

Kazuo Tatebayashi

List of Publications by Year in descending order

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26
papers

1,874
citations

361413

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552781

26
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docs citations

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times ranked

1827
citing authors

#	ARTICLE	IF	CITATIONS
1	Transmembrane mucins Hkr1 and Msb2 are putative osmosensors in the SHO1 branch of yeast HOG pathway. <i>EMBO Journal</i> , 2007, 26, 3521-3533.	7.8	204
2	Regulation of the Osmoregulatory HOG MAPK Cascade in Yeast. <i>Journal of Biochemistry</i> , 2004, 136, 267-272.	1.7	200
3	Smad-dependent GADD45beta expression mediates delayed activation of p38 MAP kinase by TGF-beta. <i>EMBO Journal</i> , 2002, 21, 6473-6482.	7.8	162
4	Adaptor functions of Cdc42, Ste50, and Sho1 in the yeast osmoregulatory HOG MAPK pathway. <i>EMBO Journal</i> , 2006, 25, 3033-3044.	7.8	148
5	Conserved Docking Site Is Essential for Activation of Mammalian MAP Kinase Kinases by Specific MAP Kinase Kinase Kinases. <i>Molecular Cell</i> , 2005, 18, 295-306.	9.7	146
6	Regulation of Initiation of S Phase, Replication Checkpoint Signaling, and Maintenance of Mitotic Chromosome Structures during S Phase by Hsk1 Kinase in the Fission Yeast. <i>Molecular Biology of the Cell</i> , 2001, 12, 1257-1274.	2.1	106
7	Yeast Osmosensors Hkr1 and Msb2 Activate the Hog1 MAPK Cascade by Different Mechanisms. <i>Science Signaling</i> , 2014, 7, ra21.	3.6	92
8	A docking site determining specificity of Pbs2 MAPKK for Ssk2/Ssk22 MAPKKs in the yeast HOG pathway. <i>EMBO Journal</i> , 2003, 22, 3624-3634.	7.8	91
9	Bloom's syndrome gene suppresses premature ageing caused by Sgs1 deficiency in yeast. <i>Genes To Cells</i> , 1999, 4, 619-625.	1.2	84
10	Dynamic Control of Yeast MAP Kinase Network by Induced Association and Dissociation between the Ste50 Scaffold and the Opy2 Membrane Anchor. <i>Molecular Cell</i> , 2010, 40, 87-98.	9.7	80
11	Glycosylation defects activate filamentous growth Kss1 MAPK and inhibit osmoregulatory Hog1 MAPK. <i>EMBO Journal</i> , 2009, 28, 1380-1391.	7.8	73
12	Isolation of a <i>Schizosaccharomyces pombe</i> rad21ts Mutant That Is Aberrant in Chromosome Segregation, Microtubule Function, DNA Repair and Sensitive to Hydroxyurea: Possible Involvement of Rad21 in Ubiquitin-Mediated Proteolysis. <i>Genetics</i> , 1998, 148, 49-57.	2.9	72
13	Phosphorylated Ssk1 Prevents Unphosphorylated Ssk1 from Activating the Ssk2 Mitogen-Activated Protein Kinase Kinase Kinase in the Yeast High-Osmolarity Glycerol Osmoregulatory Pathway. <i>Molecular and Cellular Biology</i> , 2008, 28, 5172-5183.	2.3	56
14	Two Adjacent Docking Sites in the Yeast Hog1 Mitogen-Activated Protein (MAP) Kinase Differentially Interact with the Pbs2 MAP Kinase Kinase and the Ptp2 Protein Tyrosine Phosphatase. <i>Molecular and Cellular Biology</i> , 2008, 28, 2481-2494.	2.3	52
15	Osmosensing and scaffolding functions of the oligomeric four-transmembrane domain osmosensor Sho1. <i>Nature Communications</i> , 2015, 6, 6975.	12.8	46
16	Osmostress enhances activating phosphorylation of Hog1 MAP kinase by mono-phosphorylated Pbs2 MAP 2K. <i>EMBO Journal</i> , 2020, 39, e103444.	7.8	44
17	The dhp1+ gene, encoding a putative nuclear 5'->3' exonuclease, is required for proper chromosome segregation in fission yeast. <i>Nucleic Acids Research</i> , 2001, 29, 1326-1333.	14.5	38
18	Yeast recombination pathways triggered by topoisomerase II-mediated DNA breaks. <i>Nucleic Acids Research</i> , 2003, 31, 4373-4384.	14.5	36

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19	Scaffold Protein Ahk1, Which Associates with Hkr1, Sho1, Ste11, and Pbs2, Inhibits Cross Talk Signaling from the Hkr1 Osmosensor to the Kss1 Mitogen-Activated Protein Kinase. <i>Molecular and Cellular Biology</i> , 2016, 36, 1109-1123.	2.3	24
20	The RHC21 gene of budding yeast, a homologue of the fission yeast <i>rad21</i> gene, is essential for chromosome segregation. <i>Molecular Genetics and Genomics</i> , 1998, 257, 149-156.	2.4	22
21	Binding of the Extracellular Eight-Cysteine Motif of Opy2 to the Putative Osmosensor Msb2 Is Essential for Activation of the Yeast High-Osmolarity Glycerol Pathway. <i>Molecular and Cellular Biology</i> , 2016, 36, 475-487.	2.3	22
22	The budding yeast cohesin gene <i>SCC1</i> / <i>MCD1</i> / <i>RHC21</i> genetically interacts with PKA, CDK and APC. <i>Current Genetics</i> , 1999, 36, 329-338.	1.7	20
23	Interaction between the transmembrane domains of Sho1 and Opy2 enhances the signaling efficiency of the Hog1 MAP kinase cascade in <i>Saccharomyces cerevisiae</i> . <i>PLoS ONE</i> , 2019, 14, e0211380.	2.5	18
24	Effect of the DNA topoisomerase II inhibitor VP-16 on illegitimate recombination in yeast chromosomes. <i>Gene</i> , 2002, 291, 251-257.	2.2	17
25	Structural analyses of DNA fragments integrated by illegitimate recombination in <i>Schizosaccharomyces pombe</i> . <i>Molecular Genetics and Genomics</i> , 1994, 244, 111-119.	2.4	11
26	Identification of novel suppressors for <i>Mog1</i> implies its involvement in RNA metabolism, lipid metabolism and signal transduction. <i>Gene</i> , 2007, 400, 114-121.	2.2	6