Thomas Alfred Edwards

List of Publications by Year in descending order

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66 papers 2,884 citations

186265
28
h-index

52 g-index

74 all docs

74 docs citations

times ranked

74

4142 citing authors

#	Article	IF	Citations
1	Sal-type ABC-F proteins: intrinsic and common mediators of pleuromutilin resistance by target protection in staphylococci. Nucleic Acids Research, 2022, 50, 2128-2142.	14.5	16
2	Understanding p300-transcription factor interactions using sequence variation and hybridization. RSC Chemical Biology, 2022, 3, 592-603.	4.1	2
3	The Native Orthobunyavirus Ribonucleoprotein Possesses a Helical Architecture. MBio, 2022, 13, .	4.1	10
4	Development of a multiplex assay for antibody detection in serum against pathogens affecting ruminants. Transboundary and Emerging Diseases, 2021, 68, 1229-1239.	3.0	7
5	Selective Affimers Recognise the BCLâ€2 Family Proteins BCLâ€x _L and MCLâ€1 through Noncanonical Structural Motifs**. ChemBioChem, 2021, 22, 232-240.	2.6	9
6	Query-guided protein–protein interaction inhibitor discovery. Chemical Science, 2021, 12, 4753-4762.	7.4	5
7	Towards optimizing peptide-based inhibitors of protein–protein interactions: predictive saturation variation scanning (PreSaVS). RSC Chemical Biology, 2021, 2, 1474-1478.	4.1	5
8	Identification of β-strand mediated protein–protein interaction inhibitors using ligand-directed fragment ligation. Chemical Science, 2021, 12, 2286-2293.	7.4	3
9	RAS-inhibiting biologics identify and probe druggable pockets including an SII-α3 allosteric site. Nature Communications, 2021, 12, 4045.	12.8	19
10	Structure of the 70S Ribosome from the Human Pathogen Acinetobacter baumannii in Complex with Clinically Relevant Antibiotics. Structure, 2020, 28, 1087-1100.e3.	3.3	16
11	Characterization and applications of a Crimean-Congo hemorrhagic fever virus nucleoprotein-specific Affimer: Inhibitory effects in viral replication and development of colorimetric diagnostic tests. PLoS Neglected Tropical Diseases, 2020, 14, e0008364.	3.0	4
12	A missense variant in specificity protein 6 (SP6) is associated with amelogenesis imperfecta. Human Molecular Genetics, 2020, 29, 1417-1425.	2.9	21
13	Stapled Peptides as HIFâ€1α/p300 Inhibitors: Helicity Enhancement in the Bound State Increases Inhibitory Potency. Chemistry - A European Journal, 2020, 26, 7638-7646.	3.3	16
14	Predicting and Experimentally Validating Hot-Spot Residues at Protein–Protein Interfaces. ACS Chemical Biology, 2019, 14, 2252-2263.	3.4	54
15	Reply to Rameix-Welti, "No Incongruity in Respiratory Syncytial Virus M2-1 Protein Remaining Bound to Viral mRNAs during Their Entire Life Time― MBio, 2019, 10, .	4.1	0
16	Tula orthohantavirus nucleocapsid protein is cleaved in infected cells and may sequester activated caspase-3 during persistent infection to suppress apoptosis. Journal of General Virology, 2019, 100, 1208-1221.	2.9	5
17	The Structure of the Human Respiratory Syncytial Virus M2-1 Protein Bound to the Interaction Domain of the Phosphoprotein P Defines the Orientation of the Complex. MBio, 2018, 9, .	4.1	28
18	Identification of a small molecule inhibitor of Ebola virus genome replication and transcription using in silico screening. Antiviral Research, 2018, 156, 46-54.	4.1	14

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19	Structure and Function of the Human Respiratory Syncytial Virus M2–1 Protein. Sub-Cellular Biochemistry, 2018, 88, 245-260.	2.4	4
20	Hypoxia inducible factor (HIF) as a model for studying inhibition of protein–protein interactions. Chemical Science, 2017, 8, 4188-4202.	7.4	30
21	Generation of specific inhibitors of SUMO-1– and SUMO-2/3–mediated protein-protein interactions using Affimer (Adhiron) technology. Science Signaling, 2017, 10, .	3.6	44
22	Probing Protein Surfaces: QSAR Analysis with Helix Mimetics. ChemBioChem, 2016, 17, 768-773.	2.6	5
23	Synthesis of highly functionalized oligobenzamide proteomimetic foldamers by late stage introduction of sensitive groups. Organic and Biomolecular Chemistry, 2016, 14, 3782-3786.	2.8	17
24	ABC-F Proteins Mediate Antibiotic Resistance through Ribosomal Protection. MBio, 2016, 7, e01975.	4.1	222
25	Heat Shock Protein 70 Family Members Interact with Crimean-Congo Hemorrhagic Fever Virus and Hazara Virus Nucleocapsid Proteins and Perform a Functional Role in the Nairovirus Replication Cycle. Journal of Virology, 2016, 90, 9305-9316.	3.4	36
26	Hypertrophic cardiomyopathy mutations in the calponin-homology domain of ACTN2 affect actin binding and cardiomyocyte Z-disc incorporation. Biochemical Journal, 2016, 473, 2485-2493.	3.7	42
27	Genetic analysis of Physcomitrella patens identifies ABSCISIC ACID NON-RESPONSIVE (ANR), a regulator of ABA responses unique to basal land plants and required for desiccation tolerance. Plant Cell, 2016, 28, tpc.00091.2016.	6.6	98
28	Towards "bionic―proteins: replacement of continuous sequences from HIF-1α with proteomimetics to create functional p300 binding HIF-1α mimics. Chemical Communications, 2016, 52, 5421-5424.	4.1	20
29	Hydrocarbon constrained peptides – understanding preorganisation and binding affinity. Chemical Science, 2016, 7, 3694-3702.	7.4	63
30	Selective and Potent Proteomimetic Inhibitors of Intracellular Protein–Protein Interactions. Angewandte Chemie, 2015, 127, 3003-3008.	2.0	24
31	Bespoke RNA recognition by Pumilios. Biochemical Society Transactions, 2015, 43, 801-806.	3.4	9
32	The crystal structure of the Hazara virus nucleocapsid protein. BMC Structural Biology, 2015, 15, 24.	2.3	26
33	Selective and Potent Proteomimetic Inhibitors of Intracellular Protein–Protein Interactions. Angewandte Chemie - International Edition, 2015, 54, 2960-2965.	13.8	82
34	Stereocontrolled protein surface recognition using chiral oligoamide proteomimetic foldamers. Chemical Science, 2015, 6, 2434-2443.	7.4	58
35	Exploration of the HIF- $1\hat{1}\pm/p300$ interface using peptide and Adhiron phage display technologies. Molecular BioSystems, 2015, 11, 2738-2749.	2.9	35
36	Smallâ€Molecule Proteomimetic Inhibitors of the HIFâ€1α–p300 Protein–Protein Interaction. ChemBioChem, 2014, 15, 1083-1087.	' 2.6	57

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37	Crystal structure of the essential transcription antiterminator M2-1 protein of human respiratory syncytial virus and implications of its phosphorylation. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 1580-1585.	7.1	58
38	Probing Bunyavirus N protein oligomerisation using mass spectrometry. Rapid Communications in Mass Spectrometry, 2014, 28, 793-800.	1.5	6
39	Orthogonal functionalisation of α-helix mimetics. Organic and Biomolecular Chemistry, 2014, 12, 6794-6799.	2.8	24
40	Microwave assisted solid phase synthesis of highly functionalized N-alkylated oligobenzamide \hat{l}_{\pm} -helix mimetics. Bioorganic and Medicinal Chemistry, 2013, 21, 4034-4040.	3.0	42
41	Aromatic Oligoamide Foldamers with a "Wet Edgeâ€as Inhibitors of the αâ€Helixâ€Mediated p53– <i>h</i> Protein–Protein Interaction. European Journal of Organic Chemistry, 2013, 2013, 3504-3512.	DM2 2.4	23
42	Mutagenesis Mapping of the Protein-Protein Interaction Underlying FusB-Type Fusidic Acid Resistance. Antimicrobial Agents and Chemotherapy, 2013, 57, 4640-4644.	3.2	4
43	Nucleocapsid protein structures from orthobunyaviruses reveal insight into ribonucleoprotein architecture and RNA polymerization. Nucleic Acids Research, 2013, 41, 5912-5926.	14.5	69
44	Structure, Function, and Evolution of the Crimean-Congo Hemorrhagic Fever Virus Nucleocapsid Protein. Journal of Virology, 2012, 86, 10914-10923.	3.4	94
45	Ribosome clearance by FusB-type proteins mediates resistance to the antibiotic fusidic acid. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 2102-2107.	7.1	36
46	2-O-Alkylated para-benzamide $\hat{l}\pm$ -helix mimetics: the role of scaffold curvature. Organic and Biomolecular Chemistry, 2012, 10, 6469.	2.8	46
47	ATP-Driven Remodeling of the Linker Domain in the Dynein Motor. Structure, 2012, 20, 1670-1680.	3.3	76
48	Conformational properties of O-alkylated benzamides. Tetrahedron, 2012, 68, 4485-4491.	1.9	21
49	Modeling of Arylamide Helix Mimetics in the p53 Peptide Binding Site of hDM2 Suggests Parallel and Anti-Parallel Conformations Are Both Stable. PLoS ONE, 2012, 7, e43253.	2.5	16
50	Helix-mediated protein–protein interactions as targets for intervention using foldamers. Amino Acids, 2011, 41, 743-754.	2.7	72
51	Kinked \hat{l}^2 -strands mediate high-affinity recognition of mRNA targets by the germ-cell regulator DAZL. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 18266-18271.	7.1	58
52	N-alkylated oligoamide α-helical proteomimetics. Organic and Biomolecular Chemistry, 2010, 8, 2344.	2.8	79
53	Co-occupancy of two Pumilio molecules on a single hunchback NRE. Rna, 2009, 15, 1029-1035.	3.5	24
54	Structure and RNA binding of the mouse Pumilio-2 Puf domain. Journal of Structural Biology, 2009, 167, 271-276.	2.8	26

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55	Oligobenzamide proteomimetic inhibitors of the p53–hDM2 protein–protein interaction. Chemical Communications, 2009, , 5091.	4.1	124
56	A Beta-Sheet Interaction Interface Directs the Tetramerisation of the Miz-1 POZ Domain. Journal of Molecular Biology, 2007, 373, 820-826.	4.2	33
57	Solution Structure of the Vts1 SAM Domain in the Presence of RNA. Journal of Molecular Biology, 2006, 356, 1065-1072.	4.2	15
58	Crystal Structure of the Catalytic Core of Human DNA Polymerase Kappa. Structure, 2004, 12, 1395-1404.	3.3	107
59	Crystal Structure of BstYl at 1.85Ã Resolution: A Thermophilic Restriction Endonuclease with Overlapping Specificities to BamHI and BglII. Journal of Molecular Biology, 2004, 338, 725-733.	4.2	15
60	Crystal Structure of the SF3 Helicase from Adeno-Associated Virus Type 2. Structure, 2003, 11, 1025-1035.	3.3	96
61	Model of the Brain Tumor-Pumilio translation repressor complex. Genes and Development, 2003, 17, 2508-2513.	5.9	97
62	Crystallization and characterization of Smaug: a novel RNA-binding motif. Biochemical and Biophysical Research Communications, 2002, 297, 1085-1088.	2.1	8
63	Transcription Corepressor CtBP Is an NAD+-Regulated Dehydrogenase. Molecular Cell, 2002, 10, 857-869.	9.7	252
64	Disruption of sixSaccharomyces cerevisiae ORFs on chromosome XII results in three lethal disruptants. Yeast, 2002, 19, 79-86.	1.7	1
65	Structure of Pumilio Reveals Similarity between RNA and Peptide Binding Motifs. Cell, 2001, 105, 281-289.	28.9	237
66	Crystallization and Characterization of Pumilio: A Novel RNA Binding Protein. Journal of Structural Biology, 2000, 132, 251-254.	2.8	15