

Joseph D Bauman

List of Publications by Year in descending order

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15
papers

771
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933447

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docs citations

15
times ranked

1198
citing authors

#	ARTICLE	IF	CITATIONS
1	HIV-1 reverse transcriptase complex with DNA and nevirapine reveals non-nucleoside inhibition mechanism. <i>Nature Structural and Molecular Biology</i> , 2012, 19, 253-259.	8.2	176
2	Structure of HIV-1 Reverse Transcriptase with the Inhibitor Î²-Thujaplicinol Bound at the RNase H Active Site. <i>Structure</i> , 2009, 17, 1625-1635.	3.3	135
3	Crystal engineering of HIV-1 reverse transcriptase for structure-based drug design. <i>Nucleic Acids Research</i> , 2008, 36, 5083-5092.	14.5	91
4	Detecting Allosteric Sites of HIV-1 Reverse Transcriptase by X-ray Crystallographic Fragment Screening. <i>Journal of Medicinal Chemistry</i> , 2013, 56, 2738-2746.	6.4	78
5	Crystallographic Fragment Screening and Structure-Based Optimization Yields a New Class of Influenza Endonuclease Inhibitors. <i>ACS Chemical Biology</i> , 2013, 8, 2501-2508.	3.4	76
6	Advantages of crystallographic fragment screening: Functional and mechanistic insights from a powerful platform for efficient drug discovery. <i>Progress in Biophysics and Molecular Biology</i> , 2014, 116, 92-100.	2.9	73
7	3-Hydroxyquinolin-2(1 <i>H</i>)-ones As Inhibitors of Influenza A Endonuclease. <i>ACS Medicinal Chemistry Letters</i> , 2013, 4, 547-550.	2.8	44
8	Rapid experimental SAD phasing and hot-spot identification with halogenated fragments. <i>IUCr</i> , 2016, 3, 51-60.	2.2	27
9	Fragment Screening and HIV Therapeutics. <i>Topics in Current Chemistry</i> , 2011, 317, 181-200.	4.0	20
10	A New Class of Allosteric HIV-1 Integrase Inhibitors Identified by Crystallographic Fragment Screening of the Catalytic Core Domain. <i>Journal of Biological Chemistry</i> , 2016, 291, 23569-23577.	3.4	20
11	Structure of HIV-1 reverse transcriptase/d4TTP complex: Novel DNA cross-linking site and pH-dependent conformational changes. <i>Protein Science</i> , 2019, 28, 587-597.	7.6	11
12	Integrative structural biology studies of HIV-1 reverse transcriptase binding to a high-affinity DNA aptamer. <i>Current Research in Structural Biology</i> , 2020, 2, 116-129.	2.2	8
13	Differential Isotopic Enrichment To Facilitate Characterization of Asymmetric Multimeric Proteins Using Hydrogen/Deuterium Exchange Mass Spectrometry. <i>Analytical Chemistry</i> , 2015, 87, 4015-4022.	6.5	4
14	Aryl and Arylalkyl Substituted 3-Hydroxypyridin-2(1 <i>H</i>)-ones: Synthesis and Evaluation as Inhibitors of Influenza A Endonuclease. <i>ChemMedChem</i> , 2019, 14, 1204-1223.	3.2	4
15	Crystal Structure of a Retroviral Polyprotein: Prototype Foamy Virus Protease-Reverse Transcriptase (PR-RT). <i>Viruses</i> , 2021, 13, 1495.	3.3	4