

Joaquin Arino

List of Publications by Year in descending order

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154
papers

6,088
citations

61857

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88477

70
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155
all docs

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docs citations

155
times ranked

4825
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparative Analysis of Type 1 and Type Z Protein Phosphatases Reveals D615 as a Key Residue for Ppz1 Regulation. <i>International Journal of Molecular Sciences</i> , 2022, 23, 1327.	1.8	3
2	The toxic effects of yeast Ppz1 phosphatase are counteracted by subcellular relocalization mediated by its regulatory subunit Hal3. <i>FEBS Letters</i> , 2022, 596, 1556-1566.	1.3	5
3	When Phosphatases Go Mad: The Molecular Basis for Toxicity of Yeast Ppz1. <i>International Journal of Molecular Sciences</i> , 2022, 23, 4304.	1.8	1
4	The Toxic Effects of Ppz1 Overexpression Involve Nha1-Mediated Dereglulation of K ⁺ and H ⁺ Homeostasis. <i>Journal of Fungi (Basel, Switzerland)</i> , 2021, 7, 1010.	1.5	6
5	The N-Acetylglucosamine Kinase from <i>Yarrowia lipolytica</i> Is a Moonlighting Protein. <i>International Journal of Molecular Sciences</i> , 2021, 22, 13109.	1.8	1
6	Yeast Ppz1 protein phosphatase toxicity involves the alteration of multiple cellular targets. <i>Scientific Reports</i> , 2020, 10, 15613.	1.6	18
7	Controlling Ser/Thr protein phosphatase PP1 activity and function through interaction with regulatory subunits. <i>Advances in Protein Chemistry and Structural Biology</i> , 2020, 122, 231-288.	1.0	19
8	The N-Terminal Region of Yeast Protein Phosphatase Ppz1 Is a Determinant for Its Toxicity. <i>International Journal of Molecular Sciences</i> , 2020, 21, 7733.	1.8	4
9	Overexpression of budding yeast protein phosphatase Ppz1 impairs translation. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2020, 1867, 118727.	1.9	13
10	Protein Phosphatase Ppz1 Is Not Regulated by a Hal3-Like Protein in Plant Pathogen <i>Ustilago maydis</i> . <i>International Journal of Molecular Sciences</i> , 2019, 20, 3817.	1.8	5
11	Monovalent cation transporters at the plasma membrane in yeasts. <i>Yeast</i> , 2019, 36, 177-193.	0.8	47
12	Characterization of the atypical Ppz/Hal3 phosphatase system from the pathogenic fungus <i>Cryptococcus neoformans</i> . <i>Molecular Microbiology</i> , 2019, 111, 898-917.	1.2	7
13	Ser/Thr protein phosphatases in fungi: structure, regulation and function. <i>Microbial Cell</i> , 2019, 6, 217-256.	1.4	54
14	Mutations at the hydrophobic core affect Hal3 trimer stability, reducing its Ppz1 inhibitory capacity but not its PPCDC moonlighting function. <i>Scientific Reports</i> , 2018, 8, 14701.	1.6	6
15	<i>Cryptococcus neoformans</i> can form titan-like cells in vitro in response to multiple signals. <i>PLoS Pathogens</i> , 2018, 14, e1007007.	2.1	98
16	Two NRAMP6 Isoforms Function as Iron and Manganese Transporters and Contribute to Disease Resistance in Rice. <i>Molecular Plant-Microbe Interactions</i> , 2017, 30, 385-398.	1.4	116
17	The inhibitory mechanism of Hal3 on the yeast Ppz1 phosphatase: A mutagenesis analysis. <i>Scientific Reports</i> , 2017, 7, 8819.	1.6	12
18	Improvement of biochemical methods of polyP quantification. <i>Microbial Cell</i> , 2017, 4, 6-15.	1.4	41

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19	The <i>Saccharomyces cerevisiae</i> Ptc1 protein phosphatase attenuates G2M cell cycle blockage caused by activation of the cell wall integrity pathway. <i>Molecular Microbiology</i> , 2016, 101, 671-687.	1.2	4
20	The effector AWR5 from the plant pathogen <i>Ralstonia solanacearum</i> is an inhibitor of the TOR signalling pathway. <i>Scientific Reports</i> , 2016, 6, 27058.	1.6	61
21	Genome-wide recruitment profiling of transcription factor Crz1 in response to high pH stress. <i>BMC Genomics</i> , 2016, 17, 662.	1.2	25
22	The yeast Aft2 transcription factor determines selenite toxicity by controlling the low affinity phosphate transport system. <i>Scientific Reports</i> , 2016, 6, 32836.	1.6	22
23	Polyphosphate is involved in cell cycle progression and genomic stability in <i>Saccharomyces cerevisiae</i> . <i>Molecular Microbiology</i> , 2016, 101, 367-380.	1.2	58
24	Improving Ribosomal RNA Integrity in Surgically Resected Human Brain Tumor Biopsies. <i>Biopreservation and Biobanking</i> , 2016, 14, 156-164.	0.5	6
25	Wide-Ranging Effects of the Yeast Ptc1 Protein Phosphatase Acting Through the MAPK Kinase Mkk1. <i>Genetics</i> , 2016, 202, 141-156.	1.2	24
26	Interactions Between Monovalent Cations and Nutrient Homeostasis. <i>Advances in Experimental Medicine and Biology</i> , 2016, 892, 271-289.	0.8	4
27	Regulation of the Na ⁺ /K ⁺ -ATPase Ena1 Expression by Calcineurin/Crz1 under High pH Stress: A Quantitative Study. <i>PLoS ONE</i> , 2016, 11, e0158424.	1.1	19
28	Analysis of Two Putative <i>Candida albicans</i> Phosphopantothencysteine Decarboxylase / Protein Phosphatase Z Regulatory Subunits Reveals an Unexpected Distribution of Functional Roles. <i>PLoS ONE</i> , 2016, 11, e0160965.	1.1	11
29	Complex stability and dynamic subunit interchange modulates the disparate activities of the yeast moonlighting proteins Hal3 and Vhs3. <i>Scientific Reports</i> , 2015, 5, 15774.	1.6	18
30	Coordinate responses to alkaline pH stress in budding yeast. <i>Microbial Cell</i> , 2015, 2, 182-196.	1.4	68
31	Robustness of Equations that Define Molecular Subtypes of Glioblastoma Tumors Based on Five Transcripts Measured by RT-PCR. <i>OMICS A Journal of Integrative Biology</i> , 2015, 19, 41-51.	1.0	2
32	Impact of high pH stress on yeast gene expression: A comprehensive analysis of mRNA turnover during stress responses. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2015, 1849, 653-664.	0.9	44
33	The Cytosolic pH of Individual <i>Saccharomyces cerevisiae</i> Cells Is a Key Factor in Acetic Acid Tolerance. <i>Applied and Environmental Microbiology</i> , 2015, 81, 7813-7821.	1.4	34
34	Functional interactions between potassium and phosphate homeostasis in <i>Saccharomyces cerevisiae</i> . <i>Molecular Microbiology</i> , 2015, 95, 555-572.	1.2	27
35	Systems Biology of Monovalent Cation Homeostasis in Yeast. <i>Advances in Microbial Physiology</i> , 2014, 64, 1-63.	1.0	18
36	Coregulated Expression of the Na ⁺ /Phosphate Pho89 Transporter and Ena1 Na ⁺ -ATPase Allows Their Functional Coupling under High-pH Stress. <i>Molecular and Cellular Biology</i> , 2014, 34, 4420-4435.	1.1	39

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37	Assessing Differential Expression Measurements by Highly Parallel Pyrosequencing and DNA Microarrays: A Comparative Study. <i>OMICS A Journal of Integrative Biology</i> , 2013, 17, 53-59.	1.0	2
38	The <i>Schizosaccharomyces pombe</i> fusion gene <i>hal3</i> encodes three distinct activities. <i>Molecular Microbiology</i> , 2013, 90, 367-382.	1.2	11
39	Molecular analysis of a conditional <i>hal3 vhs3</i> yeast mutant links potassium homeostasis with flocculation and invasiveness. <i>Fungal Genetics and Biology</i> , 2013, 53, 1-9.	0.9	9
40	Ptc6 Is Required for Proper Rapamycin-Induced Down-Regulation of the Genes Coding for Ribosomal and rRNA Processing Proteins in <i>S. cerevisiae</i> . <i>PLoS ONE</i> , 2013, 8, e64470.	1.1	19
41	Potassium Starvation in Yeast: Mechanisms of Homeostasis Revealed by Mathematical Modeling. <i>PLoS Computational Biology</i> , 2012, 8, e1002548.	1.5	37
42	Protein phosphatase CaPpz1 is involved in cation homeostasis, cell wall integrity and virulence of <i>Candida albicans</i> . <i>Microbiology (United Kingdom)</i> , 2012, 158, 1258-1267.	0.7	34
43	Development of robust discriminant equations for assessing subtypes of glioblastoma biopsies. <i>British Journal of Cancer</i> , 2012, 106, 1816-1825.	2.9	8
44	Modulation of Yeast Alkaline Cation Tolerance by Ypi1 Requires Calcineurin. <i>Genetics</i> , 2012, 190, 1355-1364.	1.2	14
45	Functional mapping of the disparate activities of the yeast moonlighting protein Hal3. <i>Biochemical Journal</i> , 2012, 442, 357-368.	1.7	28
46	The role of the Snf1 kinase in the adaptive response of <i>Saccharomyces cerevisiae</i> to alkaline pH stress. <i>Biochemical Journal</i> , 2012, 444, 39-49.	1.7	54
47	Protein phosphatase Z modulates oxidative stress response in fungi. <i>Fungal Genetics and Biology</i> , 2012, 49, 708-716.	0.9	26
48	Inhibition of human calcineurin and yeast calcineurin-dependent gene expression by <i>Jasminum humile</i> leaf and root extracts. <i>Journal of Ethnopharmacology</i> , 2012, 140, 293-297.	2.0	5
49	The short-term response of yeast to potassium starvation. <i>Environmental Microbiology</i> , 2012, 14, 3026-3042.	1.8	27
50	Lack of the Glc7 phosphatase regulatory subunit Ypi1 activates the morphogenetic checkpoint. <i>International Journal of Biochemistry and Cell Biology</i> , 2012, 44, 1862-1871.	1.2	6
51	Adaptation to potassium starvation of wild-type and K ⁺ transport mutant (<i>trk1,2</i>) of <i>Saccharomyces cerevisiae</i> : 2D-dimensional gel electrophoresis-based proteomic approach. <i>MicrobiologyOpen</i> , 2012, 1, 182-193.	1.2	7
52	The wheat MAP kinase phosphatase 1 confers higher lithium tolerance in yeast. <i>FEMS Yeast Research</i> , 2012, 12, 774-784.	1.1	8
53	The role of the protein kinase A pathway in the response to alkaline pH stress in yeast. <i>Biochemical Journal</i> , 2011, 438, 523-533.	1.7	36
54	Alkali-metal-cation influx and efflux systems in nonconventional yeast species. <i>FEMS Microbiology Letters</i> , 2011, 317, 1-8.	0.7	58

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55	Central nervous system gene expression changes in a transgenic mouse model for bovine spongiform encephalopathy. <i>Veterinary Research</i> , 2011, 42, 109.	1.1	10
56	Aequorin-expressing yeast emits light under electric control. <i>Journal of Biotechnology</i> , 2011, 152, 93-95.	1.9	4
57	A Genomewide Screen for Tolerance to Cationic Drugs Reveals Genes Important for Potassium Homeostasis in <i>Saccharomyces cerevisiae</i> . <i>Eukaryotic Cell</i> , 2011, 10, 1241-1250.	3.4	53
58	Type 2C Protein Phosphatases in Fungi. <i>Eukaryotic Cell</i> , 2011, 10, 21-33.	3.4	56
59	Ref2, a regulatory subunit of the yeast protein phosphatase 1, is a novel component of cation homeostasis. <i>Biochemical Journal</i> , 2010, 426, 355-364.	1.7	13
60	Lack of DNA helicase Pif1 disrupts zinc and iron homeostasis in yeast. <i>Biochemical Journal</i> , 2010, 432, 595-608.	1.7	6
61	Regulation of Trk-dependent potassium transport by the calcineurin pathway involves the Hal5 kinase. <i>FEBS Letters</i> , 2010, 584, 2415-2420.	1.3	26
62	Lack of main K ⁺ uptake systems in <i>Saccharomyces cerevisiae</i> cells affects yeast performance in both potassium-sufficient and potassium-limiting conditions. <i>FEMS Yeast Research</i> , 2010, 10, no-no.	1.1	88
63	Development of a Predictor for Human Brain Tumors Based on Gene Expression Values Obtained from Two Types of Microarray Technologies. <i>OMICS A Journal of Integrative Biology</i> , 2010, 14, 157-164.	1.0	12
64	Hal4 and Hal5 Protein Kinases Are Required for General Control of Carbon and Nitrogen Uptake and Metabolism. <i>Eukaryotic Cell</i> , 2010, 9, 1881-1890.	3.4	25
65	Integrative Responses to High pH Stress in <i>S. cerevisiae</i> . <i>OMICS A Journal of Integrative Biology</i> , 2010, 14, 517-523.	1.0	45
66	Alkali Metal Cation Transport and Homeostasis in Yeasts. <i>Microbiology and Molecular Biology Reviews</i> , 2010, 74, 95-120.	2.9	245
67	Genome-Wide Analysis of Factors Affecting Transcription Elongation and DNA Repair: A New Role for PAF and Ccr4-Not in Transcription-Coupled Repair. <i>PLoS Genetics</i> , 2009, 5, e1000364.	1.5	81
68	Automated Brain Tumor Biopsy Prediction Using Single-labeling cDNA Microarrays-based Gene Expression Profiling. <i>Diagnostic Molecular Pathology</i> , 2009, 18, 206-218.	2.1	17
69	Moonlighting proteins Hal3 and Vhs3 form a heteromeric PPCDC with Ykl088w in yeast CoA biosynthesis. <i>Nature Chemical Biology</i> , 2009, 5, 920-928.	3.9	53
70	Normal Function of the Yeast TOR Pathway Requires the Type 2C Protein Phosphatase Ptc1. <i>Molecular and Cellular Biology</i> , 2009, 29, 2876-2888.	1.1	38
71	Use of the TRP1 auxotrophic marker for gene disruption and phenotypic analysis in yeast: a note of warning. <i>FEMS Yeast Research</i> , 2008, 8, 2-5.	1.1	21
72	A role for protein kinase CK2 in plant development: evidence obtained using a dominant-negative mutant. <i>Plant Journal</i> , 2008, 55, 118-130.	2.8	60

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73	Direct Regulation of Genes Involved in Glucose Utilization by the Calcium/Calcineurin Pathway. <i>Journal of Biological Chemistry</i> , 2008, 283, 13923-13933.	1.6	52
74	Use of Yeast Genetic Tools to Define Biological Roles of Novel Protein Phosphatases. , 2007, 365, 299-308.		0
75	YPI1 and SDS22 Proteins Regulate the Nuclear Localization and Function of Yeast Type 1 Phosphatase Glc7. <i>Journal of Biological Chemistry</i> , 2007, 282, 3282-3292.	1.6	50
76	Gcn2p Regulates a G1/S Cell Cycle Checkpoint in Response to DNA Damage. <i>Cell Cycle</i> , 2007, 6, 2302-2305.	1.3	23
77	Function and Regulation of the <i>Saccharomyces cerevisiae</i> ENA Sodium ATPase System. <i>Eukaryotic Cell</i> , 2007, 6, 2175-2183.	3.4	105
78	The HAL3-PPZ1 dependent regulation of nonsense suppression efficiency in yeast and its influence on manifestation of the yeast prion-like determinant [ISP+]. <i>Genes To Cells</i> , 2007, 12, 435-445.	0.5	13
79	Disruption of iron homeostasis in <i>Saccharomyces cerevisiae</i> by high zinc levels: a genome-wide study. <i>Molecular Microbiology</i> , 2007, 65, 521-537.	1.2	96
80	Genomics and Metabolomics Research for Brain Tumour Diagnosis Based on Machine Learning. <i>Lecture Notes in Computer Science</i> , 2007, , 1012-1019.	1.0	3
81	Role of protein phosphatases 2C on tolerance to lithium toxicity in the yeast <i>Saccharomyces cerevisiae</i> . <i>Molecular Microbiology</i> , 2006, 62, 263-277.	1.2	44
82	Transcriptional Profiling of the Protein Phosphatase 2C Family in Yeast Provides Insights into the Unique Functional Roles of Ptc1. <i>Journal of Biological Chemistry</i> , 2006, 281, 35057-35069.	1.6	59
83	Heterologous Expression Implicates a GATA Factor in Regulation of Nitrogen Metabolic Genes and Ion Homeostasis in the Halotolerant Yeast <i>Debaryomyces hansenii</i> . <i>Eukaryotic Cell</i> , 2006, 5, 1388-1398.	3.4	18
84	Signaling Alkaline pH Stress in the Yeast <i>Saccharomyces cerevisiae</i> through the Wsc1 Cell Surface Sensor and the Slit2 MAPK Pathway. <i>Journal of Biological Chemistry</i> , 2006, 281, 39785-39795.	1.6	107
85	The Transcriptional Response of the Yeast Na ⁺ -ATPase ENA1 Gene to Alkaline Stress Involves Three Main Signaling Pathways*. <i>Journal of Biological Chemistry</i> , 2006, 281, 36632-36642.	1.6	80
86	Copper and Iron Are the Limiting Factors for Growth of the Yeast <i>Saccharomyces cerevisiae</i> in an Alkaline Environment. <i>Journal of Biological Chemistry</i> , 2004, 279, 19698-19704.	1.6	118
87	Characterization of the Calcium-mediated Response to Alkaline Stress in <i>Saccharomyces cerevisiae</i> . <i>Journal of Biological Chemistry</i> , 2004, 279, 43614-43624.	1.6	180
88	Functional Characterization of the Yeast Ppz1 Phosphatase Inhibitory Subunit Hal3. <i>Journal of Biological Chemistry</i> , 2004, 279, 42619-42627.	1.6	32
89	Functional Characterization of the <i>Saccharomyces cerevisiae</i> VHS3 Gene. <i>Journal of Biological Chemistry</i> , 2004, 279, 34421-34430.	1.6	45
90	The Ppz protein phosphatases regulate Trk-independent potassium influx in yeast. <i>FEBS Letters</i> , 2004, 578, 58-62.	1.3	19

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91	A new set of DNA macrochips for the yeast <i>Saccharomyces cerevisiae</i> : features and uses. <i>International Microbiology</i> , 2004, 7, 199-206.	1.1	56
92	Identification of multicopy suppressors of cell cycle arrest at the G1-S transition in <i>Saccharomyces cerevisiae</i> . <i>Yeast</i> , 2003, 20, 157-169.	0.8	40
93	Mutagenesis analysis of the yeast Nha1 Na ⁺ /H ⁺ antiporter carboxy-terminal tail reveals residues required for function in cell cycle. <i>FEBS Letters</i> , 2003, 545, 239-245.	1.3	25
94	Use of Tetracycline-Regulatable Promoters for Functional Analysis of Protein Phosphatases in Yeast. <i>Methods in Enzymology</i> , 2003, 366, 347-358.	0.4	4
95	Molecular Characterization of Ypi1, a Novel <i>Saccharomyces cerevisiae</i> Type 1 Protein Phosphatase Inhibitor. <i>Journal of Biological Chemistry</i> , 2003, 278, 47744-47752.	1.6	69
96	Regulation of ENA1 Na ⁺ -ATPase Gene Expression by the Ppz1 Protein Phosphatase Is Mediated by the Calcineurin Pathway. <i>Eukaryotic Cell</i> , 2003, 2, 937-948.	3.4	68
97	Structural Model of a Malonyl-CoA-binding Site of Carnitine Octanoyltransferase and Carnitine Palmitoyltransferase I. <i>Journal of Biological Chemistry</i> , 2002, 277, 11473-11480.	1.6	38
98	Sit4 Is Required for Proper Modulation of the Biological Functions Mediated by Pkc1 and the Cell Integrity Pathway in <i>Saccharomyces cerevisiae</i> . <i>Journal of Biological Chemistry</i> , 2002, 277, 33468-33476.	1.6	64
99	Degradation of transgenic DNA from genetically modified soya and maize in human intestinal simulations. <i>British Journal of Nutrition</i> , 2002, 87, 533-542.	1.2	49
100	Novel protein phosphatases in yeast. <i>FEBS Journal</i> , 2002, 269, 1072-1077.	0.2	37
101	The transcriptional response to alkaline pH in <i>Saccharomyces cerevisiae</i> : evidence for calcium-mediated signalling. <i>Molecular Microbiology</i> , 2002, 46, 1319-1333.	1.2	174
102	The Ppz protein phosphatases are key regulators of K ⁺ and pH homeostasis: implications for salt tolerance, cell wall integrity and cell cycle progression. <i>EMBO Journal</i> , 2002, 21, 920-929.	3.5	125
103	Degradation of transgenic DNA from genetically modified soya and maize in human intestinal simulations. <i>British Journal of Nutrition</i> , 2002, 87, 533-42.	1.2	10
104	Maize protein kinase CK2: regulation and functionality of three \hat{I}^2 regulatory subunits. <i>Plant Journal</i> , 2001, 25, 365-374.	2.8	46
105	Functional analysis of the <i>Neurospora crassa</i> PZL-1 protein phosphatase by expression in budding and fission yeast. <i>Yeast</i> , 2001, 18, 115-124.	0.8	24
106	Regulation of the Sko1 transcriptional repressor by the Hog1 MAP kinase in response to osmotic stress. <i>EMBO Journal</i> , 2001, 20, 1123-1133.	3.5	188
107	A Role for the Ppz Ser/Thr Protein Phosphatases in the Regulation of Translation Elongation Factor 1B \hat{I} . <i>Journal of Biological Chemistry</i> , 2001, 276, 14829-14834.	1.6	30
108	A Screening for High Copy Suppressors of the sit4 hal3 Synthetically Lethal Phenotype Reveals a Role for the Yeast Nha1 Antiporter in Cell Cycle Regulation. <i>Journal of Biological Chemistry</i> , 2001, 276, 29740-29747.	1.6	47

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109	Inhibition by etomoxir of rat liver carnitine octanoyltransferase is produced through the co-ordinate interaction with two histidine residues. <i>Biochemical Journal</i> , 2000, 351, 495.	1.7	7
110	Inhibition by etomoxir of rat liver carnitine octanoyltransferase is produced through the co-ordinate interaction with two histidine residues. <i>Biochemical Journal</i> , 2000, 351, 495-502.	1.7	12
111	Protein phosphatase 2A holoenzyme and its subunits from <i>Medicago sativa</i> . <i>Plant Molecular Biology</i> , 2000, 43, 527-536.	2.0	20
112	The <i>Arabidopsis thaliana</i> PPX/PP4 phosphatases: molecular cloning and structural organization of the genes and immunolocalization of the proteins to plastids. <i>Plant Molecular Biology</i> , 2000, 44, 499-511.	2.0	15
113	Trk1 and Trk2 Define the Major K ⁺ Transport System in Fission Yeast. <i>Journal of Bacteriology</i> , 2000, 182, 394-399.	1.0	45
114	The Transcriptional Response of Yeast to Saline Stress. <i>Journal of Biological Chemistry</i> , 2000, 275, 17249-17255.	1.6	353
115	Rck2 Kinase Is a Substrate for the Osmotic Stress-Activated Mitogen-Activated Protein Kinase Hog1. <i>Molecular and Cellular Biology</i> , 2000, 20, 3887-3895.	1.1	132
116	Identification of the two histidine residues responsible for the inhibition by malonyl-CoA in peroxisomal carnitine octanoyltransferase from rat liver. <i>FEBS Letters</i> , 2000, 466, 183-186.	1.3	17
117	The Yeast Ser/Thr Phosphatases Sit4 and Ppz1 Play Opposite Roles in Regulation of the Cell Cycle. <i>Molecular and Cellular Biology</i> , 1999, 19, 2408-2415.	1.1	78
118	The <i>Schizosaccharomyces pombe</i> Pzh1 protein phosphatase regulates Na ⁺ ion influx in a Trk1-independent fashion. <i>FEBS Journal</i> , 1999, 260, 31-37.	0.2	16
119	Disruption and phenotypic analysis of seven ORFs from the left arm of chromosome XV of <i>Saccharomyces cerevisiae</i> . , 1999, 15, 435-441.		13
120	Biochemical and Genetic Analyses of the Role of Yeast Casein Kinase 2 in Salt Tolerance. <i>Journal of Bacteriology</i> , 1999, 181, 6456-6462.	1.0	31
121	Molecular cloning and characterization of two phosphatase 2A catalytic subunit genes from <i>Arabidopsis thaliana</i> . <i>Gene</i> , 1998, 209, 105-112.	1.0	21
122	Role of UEV-1A, a homologue of the tumor suppressor protein TSG101, in protection from DNA damage. <i>FEBS Letters</i> , 1998, 423, 49-52.	1.3	16
123	The Pzh1 protein phosphatase and the Spm1 protein kinase are involved in the regulation of the plasma membrane H ⁺ -ATPase in fission yeast. <i>FEBS Letters</i> , 1998, 435, 241-244.	1.3	11
124	Protein Phosphatase 2A and Protein Phosphatase X Genes in <i>Arabidopsis thaliana</i> . , 1998, 93, 201-212.		2
125	The Search for the Biological Function of Novel Yeast Ser/Thr Phosphatases. , 1998, 93, 305-313.		2
126	The yeast halotolerance determinant Hal3p is an inhibitory subunit of the Ppz1p Ser/Thr protein phosphatase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1998, 95, 7357-7362.	3.3	106

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127	Regulation of Salt Tolerance in Fission Yeast by a Protein-Phosphatase-Z-Like Ser/Thr Protein Phosphatase. <i>FEBS Journal</i> , 1997, 250, 476-483.	0.2	29
128	Analysis of the DNA sequence of a 15,500 bp fragment near the left telomere of chromosome XV from <i>Saccharomyces cerevisiae</i> reveals a putative sugar transporter, a carboxypeptidase homologue and two new open reading frames. , 1996, 12, 709-714.		4
129	Sequence analysis of a 13.4 kbp fragment from the left arm of chromosome XV reveals a malate dehydrogenase gene, a putative Ser/Thr protein kinase, the ribosomal L25 gene and four new open reading frames. <i>Yeast</i> , 1996, 12, 1013-1020.	0.8	6
130	Sequence analysis of a 12 801 bp fragment of the left arm of yeast chromosome XV containing a putative 6-phosphofructo-2-kinase gene, a gene for a possible glycopospholipid-anchored surface protein and six other open reading frames. <i>Yeast</i> , 1996, 12, 1053-1058.	0.8	4
131	The NH ₂ -terminal Extension of Protein Phosphatase PPZ1 Has an Essential Functional Role. <i>Journal of Biological Chemistry</i> , 1996, 271, 26349-26355.	1.6	59
132	XV. Yeast sequencing reports. Sequence analysis of a 9873 bp fragment of the left arm of yeast chromosome XV that contains the ARG8 and CDC33 genes, a putative riboflavin synthase beta chain gene, and four new open reading frames. <i>Yeast</i> , 1995, 11, 1061-1067.	0.8	6
133	XV. Yeast sequencing reports. DNA sequence analysis of a 13 kbp fragment of the left arm of yeast chromosome XV containing seven new open reading frames. <i>Yeast</i> , 1995, 11, 1281-1288.	0.8	14
134	The PPZ Protein Phosphatases Are Important Determinants of Salt Tolerance in Yeast Cells. <i>Journal of Biological Chemistry</i> , 1995, 270, 13036-13041.	1.6	138
135	Biochemical characterization of recombinant yeast PPZ1, a protein phosphatase involved in salt tolerance. <i>FEBS Letters</i> , 1995, 368, 39-44.	1.3	28
136	Molecular characterization of a fourth isoform of the catalytic subunit of protein phosphatase 2A from <i>Arabidopsis thaliana</i> . <i>Plant Molecular Biology</i> , 1994, 26, 523-528.	2.0	39
137	Effects of Glucose on the Activation and Translocation of Glycogen Synthase in Diabetic Rat Hepatocytes. <i>FEBS Journal</i> , 1994, 226, 665-671.	0.2	23
138	Protein phosphatases in higher plants: multiplicity of type 2A phosphatases in <i>Arabidopsis thaliana</i> . <i>Plant Molecular Biology</i> , 1993, 21, 475-485.	2.0	75
139	Identification and molecular cloning of two homologues of protein phosphatase X from <i>Arabidopsis thaliana</i> . <i>Plant Molecular Biology</i> , 1993, 23, 1177-1185.	2.0	24
140	The PPZ protein phosphatases are involved in the maintenance of osmotic stability of yeast cells. <i>FEBS Letters</i> , 1993, 318, 282-286.	1.3	87
141	Glycogen metabolism in a <i>Saccharomyces cerevisiae</i> phosphoglucose isomerase (pgi1) disruption mutant. <i>FEBS Letters</i> , 1992, 310, 182-186.	1.3	18
142	<i>Saccharomyces cerevisiae</i> gene SIT4 is involved in the control of glycogen metabolism. <i>FEBS Letters</i> , 1991, 279, 341-345.	1.3	36
143	Glycogen hyperaccumulation in <i>Saccharomyces cerevisiae</i> ras2 mutant A biochemical study. <i>FEBS Letters</i> , 1991, 290, 38-42.	1.3	9
144	The gene DIS2S1 is essential in <i>Saccharomyces cerevisiae</i> and is involved in glycogen phosphorylase activation. <i>Current Genetics</i> , 1991, 19, 339-342.	0.8	47

#	ARTICLE	IF	CITATIONS
145	Nucleotide sequence of a rat heart cDNA encoding the isotype \hat{I}^2 of the catalytic subunit of protein phosphatase 2A. <i>Nucleic Acids Research</i> , 1989, 17, 8370-8370.	6.5	4
146	Nucleotide sequence of a rat heart cDNA encoding the isotype \hat{I}^1 of the catalytic subunit of protein phosphatase 2A. <i>Nucleic Acids Research</i> , 1989, 17, 8369-8369.	6.5	3
147	Control of glycogen synthase and phosphorylase in hepatocytes from diabetic rats. Effects of glucagon, vasopressin, and vanadate. <i>Diabetes</i> , 1989, 38, 793-798.	0.3	4
148	Human liver phosphatase 2A: cDNA and amino acid sequence of two catalytic subunit isotypes.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1988, 85, 4252-4256.	3.3	137
149	Sequence Homologies between Type 1 and Type 2A Protein Phosphatases. <i>Molecular Endocrinology</i> , 1987, 1, 745-748.	3.7	14
150	Phosphorylation and inactivation of rat hepatocyte glycogen synthase by phorbol esters and mezerein. <i>Biochemical and Biophysical Research Communications</i> , 1986, 134, 113-119.	1.0	22
151	Benzomorphan-related compounds. Part 21. Synthesis of 7,8-benzomorphans via 2-aryl-4-piperidones. <i>Journal of the Chemical Society Perkin Transactions 1</i> , 1986, , 1533-1539.	0.9	16
152	Threonine phosphorylation of rat liver glycogen synthase. <i>Biochemical and Biophysical Research Communications</i> , 1985, 130, 987-993.	1.0	7
153	Hormonal effects on the phosphorylation of glycogen synthase in rat hepatocytes. <i>FEBS Letters</i> , 1984, 170, 310-314.	1.3	17
154	Functional mapping of the N-terminal region of the yeast moonlighting protein Sis2/Hal3 reveals crucial residues for Ppz1 regulation. <i>FEBS Journal</i> , 0, , .	2.2	1