Claudio De Virgilio

List of Publications by Year in descending order

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87	10,980	51 h-index	87
papers	citations		g-index
91	91	91	10487
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Global analysis of protein phosphorylation in yeast. Nature, 2005, 438, 679-684.	27.8	915
2	Sch9 Is a Major Target of TORC1 in Saccharomyces cerevisiae. Molecular Cell, 2007, 26, 663-674.	9.7	723
3	The septins: roles in cytokinesis and other processes. Current Opinion in Cell Biology, 1996, 8, 106-119.	5.4	455
4	The Vam6 GEF Controls TORC1 by Activating the EGO Complex. Molecular Cell, 2009, 35, 563-573.	9.7	398
5	Ste20-like protein kinases are required for normal localization of cell growth and for cytokinesis in budding yeast Genes and Development, 1995, 9, 1817-1830.	5.9	370
6	Deciphering Protein Kinase Specificity Through Large-Scale Analysis of Yeast Phosphorylation Site Motifs. Science Signaling, 2010, 3, ra12.	3.6	341
7	Leucyl-tRNA Synthetase Controls TORC1 via the EGO Complex. Molecular Cell, 2012, 46, 105-110.	9.7	308
8	The TOR and EGO Protein Complexes Orchestrate Microautophagy in Yeast. Molecular Cell, 2005, 19, 15-26.	9.7	305
9	A Septin-based Hierarchy of Proteins Required for Localized Deposition of Chitin in the Saccharomyces cerevisiae Cell Wall. Journal of Cell Biology, 1997, 139, 75-93.	5.2	301
10	The role of trehalose synthesis for the acquisition of thermotolerance in yeast. II. Physiological concentrations of trehalose increase the thermal stability of proteins in vitro. FEBS Journal, 1994, 219, 187-193.	0.2	295
11	The role of trehalose synthesis for the acquisition of thermotolerance in yeast. I. Genetic evidence that trehalose is a thermoprotectant. FEBS Journal, 1994, 219, 179-186.	0.2	279
12	TOR and PKA Signaling Pathways Converge on the Protein Kinase Rim15 to Control Entry into GO. Molecular Cell, 2003, 12, 1607-1613.	9.7	277
13	Role for the Rho-family GTPase Cdc42 in yeast mating-pheromone signal pathway. Nature, 1995, 376, 702-705.	27.8	251
14	Amino Acid Deprivation Inhibits TORC1 Through a GTPase-Activating Protein Complex for the Rag Family GTPase Gtr1. Science Signaling, 2013, 6, ra42.	3.6	237
15	Cell growth control: little eukaryotes make big contributions. Oncogene, 2006, 25, 6392-6415.	5.9	223
16	Disruption of TPS2, the gene encoding the 100-kDa subunit of the trehalose-6-phosphate synthase/phosphatase complex in Saccharomyces cerevisiae, causes accumulation of trehalose-6-phosphate and loss of trehalose-6-phosphate phosphatase activity. FEBS Journal, 1993, 212, 315-323.	0.2	213
17	<i>Saccharomyces cerevisiae</i> cAMP-dependent protein kinase controls entry into stationary phase through the Rim15p protein kinase. Genes and Development, 1998, 12, 2943-2955.	5.9	197
18	The TOR signalling network from yeast to man. International Journal of Biochemistry and Cell Biology, 2006, 38, 1476-1481.	2.8	194

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19	Composition and Functional Analysis of the Saccharomyces cerevisiae Trehalose Synthase Complex. Journal of Biological Chemistry, 1998, 273, 33311-33319.	3.4	189
20	Life in the midst of scarcity: adaptations to nutrient availability in Saccharomyces cerevisiae. Current Genetics, 2010, 56, 1-32.	1.7	189
21	The essence of yeast quiescence. FEMS Microbiology Reviews, 2012, 36, 306-339.	8.6	189
22	Caffeine extends yeast lifespan by targeting TORC1. Molecular Microbiology, 2008, 69, 277-285.	2.5	186
23	Establishment of Cell Polarity in Yeast. Cold Spring Harbor Symposia on Quantitative Biology, 1995, 60, 729-744.	1.1	185
24	PKA and Sch9 control a molecular switch important for the proper adaptation to nutrient availability. Molecular Microbiology, 2005, 55, 862-880.	2.5	170
25	The Novel Yeast PAS Kinase Rim15 Orchestrates GO-Associated Antioxidant Defense Mechanisms. Cell Cycle, 2004, 3, 460-466.	2.6	154
26	Amino Acids Stimulate TORC1 through Lst4-Lst7, a GTPase-Activating Protein Complex for the Rag Family GTPase Gtr2. Cell Reports, 2015, 13, 1-7.	6.4	145
27	Regulation of GO entry by the Pho80–Pho85 cyclin–CDK complex. EMBO Journal, 2005, 24, 4271-4278.	7.8	135
28	Rim 15 and the crossroads of nutrient signalling pathways in Saccharomyces cerevisiae. Cell Division, 2006, $1,3$.	2.4	129
29	Functional mapping of yeast genomes by saturated transposition. ELife, 2017, 6, .	6.0	126
30	Mutation of RGA1, which encodes a putative GTPase-activating protein for the polarity-establishment protein Cdc42p, activates the pheromone-response pathway in the yeast Saccharomyces cerevisiae Genes and Development, 1995, 9, 2949-2963.	5.9	124
31	Initiation of the TORC1-Regulated GO Program Requires Igo1/2, which License Specific mRNAs to Evade Degradation via the 5′-3′ mRNA Decay Pathway. Molecular Cell, 2010, 38, 345-355.	9.7	106
32	Phosphorylation, lipid raft interaction and traffic of \hat{l}_{\pm} -synuclein in a yeast model for Parkinson. Biochimica Et Biophysica Acta - Molecular Cell Research, 2008, 1783, 1767-1780.	4.1	104
33	Structural analysis of the subunits of the trehaloseâ€6â€phosphate synthase/phosphatase complex in Saccharomyces cerevisiae and their function during heat shock. Molecular Microbiology, 1997, 24, 687-696.	2.5	101
34	Heat shock induces enzymes of trehalose metabolism, trehalose accumulation, and thermotolerance inSchizosaccharomyces pombe, even in the presence of cycloheximide. FEBS Letters, 1990, 273, 107-110.	2.8	99
35	SEACing the GAP that nEGOCiates TORC1 activation. Cell Cycle, 2013, 12, 2948-2952.	2.6	98
36	Spatially Distinct Pools of TORC1 Balance Protein Homeostasis. Molecular Cell, 2019, 73, 325-338.e8.	9.7	95

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37	Bud8p and Bud9p, Proteins That May Mark the Sites for Bipolar Budding in Yeast. Molecular Biology of the Cell, 2001, 12, 2497-2518.	2.1	90
38	Multilayered Control of Protein Turnover by TORC1 and Atg1. Cell Reports, 2019, 28, 3486-3496.e6.	6.4	87
39	Cloning and disruption of a gene required for growth on acetate but not on ethanol: The acetyl-coenzyme a synthetase gene of Saccharmoyces cerevisiae. Yeast, 1992, 8, 1043-1051.	1.7	84
40	Conserved regulators of Rag GTPases orchestrate amino acid-dependent TORC1 signaling. Cell Discovery, 2016, 2, 15049.	6.7	84
41	The novel yeast PAS kinase Rim 15 orchestrates GO-associated antioxidant defense mechanisms. Cell Cycle, 2004, 3, 462-8.	2.6	84
42	TORC1 controls G1–S cell cycle transition in yeast via Mpk1 and the greatwall kinase pathway. Nature Communications, 2015, 6, 8256.	12.8	79
43	Yeast Endosulfines Control Entry into Quiescence and Chronological Life Span by Inhibiting Protein Phosphatase 2A. Cell Reports, 2013, 3, 16-22.	6.4	77
44	Acquisition of thermotolerance in Saccharomyces cerevisiae without heat shock protein hsp104 and in the absence of protein synthesis. FEBS Letters, 1991, 288, 86-90.	2.8	74
45	Saccharomyces cerevisiae Ccr4-Not complex contributes to the control of Msn2p-dependent transcription by the Ras/cAMP pathway. Molecular Microbiology, 2002, 43, 1023-1037.	2.5	72
46	Crystal structure of the Ego1-Ego2-Ego3 complex and its role in promoting Rag GTPase-dependent TORC1 signaling. Cell Research, 2015, 25, 1043-1059.	12.0	71
47	Trehalose synthesis is important for the acquisition of thermotolerance in Schizosaccharomyces pombe. Molecular Microbiology, 1997, 25, 571-581.	2.5	67
48	Bni5p, a Septin-Interacting Protein, Is Required for Normal Septin Function and Cytokinesis in Saccharomyces cerevisiae. Molecular and Cellular Biology, 2002, 22, 6906-6920.	2.3	65
49	The Ccr4-Not Complex Independently Controls both Msn2-Dependent Transcriptional Activation—via a Newly Identified Glc7/Bud14 Type I Protein Phosphatase Module—and TFIID Promoter Distribution. Molecular and Cellular Biology, 2005, 25, 488-498.	2.3	61
50	The Architecture of the Rag GTPase Signaling Network. Biomolecules, 2017, 7, 48.	4.0	59
51	Disruption in Candida albicans of the TPS2 gene encoding trehalose-6-phosphate phosphatase affects cell integrity and decreases infectivity The EMBL accession number for the sequence reported in this paper is AJ242990 Microbiology (United Kingdom), 2002, 148, 1281-1290.	1.8	59
52	Ego3 Functions as a Homodimer to Mediate the Interaction between Gtr1-Gtr2 and Ego1 in the EGO Complex to Activate TORC1. Structure, 2012, 20, 2151-2160.	3.3	56
53	The 70-kilodalton heat-shock proteins of the SSA subfamily negatively modulate heat-shock-induced accumulation of trehalose and promote recovery from heat stress in the yeast, Saccharomyces cerevisiae. FEBS Journal, 1992, 210, 125-132.	0.2	55
54	TORC1 Regulates Pah1 Phosphatidate Phosphatase Activity via the Nem1/Spo7 Protein Phosphatase Complex. PLoS ONE, 2014, 9, e104194.	2.5	53

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55	Expression of a functional barley sucrose-fructan 6-fructosyltransferase in the methylotrophic yeast Pichia pastoris. FEBS Letters, 1998, 440, 356-360.	2.8	51
56	Modulation of Ubc4p/Ubc5p-Mediated Stress Responses by the RING-Finger-Dependent Ubiquitin-Protein Ligase Not4p in Saccharomyces cerevisiae. Genetics, 2007, 176, 181-192.	2.9	48
57	Retromer and TBC1D5 maintain late endosomal RAB7 domains to enable amino acid–induced mTORC1 signaling. Journal of Cell Biology, 2019, 218, 3019-3038.	5. 2	46
58	The yeast protein kinase Sch9 adjusts V-ATPase assembly/disassembly to control pH homeostasis and longevity in response to glucose availability. PLoS Genetics, 2017, 13, e1006835.	3.5	45
59	The Thermophilic Yeast <i>Hansenula polymorpha</i> Does Not Require Trehalose Synthesis for Growth at High Temperatures but Does for Normal Acquisition of Thermotolerance. Journal of Bacteriology, 1999, 181, 4665-4668.	2.2	44
60	The I-BAR protein lvy1 is an effector of the Rab7 GTPase Ypt7 involved in vacuole membrane homeostasis. Journal of Cell Science, 2015, 128, 2278-2292.	2.0	40
61	Unsolved mysteries of Rag GTPase signaling in yeast. Small GTPases, 2016, 7, 239-246.	1.6	38
62	The Bud14p–Glc7p complex functions as a cortical regulator of dynein in budding yeast. EMBO Journal, 2005, 24, 3000-3011.	7.8	36
63	Membrane stress is coupled to a rapid translational control of gene expression in chlorpromazine-treated cells. Current Genetics, 2007, 52, 171-185.	1.7	36
64	CNE1, aSaccharomyces cerevisiae Homologue of the Genes Encoding Mammalian Calnexin and Calreticulin. Yeast, 1993, 9, 185-188.	1.7	34
65	A method to study the rapid phosphorylation-related modulation of neutral trehalase activity by temperature shifts in yeast. FEBS Letters, 1991, 291, 355-358.	2.8	33
66	TORC1 specifically inhibits microautophagy through ESCRT-0. Current Genetics, 2019, 65, 1243-1249.	1.7	32
67	Indole-3-acetic acid is a physiological inhibitor of TORC1 in yeast. PLoS Genetics, 2021, 17, e1009414.	3.5	32
68	TORC1 Determines Fab1 Lipid Kinase Function at Signaling Endosomes and Vacuoles. Current Biology, 2021, 31, 297-309.e8.	3.9	31
69	TORC1 coordinates the conversion of Sic1 from a target to an inhibitor of cyclin-CDK-Cks1. Cell Discovery, 2017, 3, 17012.	6.7	30
70	Cyclin-dependent kinase 5 (CDK5) regulates the circadian clock. ELife, 2019, 8, .	6.0	30
71	Control of Cellular Physiology by TM9 Proteins in Yeast and Dictyostelium. Journal of Biological Chemistry, 2008, 283, 6764-6772.	3.4	29
72	Mitochondrial Genomic Dysfunction Causes Dephosphorylation of Sch9 in the Yeast Saccharomyces cerevisiae. Eukaryotic Cell, 2011, 10, 1367-1369.	3.4	29

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73	Initiation of the yeast G0program requires $\log 1$ and $\log 2$, which antagonize activation of decapping of specific nutrient-regulated mRNAs. RNA Biology, 2011, 8, 14-17.	3.1	26
74	A spatially and functionally distinct pool of TORC1 defines signaling endosomes in yeast. Autophagy, 2019, 15, 915-916.	9.1	24
75	Identification of a Small Molecule Yeast TORC1 Inhibitor with a Multiplex Screen Based on Flow Cytometry. ACS Chemical Biology, 2012, 7, 715-722.	3.4	22
76	Feedback Inhibition of the Rag GTPase GAP Complex Lst4-Lst7 Safeguards TORC1 from Hyperactivation by Amino Acid Signals. Cell Reports, 2017, 20, 281-288.	6.4	22
77	Phosphoproteomic responses of TORC1 target kinases reveal discrete and convergent mechanisms that orchestrate the quiescence program in yeast. Cell Reports, 2021, 37, 110149.	6.4	20
78	The Impact of ESCRT on Aβ1-42 Induced Membrane Lesions in a Yeast Model for Alzheimer's Disease. Frontiers in Molecular Neuroscience, 2018, 11, 406.	2.9	19
79	An EGOcentric view of TORC1 signaling. Cell Cycle, 2010, 9, 221-222.	2.6	18
80	Global phosphoproteomics pinpoints uncharted Gcn2-mediated mechanisms of translational control. Molecular Cell, 2021, 81, 1879-1889.e6.	9.7	16
81	Structural insights into the EGO-TC–mediated membrane tethering of the TORC1-regulatory Rag GTPases. Science Advances, 2019, 5, eaax8164.	10.3	15
82	Phosphatidylinositol 4-Phosphate Is Required for Translation Initiation in Saccharomyces cerevisiae. Journal of Biological Chemistry, 2006, 281, 38139-38149.	3.4	13
83	Quantification of mRNA stability of stress-responsive yeast genes following conditional excision of open reading frames. RNA Biology, 2013, 10, 1299-1306.	3.1	10
84	Genetic and physical localization of the acetyl-coenzyme A synthetase geneACS1 on chromosome I ofSaccharomyces cerevisiae. Yeast, 1993, 9, 419-421.	1.7	6
85	The evolutionary conserved BER1 gene is involved in microtubule stability in yeast. Current Genetics, 2008, 53, 107-115.	1.7	6
86	N- and C-terminal Gln3 \hat{a} "Tor1 interaction sites: one acting negatively and the other positively to regulate nuclear Gln3 localization. Genetics, 2021, 217, .	2.9	6
87	The HOPS tethering complex is required to maintain signaling endosome identity and TORC1 activity. Journal of Cell Biology, 2022, 221, .	5.2	6