i>Genes To Cells</i> , 2002, 7, 75-90.	0.5	109