

Yasuyuki Suda

List of Publications by Year in descending order

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papers

826
citations

567281

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27
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all docs

32
docs citations

32
times ranked

1077
citing authors

#	ARTICLE	IF	CITATIONS
1	Dynamic Behavior of the trans-Golgi Network in Root Tissues of Arabidopsis Revealed by Super-Resolution Live Imaging. <i>Plant and Cell Physiology</i> , 2014, 55, 694-703.	3.1	94
2	Visualization of secretory cargo transport within the Golgi apparatus. <i>Journal of Cell Biology</i> , 2019, 218, 1602-1618.	5.2	63
3	Rab GAP cascade regulates dynamics of Ypt6 in the Golgi traffic. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 18976-18981.	7.1	62
4	Phosphatidylinositol 3-Kinase and 4-Kinase Have Distinct Roles in Intracellular Trafficking of Cellulose Synthase Complexes in Arabidopsis thaliana. <i>Plant and Cell Physiology</i> , 2015, 56, 287-298.	3.1	60
5	Alternative Modes of Organellar Segregation during Sporulation in <i>Saccharomyces cerevisiae</i> . <i>Eukaryotic Cell</i> , 2007, 6, 2009-2017.	3.4	52
6	Erv14 family cargo receptors are necessary for ER exit during sporulation in <i>Saccharomyces cerevisiae</i> . <i>Journal of Cell Science</i> , 2007, 120, 908-916.	2.0	50
7	COPI is essential for Golgi cisternal maturation and dynamics. <i>Journal of Cell Science</i> , 2016, 129, 3251-61.	2.0	49
8	Live Cell Visualization of Golgi Membrane Dynamics by Super-resolution Confocal Live Imaging Microscopy. <i>Methods in Cell Biology</i> , 2013, 118, 235-242.	1.1	47
9	The Yeast <i>Golgi</i> Apparatus. <i>Traffic</i> , 2012, 13, 505-510.	2.7	46
10	Spatiotemporal dissection of the trans-Golgi network. <i>Journal of Cell Science</i> , 2019, 132, .	2.0	42
11	A Screen for Spore Wall Permeability Mutants Identifies a Secreted Protease Required for Proper Spore Wall Assembly. <i>PLoS ONE</i> , 2009, 4, e7184.	2.5	36
12	<i>Saccharomyces cerevisiae</i> QNS1 codes for NAD ⁺ synthetase that is functionally conserved in mammals. <i>Yeast</i> , 2003, 20, 995-1005.	1.7	30
13	Sar1 localizes at the rims of COPII-coated membranes in vivo. <i>Journal of Cell Science</i> , 2016, 129, 3231-7.	2.0	30
14	Regulation of ER-Golgi Transport Dynamics by GTPases in Budding Yeast. <i>Frontiers in Cell and Developmental Biology</i> , 2017, 5, 122.	3.7	29
15	Vesicle Docking to the Spindle Pole Body Is Necessary to Recruit the Exocyst During Membrane Formation in <i>Saccharomyces cerevisiae</i> . <i>Molecular Biology of the Cell</i> , 2010, 21, 3693-3707.	2.1	19
16	Cytoplasmic deadenylase Ccr4 is required for translational repression of LRG1 mRNA in the stationary phase. <i>PLoS ONE</i> , 2017, 12, e0172476.	2.5	15
17	Analysis of the Physiological Activities of Scd6 through Its Interaction with Hmt1. <i>PLoS ONE</i> , 2016, 11, e0164773.	2.5	12
18	Different Regulations of ROM2 and LRG1 Expression by Ccr4, Pop2, and Dhh1 in the <i>Saccharomyces cerevisiae</i> Cell Wall Integrity Pathway. <i>MSphere</i> , 2016, 1, .	2.9	12

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19	Suppression of Vps13 adaptor protein mutants reveals a central role for PI4P in regulating prospore membrane extension. <i>PLoS Genetics</i> , 2021, 17, e1009727.	3.5	12
20	Protein Phosphatase Type 1-Interacting Protein Ysw1 Is Involved in Proper Septin Organization and Prospore Membrane Formation during Sporulation. <i>Eukaryotic Cell</i> , 2009, 8, 1027-1037.	3.4	11
21	The Dysferlin Domain-Only Protein, Spo73, Is Required for Prospore Membrane Extension in <i>Saccharomyces cerevisiae</i> . <i>MSphere</i> , 2016, 1, .	2.9	10
22	Dynamic localization of a yeast development-specific PP1 complex during prospore membrane formation is dependent on multiple localization signals and complex formation. <i>Molecular Biology of the Cell</i> , 2017, 28, 3881-3895.	2.1	9
23	Regulation of LRG1 expression by RNA-binding protein Puf5 in the budding yeast cell wall integrity pathway. <i>Genes To Cells</i> , 2018, 23, 988-997.	1.2	8
24	Activation of Rab GTPase Sec4 by its GEF Sec2 is required for prospore membrane formation during sporulation in yeast <i>Saccharomyces cerevisiae</i> . <i>FEMS Yeast Research</i> , 2018, 18, .	2.3	6
25	Pop2 phosphorylation at S39 contributes to the glucose repression of stress response genes, HSP12 and HSP26. <i>PLoS ONE</i> , 2019, 14, e0215064.	2.5	5
26	Pbp1 mediates the aberrant expression of genes involved in growth defect of <i>ccr4</i> and <i>pop2</i> mutants in yeast <i>Saccharomyces cerevisiae</i> . <i>Genes To Cells</i> , 2021, 26, 381-398.	1.2	5
27	Yeast Dop1 is required for glycosyltransferase retrieval from the trans-Golgi network. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2019, 1863, 1147-1157.	2.4	4
28	Pbp1, the yeast ortholog of human Ataxin-2, functions in the cell growth on non-fermentable carbon sources. <i>PLoS ONE</i> , 2021, 16, e0251456.	2.5	4
29	Regulation of CLB6 expression by the cytoplasmic deadenylase Ccr4 through its coding and 3' UTR regions. <i>PLoS ONE</i> , 2022, 17, e0268283.	2.5	2
30	The eIF4E-binding protein Eap1 has similar but independent roles in cell growth and gene expression with the cytoplasmic deadenylase Ccr4. <i>Bioscience, Biotechnology and Biochemistry</i> , 2021, 85, 1452-1459.	1.3	1
31	Pan2-Pan3 complex, together with Ccr4-Not complex, has a role in the cell growth on non-fermentable carbon sources. <i>Biochemical and Biophysical Research Communications</i> , 2021, 570, 125-130.	2.1	1