

Colin R Groom

List of Publications by Year in descending order

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32
papers

15,284
citations

331259

21
h-index

395343

33
g-index

37
all docs

37
docs citations

37
times ranked

18496
citing authors

#	ARTICLE	IF	CITATIONS
1	The Cambridge Structural Database. <i>Acta Crystallographica Section B: Structural Science, Crystal Engineering and Materials</i> , 2016, 72, 171-179.	0.5	7,159
2	The druggable genome. <i>Nature Reviews Drug Discovery</i> , 2002, 1, 727-730.	21.5	2,918
3	Ligand efficiency: a useful metric for lead selection. <i>Drug Discovery Today</i> , 2004, 9, 430-431.	3.2	1,687
4	The Cambridge Structural Database in Retrospect and Prospect. <i>Angewandte Chemie - International Edition</i> , 2014, 53, 662-671.	7.2	1,004
5	Report on the sixth blind test of organic crystal structure prediction methods. <i>Acta Crystallographica Section B: Structural Science, Crystal Engineering and Materials</i> , 2016, 72, 439-459.	0.5	445
6	Pheromone binding to two rodent urinary proteins revealed by X-ray crystallography. <i>Nature</i> , 1992, 360, 186-188.	13.7	374
7	Three-dimensional structure of diferric bovine lactoferrin at 2.8 Å... resolution. <i>Journal of Molecular Biology</i> , 1997, 274, 222-236.	2.0	361
8	Heteroaromatic Rings of the Future. <i>Journal of Medicinal Chemistry</i> , 2009, 52, 2952-2963.	2.9	286
9	Atomic Interactions and Profile of Small Molecules Disrupting Protein-Protein Interfaces: the TIMBAL Database. <i>Chemical Biology and Drug Design</i> , 2009, 74, 457-467.	1.5	144
10	Evaluation of molecular crystal structures using Full Interaction Maps. <i>CrystEngComm</i> , 2013, 15, 65-72.	1.3	109
11	Experimental and computational mapping of the binding surface of a crystalline protein. <i>Protein Engineering, Design and Selection</i> , 2001, 14, 47-59.	1.0	104
12	Locating interaction sites on proteins: The crystal structure of thermolysin soaked in 2% to 100% isopropanol. , 1999, 37, 628-640.		99
13	The good, the bad and the twisted: a survey of ligand geometry in protein crystal structures. <i>Journal of Computer-Aided Molecular Design</i> , 2012, 26, 169-183.	1.3	92
14	The Hydrogen Bond Environments of 1 <i>H</i> -Tetrazole and Tetrazolate Rings: The Structural Basis for Tetrazole-Carboxylic Acid Bioisosterism. <i>Journal of Chemical Information and Modeling</i> , 2012, 52, 857-866.	2.5	65
15	Identification, classification and relative stability of tautomers in the cambridge structural database. <i>CrystEngComm</i> , 2011, 13, 93-98.	1.3	58
16	IRAK-4 inhibitors. Part II: A structure-based assessment of imidazo[1,2-a]pyridine binding. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2008, 18, 3291-3295.	1.0	47
17	One in half a million: a solid form informatics study of a pharmaceutical crystal structure. <i>CrystEngComm</i> , 2012, 14, 2391-2403.	1.3	42
18	A crystallographic perspective on sharing data and knowledge. <i>Journal of Computer-Aided Molecular Design</i> , 2014, 28, 1015-1022.	1.3	27

#	ARTICLE	IF	CITATIONS
19	The use of small-molecule structures to complement protein-ligand crystal structures in drug discovery. <i>Acta Crystallographica Section D: Structural Biology</i> , 2017, 73, 240-245.	1.1	25
20	Hydrogen bonding at C=Se acceptors in selenoureas, selenoamides and selones. <i>Acta Crystallographica Section B: Structural Science, Crystal Engineering and Materials</i> , 2016, 72, 317-325.	0.5	23
21	The Cambridge Structural Database: experimental three-dimensional information on small molecules is a vital resource for interdisciplinary research and learning. <i>Wiley Interdisciplinary Reviews: Computational Molecular Science</i> , 2011, 1, 368-376.	6.2	22
22	Generation of crystal structures using known crystal structures as analogues. <i>Acta Crystallographica Section B: Structural Science, Crystal Engineering and Materials</i> , 2016, 72, 530-541.	0.5	18
23	Protein kinase drugs – optimism doesn't wait on facts –¾. <i>Drug Discovery Today</i> , 2002, 7, 801-802.	3.2	15
24	Capturing neon – the first experimental structure of neon trapped within a metal-organic environment. <i>Chemical Communications</i> , 2016, 52, 10048-10051.	2.2	13
25	Data to knowledge: how to get meaning from your result. <i>IUCr</i> , 2015, 2, 45-58.	1.0	12
26	Knowledge-Based Optimization of Molecular Geometries Using Crystal Structures. <i>Journal of Chemical Information and Modeling</i> , 2016, 56, 652-661.	2.5	12
27	Up the Garden Path: A Chemical Trail through the Cambridge University Botanic Garden. <i>Journal of Chemical Education</i> , 2012, 89, 1390-1394.	1.1	9
28	Data-Driven High-Throughput Prediction of the 3-D Structure of Small Molecules: Review and Progress. A Response from The Cambridge Crystallographic Data Centre. <i>Journal of Chemical Information and Modeling</i> , 2011, 51, 2787-2787.	2.5	8
29	Using more than 801,296 small-molecule crystal structures to aid in protein structure refinement and analysis. <i>Acta Crystallographica Section D: Structural Biology</i> , 2017, 73, 234-239.	1.1	6
30	Institutional Profile: Crystal structure information in drug discovery and development: current perspectives and new possibilities from the Cambridge Crystallographic Data Centre. <i>Future Medicinal Chemistry</i> , 2010, 2, 933-939.	1.1	5
31	Small Molecule Crystal Structures in Drug Discovery. <i>NATO Science for Peace and Security Series A: Chemistry and Biology</i> , 2015, , 107-114.	0.5	2
32	The Cambridge Structural Database (CSD). , 2019, , .		2