

# Markus J Tamas

## List of Publications by Year in descending order

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58  
papers

3,886  
citations

147566

31  
h-index

168136

53  
g-index

64  
all docs

64  
docs citations

64  
times ranked

4247  
citing authors

#	ARTICLE	IF	CITATIONS
1	A subgroup of plant aquaporins facilitate the bi-directional diffusion of As(OH) <sub>3</sub> and Sb(OH) <sub>3</sub> across membranes. <i>BMC Biology</i> , 2008, 6, 26.	1.7	379
2	Fps1p controls the accumulation and release of the compatible solute glycerol in yeast osmoregulation. <i>Molecular Microbiology</i> , 1999, 31, 1087-1104.	1.2	357
3	Heavy Metals and Metalloids As a Cause for Protein Misfolding and Aggregation. <i>Biomolecules</i> , 2014, 4, 252-267.	1.8	316
4	The glycerol channel Fps1p mediates the uptake of arsenite and antimonite in <i>Saccharomyces cerevisiae</i> . <i>Molecular Microbiology</i> , 2001, 40, 1391-1401.	1.2	306
5	How <i>Saccharomyces cerevisiae</i> copes with toxic metals and metalloids. <i>FEMS Microbiology Reviews</i> , 2010, 34, 925-951.	3.9	254
6	The MAPK Hog1p Modulates Fps1p-dependent Arsenite Uptake and Tolerance in Yeast. <i>Molecular Biology of the Cell</i> , 2006, 17, 4400-4410.	0.9	177
7	The <i>Saccharomyces cerevisiae</i> Sko1p transcription factor mediates HOG pathway-dependent osmotic regulation of a set of genes encoding enzymes implicated in protection from oxidative damage. <i>Molecular Microbiology</i> , 2001, 40, 1067-1083.	1.2	161
8	Arsenite interferes with protein folding and triggers formation of protein aggregates in yeast. <i>Journal of Cell Science</i> , 2012, 125, 5073-83.	1.2	121
9	Distinct stress conditions result in aggregation of proteins with similar properties. <i>Scientific Reports</i> , 2016, 6, 24554.	1.6	117
10	Quantitative transcriptome, proteome, and sulfur metabolite profiling of the <i>Saccharomyces cerevisiae</i> response to arsenite. <i>Physiological Genomics</i> , 2007, 30, 35-43.	1.0	109
11	Genetic basis of arsenite and cadmium tolerance in <i>Saccharomyces cerevisiae</i> . <i>BMC Genomics</i> , 2009, 10, 105.	1.2	100
12	A Short Regulatory Domain Restricts Glycerol Transport through Yeast Fps1p. <i>Journal of Biological Chemistry</i> , 2003, 278, 6337-6345.	1.6	87
13	Transcriptional Activation of Metalloid Tolerance Genes in <i>Saccharomyces cerevisiae</i> Requires the AP-1-like Proteins Yap1p and Yap8p. <i>Molecular Biology of the Cell</i> , 2004, 15, 2049-2060.	0.9	84
14	Stimulation of the yeast high osmolarity glycerol (HOG) pathway: evidence for a signal generated by a change in turgor rather than by water stress. <i>FEBS Letters</i> , 2000, 472, 159-165.	1.3	81
15	Mechanisms involved in metalloid transport and tolerance acquisition. <i>Current Genetics</i> , 2001, 40, 2-12.	0.8	65
16	Yeast reveals unexpected roles and regulatory features of aquaporins and aquaglyceroporins. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2014, 1840, 1482-1491.	1.1	59
17	Molecular and physiological characterization of the NAD-dependent glycerol 3-phosphate dehydrogenase in the filamentous fungus <i>Aspergillus nidulans</i> . <i>Molecular Microbiology</i> , 2001, 39, 145-157.	1.2	58
18	Determination of primary sequence specificity of <i>Arabidopsis</i> MAPKs MPK3 and MPK6 leads to identification of new substrates. <i>Biochemical Journal</i> , 2012, 446, 271-278.	1.7	58

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19	Cadmium Causes Misfolding and Aggregation of Cytosolic Proteins in Yeast. <i>Molecular and Cellular Biology</i> , 2017, 37, .	1.1	58
20	A Regulatory Domain in the C-terminal Extension of the Yeast Glycerol Channel Fps1p. <i>Journal of Biological Chemistry</i> , 2004, 279, 14954-14960.	1.6	54
21	Evolutionary Forces Act on Promoter Length: Identification of Enriched Cis-Regulatory Elements. <i>Molecular Biology and Evolution</i> , 2009, 26, 1299-1307.	3.5	53
22	Modulation of <i>Leishmania major</i> aquaglyceroporin activity by a mitogen-activated protein kinase. <i>Molecular Microbiology</i> , 2012, 85, 1204-1218.	1.2	52
23	Misfolding and aggregation of nascent proteins: a novel mode of toxic cadmium action in vivo. <i>Current Genetics</i> , 2018, 64, 177-181.	0.8	52
24	Metalloid tolerance based on phytochelatins is not functionally equivalent to the arsenite transporter Acr3p. <i>Biochemical and Biophysical Research Communications</i> , 2003, 304, 293-300.	1.0	51
25	Glutathione serves an extracellular defence function to decrease arsenite accumulation and toxicity in yeast. <i>Molecular Microbiology</i> , 2012, 84, 1177-1188.	1.2	48
26	Amplification of the CUP1 gene is associated with evolution of copper tolerance in <i>Saccharomyces cerevisiae</i> . <i>Microbiology (United Kingdom)</i> , 2012, 158, 2325-2335.	0.7	47
27	Arsenic Transport in Prokaryotes and Eukaryotic Microbes. <i>Advances in Experimental Medicine and Biology</i> , 2010, 679, 47-55.	0.8	44
28	Arsenic Directly Binds to and Activates the Yeast AP-1-Like Transcription Factor Yap8. <i>Molecular and Cellular Biology</i> , 2016, 36, 913-922.	1.1	42
29	Regulation of the arsenic-responsive transcription factor Yap8p involves the ubiquitin-proteasome pathway. <i>Journal of Cell Science</i> , 2007, 120, 256-264.	1.2	37
30	Global analysis of protein aggregation in yeast during physiological conditions and arsenite stress. <i>Biology Open</i> , 2014, 3, 913-923.	0.6	36
31	Characterization of the DNA-binding motif of the arsenic-responsive transcription factor Yap8p. <i>Biochemical Journal</i> , 2008, 415, 467-475.	1.7	35
32	Identification of residues controlling transport through the yeast aquaglyceroporin Fps1 using a genetic screen. <i>FEBS Journal</i> , 2004, 271, 771-779.	0.2	32
33	Probing Conserved Regions of the Cytoplasmic LOOP1 Segment Linking Transmembrane Segments 2 and 3 of the <i>Saccharomyces cerevisiae</i> Plasma Membrane H <sup>+</sup> -ATPase. <i>Journal of Biological Chemistry</i> , 1996, 271, 25438-25445.	1.6	28
34	Nuclear envelope budding is a response to cellular stress. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	28
35	The osmotic stress response of <i>Saccharomyces cerevisiae</i> . , 2003, , 121-200.		27
36	Mechanisms of toxic metal tolerance in yeast. <i>Topics in Current Genetics</i> , 2005, , 395-454.	0.7	27

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37	Mitogen-Activated Protein Kinase Hog1 Mediates Adaptation to G <sub>1</sub> Checkpoint Arrest during Arsenite and Hyperosmotic Stress. <i>Eukaryotic Cell</i> , 2008, 7, 1309-1317.	3.4	27
38	Design, Synthesis, and Characterization of a Highly Effective Hog1 Inhibitor: A Powerful Tool for Analyzing MAP Kinase Signaling in Yeast. <i>PLoS ONE</i> , 2011, 6, e20012.	1.1	23
39	The mitogen-activated protein kinase Slt2 modulates arsenite transport through the aquaglyceroporin Fps1. <i>FEBS Letters</i> , 2016, 590, 3649-3659.	1.3	21
40	Sequence-specific dynamics of DNA response elements and their flanking sites regulate the recognition by AP-1 transcription factors. <i>Nucleic Acids Research</i> , 2021, 49, 9280-9293.	6.5	21
41	HwHog1 kinase activity is crucial for survival of <i>Hortaea werneckii</i> in extremely hyperosmolar environments. <i>Fungal Genetics and Biology</i> , 2015, 74, 45-58.	0.9	18
42	Mathematical modelling of arsenic transport, distribution and detoxification processes in yeast. <i>Molecular Microbiology</i> , 2014, 92, 1343-1356.	1.2	15
43	Yeast Aquaglyceroporins Use the Transmembrane Core to Restrict Glycerol Transport. <i>Journal of Biological Chemistry</i> , 2012, 287, 23562-23570.	1.6	14
44	Cellular and molecular mechanisms of antimony transport, toxicity and resistance. <i>Environmental Chemistry</i> , 2016, 13, 955.	0.7	13
45	Editorial: Molecular Mechanisms of Metalloid Transport, Toxicity and Tolerance. <i>Frontiers in Cell and Developmental Biology</i> , 2018, 6, 99.	1.8	13
46	<i>Saccharomyces cerevisiae</i> as a Model Organism for Elucidating Arsenic Tolerance Mechanisms. , 2011, , 87-112.		13
47	Effects of the Toxic Metals Arsenite and Cadmium on $\alpha$ -Synuclein Aggregation In Vitro and in Cells. <i>International Journal of Molecular Sciences</i> , 2021, 22, 11455.	1.8	13
48	Genome-wide imaging screen uncovers molecular determinants of arsenite-induced protein aggregation and toxicity. <i>Journal of Cell Science</i> , 2021, 134, .	1.2	11
49	Disentangling genetic and epigenetic determinants of ultrafast adaptation. <i>Molecular Systems Biology</i> , 2016, 12, 892.	3.2	9
50	Positional Scanning Peptide Libraries for Kinase Substrate Specificity Determinations: Straightforward and Reproducible Synthesis Using Pentafluorophenyl Esters. <i>ACS Combinatorial Science</i> , 2010, 12, 733-742.	3.3	8
51	Elucidating the response of <i>Kluyveromyces lactis</i> to arsenite and peroxide stress and the role of the transcription factor K1Yap8. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2014, 1839, 1295-1306.	0.9	8
52	The ancillary N-terminal region of the yeast AP-1 transcription factor Yap8 contributes to its DNA binding specificity. <i>Nucleic Acids Research</i> , 2020, 48, 5426-5441.	6.5	7
53	Application of a peptide-based assay to characterize inhibitors targeting protein kinases from yeast. <i>Current Genetics</i> , 2014, 60, 193-200.	0.8	6
54	Etp1 confers arsenite resistance by affecting <i>ACR3</i> expression. <i>FEMS Yeast Research</i> , 2022, , .	1.1	1

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55	Identification of novel arsenic resistance genes in yeast. <i>MicrobiologyOpen</i> , 2022, 11, .	1.2	1
56	An Investigation of the Possible Existence of Homologues of FPS1, a Glycerol Facilitator of <i>Saccharomyces Cerevisiae</i> , in the Osmotolerant Yeast <i>Zygosaccharomyces Rouxii</i> . , 2000, , 393-403.		0
57	Function and Regulation of the Yeast MIP Glycerol Export Channel Fps1p. , 2000, , 423-430.		0
58	Physical, genetic and functional interactions between the eisosome protein Pil1 and the MBOAT O-acyltransferase Gup1. <i>FEMS Yeast Research</i> , 2021, 21, .	1.1	0