

Michael Downey

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4901680/publications.pdf>

Version: 2024-02-01

27
papers

1,194
citations

567144

15
h-index

526166

27
g-index

36
all docs

36
docs citations

36
times ranked

1841
citing authors

#	ARTICLE	IF	CITATIONS
1	A phosphatase complex that dephosphorylates γ -H2AX regulates DNA damage checkpoint recovery. <i>Nature</i> , 2006, 439, 497-501.	13.7	439
2	A Genome-Wide Screen Identifies the Evolutionarily Conserved KEOPS Complex as a Telomere Regulator. <i>Cell</i> , 2006, 124, 1155-1168.	13.5	158
3	A Screen for Candidate Targets of Lysine Polyphosphorylation Uncovers a Conserved Network Implicated in Ribosome Biogenesis. <i>Cell Reports</i> , 2018, 22, 3427-3439.	2.9	61
4	Acetylome Profiling Reveals Overlap in the Regulation of Diverse Processes by Sirtuins, Gcn5, and Esa1. <i>Molecular and Cellular Proteomics</i> , 2015, 14, 162-176.	2.5	59
5	γ -H2AX as a Checkpoint Maintenance Signal. <i>Cell Cycle</i> , 2006, 5, 1376-1381.	1.3	50
6	Gcn5 and Sirtuins Regulate Acetylation of the Ribosomal Protein Transcription Factor Ifh1. <i>Current Biology</i> , 2013, 23, 1638-1648.	1.8	43
7	Chromatin and DNA repair: the benefits of relaxation. <i>Nature Cell Biology</i> , 2006, 8, 9-10.	4.6	39
8	Polymerase Stalling during Replication, Transcription and Translation. <i>Current Biology</i> , 2014, 24, R445-R452.	1.8	36
9	A Broad Response to Intracellular Long-Chain Polyphosphate in Human Cells. <i>Cell Reports</i> , 2020, 33, 108318.	2.9	33
10	SIRT3 controls brown fat thermogenesis by deacetylation regulation of pathways upstream of UCP1. <i>Molecular Metabolism</i> , 2019, 25, 35-49.	3.0	30
11	Nonhistone targets of KAT2A and KAT2B implicated in cancer biology. <i>Biochemistry and Cell Biology</i> , 2019, 97, 30-45.	0.9	29
12	Hmgcs2-mediated ketogenesis modulates high-fat diet-induced hepatosteatosis. <i>Molecular Metabolism</i> , 2022, 61, 101494.	3.0	28
13	Heart failure drug proscillaridin A targets MYC overexpressing leukemia through global loss of lysine acetylation. <i>Journal of Experimental and Clinical Cancer Research</i> , 2019, 38, 251.	3.5	27
14	Model systems for studying polyphosphate biology: a focus on microorganisms. <i>Current Genetics</i> , 2021, 67, 331-346.	0.8	26
15	Non-histone protein acetylation by the evolutionarily conserved GCN5 and PCAF acetyltransferases. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2021, 1864, 194608.	0.9	23
16	Building a KATalogue of acetyllysine targeting and function. <i>Briefings in Functional Genomics</i> , 2016, 15, 109-118.	1.3	21
17	Proteins required for vacuolar function are targets of lysine polyphosphorylation in yeast. <i>FEBS Letters</i> , 2020, 594, 21-30.	1.3	17
18	From underlying chemistry to therapeutic potential: open questions in the new field of lysine polyphosphorylation. <i>Current Genetics</i> , 2019, 65, 57-64.	0.8	16

#	ARTICLE	IF	CITATIONS
19	The promises of lysine polyphosphorylation as a regulatory modification in mammals are tempered by conceptual and technical challenges. <i>BioEssays</i> , 2021, 43, e2100058.	1.2	10
20	Ddp1 Cooperates with Ppx1 to Counter a Stress Response Initiated by Nonvacuolar Polyphosphate. <i>MBio</i> , 2022, 13, .	1.8	10
21	A synthetic non-histone substrate to study substrate targeting by the Gcn5 HAT and sirtuin HDACs. <i>Journal of Biological Chemistry</i> , 2019, 294, 6227-6239.	1.6	9
22	A Stringent Analysis of Polyphosphate Dynamics in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2019, 201, .	1.0	8
23	Nicotinamide Suppresses the DNA Damage Sensitivity of <i>Saccharomyces cerevisiae</i> Independently of Sirtuin Deacetylases. <i>Genetics</i> , 2016, 204, 569-579.	1.2	6
24	Vtc5 Is Localized to the Vacuole Membrane by the Conserved AP-3 Complex to Regulate Polyphosphate Synthesis in Budding Yeast. <i>MBio</i> , 2021, 12, e0099421.	1.8	4
25	Targeting Polyphosphate Kinases in the Fight against <i>Pseudomonas aeruginosa</i> . <i>MBio</i> , 2021, 12, e0147721.	1.8	3
26	Repair Scaffolding Reaches New Heights at Blocked Replication Forks. <i>Molecular Cell</i> , 2010, 39, 162-164.	4.5	2
27	A screen for novel targets casts polyphosphorylation of lysine as a common post-translational modification. <i>FASEB Journal</i> , 2018, 32, 791.12.	0.2	0