

Philip Coffino

List of Publications by Year in descending order

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72
papers

4,109
citations

117625

34
h-index

114465

63
g-index

105
all docs

105
docs citations

105
times ranked

2441
citing authors

#	ARTICLE	IF	CITATIONS
1	Allostery Modulates Interactions between Proteasome Core Particles and Regulatory Particles. <i>Biomolecules</i> , 2022, 12, 764.	4.0	3
2	Identification of Novel Therapeutic Targets for Fibrolamellar Carcinoma Using Patient-Derived Xenografts and Direct-from-Patient Screening. <i>Cancer Discovery</i> , 2021, 11, 2544-2563.	9.4	27
3	Allosteric coupling between $\hat{\pm}$ -rings of the 20S proteasome. <i>Nature Communications</i> , 2020, 11, 4580.	12.8	16
4	Slippery Substrates Impair ATP-dependent Protease Function by Slowing Unfolding. <i>Journal of Biological Chemistry</i> , 2014, 289, 3826.	3.4	2
5	Ubiquitin-independent proteasomal degradation. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2014, 1843, 216-221.	4.1	190
6	Slippery Substrates Impair Function of a Bacterial Protease ATPase by Unbalancing Translocation versus Exit. <i>Journal of Biological Chemistry</i> , 2013, 288, 13243-13257.	3.4	26
7	Functional Asymmetries of Proteasome Translocase Pore. <i>Journal of Biological Chemistry</i> , 2012, 287, 18535-18543.	3.4	49
8	Ubiquitin Proteasome System in Stress and Disease. <i>Biochemistry Research International</i> , 2012, 2012, 1-2.	3.3	1
9	Dependence of Proteasome Processing Rate on Substrate Unfolding. <i>Journal of Biological Chemistry</i> , 2011, 286, 17495-17502.	3.4	41
10	Ordering an Engagement Ring. <i>Molecular Cell</i> , 2010, 38, 319-320.	9.7	0
11	A genetic screen for <i>Saccharomyces cerevisiae</i> mutants affecting proteasome function, using a ubiquitin-independent substrate. <i>Yeast</i> , 2008, 25, 199-217.	1.7	18
12	Structural elements of the ubiquitin-independent proteasome degnon of ornithine decarboxylase. <i>Biochemical Journal</i> , 2008, 410, 401-407.	3.7	53
13	The Cytoplasmic Hsp70 Chaperone Machinery Subjects Misfolded and Endoplasmic Reticulum Import-incompetent Proteins to Degradation via the Ubiquitin-Proteasome System. <i>Molecular Biology of the Cell</i> , 2007, 18, 153-165.	2.1	148
14	Proteasome substrate degradation requires association plus extended peptide. <i>EMBO Journal</i> , 2007, 26, 123-131.	7.8	113
15	Glycine-alanine repeats impair proper substrate unfolding by the proteasome. <i>EMBO Journal</i> , 2006, 25, 1720-1729.	7.8	73
16	Probing the Ubiquitin/Proteasome System with Ornithine Decarboxylase, a Ubiquitin-independent Substrate. <i>Methods in Enzymology</i> , 2005, 398, 399-413.	1.0	29
17	Proteasomes Begin Ornithine Decarboxylase Digestion at the C Terminus. <i>Journal of Biological Chemistry</i> , 2004, 279, 20959-20965.	3.4	49
18	Repeat Sequence of Epstein-Barr Virus-encoded Nuclear Antigen 1 Protein Interrupts Proteasome Substrate Processing. <i>Journal of Biological Chemistry</i> , 2004, 279, 8635-8641.	3.4	64

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19	Ubistatins Inhibit Proteasome-Dependent Degradation by Binding the Ubiquitin Chain. <i>Science</i> , 2004, 306, 117-120.	12.6	183
20	Development of a method for screening short-lived proteins using green fluorescent protein. <i>Genome Biology</i> , 2004, 5, R81.	9.6	16
21	Determinants of proteasome recognition of ornithine decarboxylase, a ubiquitin-independent substrate. <i>EMBO Journal</i> , 2003, 22, 1488-1496.	7.8	189
22	Transcriptional regulation of the ornithine decarboxylase gene by c-Myc/Max/Mad network and retinoblastoma protein interacting with c-Myc. <i>International Journal of Biochemistry and Cell Biology</i> , 2003, 35, 496-521.	2.8	32
23	Ubiquitin-independent Mechanisms of Mouse Ornithine Decarboxylase Degradation Are Conserved between Mammalian and Fungal Cells. <i>Journal of Biological Chemistry</i> , 2003, 278, 12135-12143.	3.4	76
24	Structural Elements of Antizymes 1 and 2 Are Required for Proteasomal Degradation of Ornithine Decarboxylase. <i>Journal of Biological Chemistry</i> , 2002, 277, 45957-45961.	3.4	44
25	An Easily Dissociated 26 S Proteasome Catalyzes an Essential Ubiquitin-mediated Protein Degradation Pathway in <i>Trypanosoma brucei</i> . <i>Journal of Biological Chemistry</i> , 2002, 277, 15486-15498.	3.4	59
26	Antizyme, a mediator of ubiquitin-independent proteasomal degradation. <i>Biochimie</i> , 2001, 83, 319-323.	2.6	69
27	Regulation of cellular polyamines by antizyme. <i>Nature Reviews Molecular Cell Biology</i> , 2001, 2, 188-194.	37.0	325
28	Regulated Degradation of Yeast Ornithine Decarboxylase. <i>Journal of Biological Chemistry</i> , 1999, 274, 25921-25926.	3.4	33
29	Antizyme2 Is a Negative Regulator of Ornithine Decarboxylase and Polyamine Transport. <i>Journal of Biological Chemistry</i> , 1999, 274, 26425-26430.	3.4	71
30	Structure of mammalian ornithine decarboxylase at 1.6 Å... resolution: stereochemical implications of PLP-dependent amino acid decarboxylases. <i>Structure</i> , 1999, 7, 567-581.	3.3	145
31	Î±5 subunit in <i>Trypanosoma brucei</i> proteasome can self-assemble to form a cylinder of four stacked heptamer rings. <i>Biochemical Journal</i> , 1999, 344, 349-358.	3.7	36
32	Î±5 subunit in <i>Trypanosoma brucei</i> proteasome can self-assemble to form a cylinder of four stacked heptamer rings. <i>Biochemical Journal</i> , 1999, 344, 349.	3.7	26
33	Developmental effect of polyamine depletion in <i>Caenorhabditis elegans</i> . <i>Biochemical Journal</i> , 1998, 333, 309-315.	3.7	30
34	The N Terminus of Antizyme Promotes Degradation of Heterologous Proteins. <i>Journal of Biological Chemistry</i> , 1996, 271, 4441-4446.	3.4	32
35	Identification of a Region of p53 That Confers Lability. <i>Journal of Biological Chemistry</i> , 1996, 271, 4447-4451.	3.4	27
36	Crystallization of a mammalian ornithine decarboxylase. , 1996, 24, 266-268.		5

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37	Degradation of Ornithine Decarboxylase by the Mammalian and Yeast 26S Proteasome Complexes Requires all the Components of the Protease. <i>FEBS Journal</i> , 1995, 229, 276-283.	0.2	38
38	Rat Antizyme Inhibits the Activity but Does Not Promote the Degradation of Mouse Ornithine Decarboxylase in <i>Trypanosoma brucei</i> . <i>Journal of Biological Chemistry</i> , 1995, 270, 10264-10271.	3.4	19
39	Expression and post-transcriptional regulation of ornithine decarboxylase during early <i>Xenopus</i> development. <i>FEBS Journal</i> , 1991, 202, 575-581.	0.2	43
40	Killer polyamines?. <i>Journal of Cellular Biochemistry</i> , 1991, 45, 54-58.	2.6	45
41	Ornithine decarboxylase of African trypanosomes. <i>Biochemical Society Transactions</i> , 1990, 18, 739-740.	3.4	2
42	Regulation of mouse ornithine decarboxylase activity by cell growth, serum and tetradecanoyl phorbol acetate is governed primarily by sequences within the coding region of the gene. <i>Nucleic Acids Research</i> , 1989, 17, 9843-9860.	14.5	57
43	Linkage genetics of mouse ornithine decarboxylase (Odc). <i>Genomics</i> , 1989, 5, 636-638.	2.9	15
44	Probable cloning artefacts previously interpreted as unusual leader sequences of rodent ornithine decarboxylase mRNAs – a cautionary tale. <i>Gene</i> , 1988, 69, 365-368.	2.2	15
45	Nucleotide sequence of the mouse ornithine decarboxylase gene. <i>Nucleic Acids Research</i> , 1988, 16, 2731-2732.	14.5	30
46	[2] Cultured S49 mouse T lymphoma cells. <i>Methods in Enzymology</i> , 1987, 151, 9-19.	1.0	19
47	Revertants of a trans-dominant S49 mouse lymphoma mutant that affects expression of cAMP-dependent protein kinase. <i>Cell</i> , 1983, 35, 311-320.	28.9	23
48	Increased histone mRNA levels during inhibition of protein synthesis. <i>Biochemical and Biophysical Research Communications</i> , 1983, 114, 131-137.	2.1	50
49	Quantitative forward-mutation specificity of mono-functional alkylating agents, ICR-191, and aflatoxin B1 in mouse lymphoma cells. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 1982, 95, 297-311.	1.0	13
50	Hormonal regulation of cloned genes. <i>Nature</i> , 1981, 292, 492-493.	27.8	2
51	G1 and S phase mammalian cells synthesize histones at equivalent rates. <i>Cell</i> , 1980, 21, 195-204.	28.9	109
52	Regulation of phosphodiesterase and ornithine decarboxylase by cAMP is cell cycle independent. <i>Journal of Cellular Physiology</i> , 1979, 101, 369-374.	4.1	24
53	Two-dimensional gel analysis of cyclic amp effects in cultured s49 mouse lymphoma cells: Protein modifications, inductions and repressions. <i>Cell</i> , 1979, 18, 719-733.	28.9	100
54	Regulation of Lymphoma Cell Growth by Cyclic AMP. , 1979, , 43-47.		0

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55	Complementation analysis of hormone-sensitive adenylate cyclase. <i>Nature</i> , 1978, 272, 720-722.	27.8	34
56	Studies of cyclic AMP action using mutant tissue culture cells. <i>In Vitro</i> , 1978, 14, 140-145.	1.2	31
57	Kinase-negative mutants of S49 mouse lymphoma cells carry a trans-dominant mutation affecting expression of cAMP-dependent protein kinase. <i>Cell</i> , 1978, 15, 1351-1361.	28.9	102
58	Subunit interaction in cyclic AMP-dependent protein kinase of mutant lymphoma cells.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1977, 74, 1167-1171.	7.1	6
59	Cyclic amp-induced cytolysis in S49 cells: selection of an unresponsive "deathless" mutant. <i>Cell</i> , 1977, 11, 149-155.	28.9	48
60	Mutations causing charge alterations in regulatory subunits of the cAMP-dependent protein kinase of cultured S49 lymphoma cells. <i>Cell</i> , 1977, 10, 381-391.	28.9	224
61	Hypoxanthine-guanine phosphoribosyl transferase with altered substrate affinity in mutant mouse lymphoma cells. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1977, 483, 70-78.	2.6	4
62	Coexpression of mutant and wild type protein kinase in lymphoma cells resistant to dibutyryl cyclic AMP. <i>Journal of Cellular Physiology</i> , 1977, 92, 437-445.	4.1	23
63	Receptors for Low-Molecular-Weight Hormones on Lymphocytes. , 1977, , 331-356.		2
64	CYCLIC NUCLEOTIDES AND CONTROL OF CELL PROLIFERATION. , 1977, , 536.		0
65	SUMMARY REMARKS OF CHAIRMAN. , 1977, , 49-51.		0
66	Hormone-Mediated Lymphoma Cell Death: Mechanisms Of Glucocorticoid And Cyclic Amp Action. <i>Journal of Investigative Dermatology</i> , 1976, 67, 648-649.	0.7	1
67	Molecular Mechanisms of Cyclic AMP Action: A Genetic Approach. , 1976, 32, 669-682.		25
68	A structural gene mutation affecting the regulatory subunit of cyclic AMP-dependent protein kinase in mouse lymphoma cells.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1975, 72, 5051-5055.	7.1	58
69	Somatic genetic analysis of cyclic AMP action: Selection of unresponsive mutants. <i>Journal of Cellular Physiology</i> , 1975, 85, 603-609.	4.1	124
70	Somatic genetic analysis of cyclic AMP action: Characterization of unresponsive mutants. <i>Journal of Cellular Physiology</i> , 1975, 85, 611-619.	4.1	96
71	Cyclic AMP-dependent protein kinase: pivotal role in regulation of enzyme induction and growth. <i>Science</i> , 1975, 190, 896-898.	12.6	166
72	Cloning of mouse myeloma cells and detection of rare variants. <i>Journal of Cellular Physiology</i> , 1972, 79, 429-440.	4.1	246