## Martin A Wear

List of Publications by Year in descending order

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279798 2,194 46 23 citations h-index papers

g-index 46 46 46 3141 citing authors all docs docs citations times ranked

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#	Article	IF	CITATIONS
1	Pilot scale production, extraction and purification of a thermostable phycocyanin from Synechocystis sp. PCC 6803. Bioresource Technology, 2022, 345, 126459.	9.6	8
2	Fast acting allosteric phosphofructokinase inhibitors block trypanosome glycolysis and cure acute African trypanosomiasis in mice. Nature Communications, 2021, 12, 1052.	12.8	21
3	Evaluation of novel 3D-printed monolithic adsorbers against conventional chromatography columns for the purification of c-phycocyanin from Spirulina. Algal Research, 2021, 55, 102253.	4.6	10
4	Kinetic and structural studies of <i>Trypanosoma</i> and <i>Leishmania</i> phosphofructokinases show evolutionary divergence and identify AMP as a switch regulating glycolysis <i>versus</i> gluconeogenesis. FEBS Journal, 2020, 287, 2847-2861.	4.7	8
5	A Truncated Form of HpARI Stabilizes IL-33, Amplifying Responses to the Cytokine. Frontiers in Immunology, 2020, 11, 1363.	4.8	18
6	A helminth-derived suppressor of ST2 blocks allergic responses. ELife, 2020, 9, .	6.0	39
7	A computationally designed binding mode flip leads to a novel class of potent tri-vector cyclophilin inhibitors. Chemical Science, 2019, 10, 542-547.	7.4	17
8	Borealin–nucleosome interaction secures chromosome association of the chromosomal passenger complex. Journal of Cell Biology, 2019, 218, 3912-3925.	5.2	34
9	A chicken bioreactor for efficient production of functional cytokines. BMC Biotechnology, 2018, 18, 82.	3.3	33
10	Redox regulation of pyruvate kinase M2 by cysteine oxidation and S-nitrosation. Biochemical Journal, 2018, 475, 3275-3291.	3.7	24
11	An allostatic mechanism for M2 pyruvate kinase as an amino-acid sensor. Biochemical Journal, 2018, 475, 1821-1837.	3.7	44
12	Thermo-kinetic analysis space expansion for cyclophilin-ligand interactions - identification of a new nonpeptide inhibitor using Biacoreâ,,¢ T200. FEBS Open Bio, 2017, 7, 533-549.	2.3	8
13	Molecular basis for Cdk1â€regulated timing of Mis18 complex assembly and CENPâ€A deposition. EMBO Reports, 2017, 18, 894-905.	4.5	51
14	HpARI Protein Secreted by a Helminth Parasite Suppresses Interleukin-33. Immunity, 2017, 47, 739-751.e5.	14.3	130
15	Pushing the Limits of Detection of Weak Binding Using Fragment-Based Drug Discovery: Identification of New Cyclophilin Binders. Journal of Molecular Biology, 2017, 429, 2556-2570.	4.2	16
16	Biophysical Characterization and Activity of Lymphostatin, a Multifunctional Virulence Factor of Attaching and Effacing Escherichia coli. Journal of Biological Chemistry, 2016, 291, 5803-5816.	3.4	9
17	Cyclophilin40 isomerase activity is regulated by a temperature-dependent allosteric interaction with Hsp90. Bioscience Reports, 2015, 35, .	2.4	11
18	A Streamlined, Automated Protocol for the Production of Milligram Quantities of Untagged Recombinant Rat Lactate Dehydrogenase A Using ÄKTAxpressTM. PLoS ONE, 2015, 10, e0146164.	2.5	9

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19	Inhibition of the ERCC1–XPF structure-specific endonuclease to overcome cancer chemoresistance. DNA Repair, 2015, 31, 19-28.	2.8	56
20	Trypanosomatid phosphoglycerate mutases have multiple conformational and oligomeric states. Biochemical and Biophysical Research Communications, 2014, 450, 936-941.	2.1	9
21	Wzi Is an Outer Membrane Lectin that Underpins Group 1 Capsule Assembly in Escherichia coli. Structure, 2013, 21, 844-853.	3.3	63
22	Triazole biotin: a tight-binding biotinidase-resistant conjugate. Organic and Biomolecular Chemistry, 2013, 11, 7700.	2.8	18
23	M2 pyruvate kinase provides a mechanism for nutrient sensing and regulation of cell proliferation. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 5881-5886.	7.1	132
24	The Mitosis and Neurodevelopment Proteins NDE1 and NDEL1 Form Dimers, Tetramers, and Polymers with a Folded Back Structure in Solution. Journal of Biological Chemistry, 2012, 287, 32381-32393.	3.4	38
25	A t(1;11) translocation linked to schizophrenia and affective disorders gives rise to aberrant chimeric DISC1 transcripts that encode structurally altered, deleterious mitochondrial proteins. Human Molecular Genetics, 2012, 21, 3374-3386.	2.9	61
26	Design and Synthesis of Conformationally Constrained Cyclophilin Inhibitors Showing a Cyclosporinâ€A Phenotype in <i>C. elegans</i> . ChemBioChem, 2011, 12, 802-810.	2.6	10
27	Streamlined, automated protocols for the production of milligram quantities of untagged recombinant human cyclophilin-A (hCypA) and untagged human proliferating cell nuclear antigen (hPCNA) using ÄKTAxpress™. Protein Expression and Purification, 2010, 71, 54-61.	1.3	12
28	A global benchmark study using affinity-based biosensors. Analytical Biochemistry, 2009, 386, 194-216.	2.4	85
29	Specificity of Atonal and Scute bHLH factors: analysis of cognate E box binding sites and the influence of Senseless. Genes To Cells, 2008, 13, 915-929.	1.2	29
30	Sequence specificity of single-stranded DNA-binding proteins: a novel DNA microarray approach. Nucleic Acids Research, 2007, 35, e75.	14.5	22
31	A kinetically trapped intermediate of FK506 binding protein forms in vitro: Chaperone machinery dominates protein folding in vivo. Protein Expression and Purification, 2007, 51, 80-95.	1.3	18
32	Structure-based discovery of a family of synthetic cyclophilin inhibitors showing a cyclosporin-A phenotype in Caenorhabditis elegans. Biochemical and Biophysical Research Communications, 2007, 363, 1013-1019.	2.1	23
33	Actin Filament Severing by Cofilin. Journal of Molecular Biology, 2007, 365, 1350-1358.	4.2	164
34	Experimental Determination of van der Waals Energies in a Biological System. Angewandte Chemie - International Edition, 2007, 46, 6453-6456.	13.8	15
35	Determination of the rate constants for the FK506 binding protein/rapamycin interaction using surface plasmon resonance: An alternative sensor surface for Ni2+–nitrilotriacetic acid immobilization of His-tagged proteins. Analytical Biochemistry, 2007, 371, 250-252.	2.4	16
36	Thermodynamics of the cyclophilin-A/cyclosporin-A interaction: A direct comparison of parameters determined by surface plasmon resonance using Biacore T100 and isothermal titration calorimetry. Analytical Biochemistry, 2006, 359, 285-287.	2.4	31

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37	A surface plasmon resonance-based assay for small molecule inhibitors of human cyclophilin A. Analytical Biochemistry, 2005, 345, 214-226.	2.4	61
38	Mammalian CARMIL Inhibits Actin Filament Capping by Capping Protein. Developmental Cell, 2005, 9, 209-221.	7.0	114
39	Capping protein binding to actin in yeast. Journal of Cell Biology, 2004, 164, 567-580.	5.2	90
40	Capping Protein Binding to S100B. Journal of Biological Chemistry, 2004, 279, 14382-14390.	3.4	14
41	Biological role and structural mechanism of twinfilin–capping protein interaction. EMBO Journal, 2004, 23, 3010-3019.	7.8	71
42	Capping protein: new insights into mechanism and regulation. Trends in Biochemical Sciences, 2004, 29, 418-428.	7.5	114
43	How Capping Protein Binds the Barbed End of the Actin Filament. Current Biology, 2003, 13, 1531-1537.	3.9	143
44	Arp2/3 Complex. Cell, 2001, 107, 703-705.	28.9	32
45	Interactions with PIP2, ADP-actin monomers, and capping protein regulate the activity and localization of yeast twinfilin. Journal of Cell Biology, 2001, 155, 251-260.	5.2	156
46	Actin dynamics: Assembly and disassembly of actin networks. Current Biology, 2000, 10, R891-R895.	3.9	107