Daniel J Deredge

List of Publications by Year in descending order

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567281 580821 30 701 15 25 citations h-index g-index papers 34 34 34 1297 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Conformational dynamics of a neurotransmitter:sodium symporter in a lipid bilayer. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E1786-E1795.	7.1	76
2	Evolution of Structural Biology through the Lens of Mass Spectrometry. Analytical Chemistry, 2019, 91, 142-155.	6.5	49
3	IL-1 Family Cytokines Use Distinct Molecular Mechanisms to Signal through Their Shared Co-receptor. Immunity, 2017, 47, 510-523.e4.	14.3	48
4	A Dimer Interface Mutation in Glyceraldehyde-3-Phosphate Dehydrogenase Regulates Its Binding to AU-rich RNA. Journal of Biological Chemistry, 2015, 290, 1770-1785.	3.4	47
5	The <i>Helicobacter pylori</i> adhesin protein HopQ exploits the dimer interface of human CEACAMs to facilitate translocation of the oncoprotein CagA. EMBO Journal, 2018, 37, .	7.8	47
6	Common coding variant in <i>SERPINA1</i> increases the risk for large artery stroke. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 3613-3618.	7.1	46
7	Bacterial flagellar capping proteins adopt diverse oligomeric states. ELife, 2016, 5, .	6.0	46
8	Interlaboratory Comparison of Hydrogen–Deuterium Exchange Mass Spectrometry Measurements of the Fab Fragment of NISTmAb. Analytical Chemistry, 2019, 91, 7336-7345.	6.5	44
9	Current Trends and Limitations in Dengue Antiviral Research. Tropical Medicine and Infectious Disease, 2021, 6, 180.	2.3	35
10	Structure and dynamics of an \hat{l}_{\pm} -fucosidase reveal a mechanism for highly efficient lgG transfucosylation. Nature Communications, 2020, 11, 6204.	12.8	29
11	Gallium(III)–Salophen as a Dual Inhibitor of <i>Pseudomonas aeruginosa</i> Heme Sensing and Iron Acquisition. ACS Infectious Diseases, 2020, 6, 2073-2085.	3.8	29
11	Gallium(III)–Salophen as a Dual Inhibitor of <i>Pseudomonas aeruginosa</i> Acquisition. ACS Infectious Diseases, 2020, 6, 2073-2085. Structure and Dynamics of FosA-Mediated Fosfomycin Resistance in Klebsiella pneumoniae and Escherichia coli. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.8	29
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12	Acquisition. ACS Infectious Diseases, 2020, 6, 2073-2085. Structure and Dynamics of FosA-Mediated Fosfomycin Resistance in Klebsiella pneumoniae and Escherichia coli. Antimicrobial Agents and Chemotherapy, 2017, 61, . Molecular Basis of Broad Spectrum <i>N</i> -Glycan Specificity and Processing of Therapeutic IgG	3.2	28
12	Acquisition. ACS Infectious Diseases, 2020, 6, 2073-2085. Structure and Dynamics of FosA-Mediated Fosfomycin Resistance in Klebsiella pneumoniae and Escherichia coli. Antimicrobial Agents and Chemotherapy, 2017, 61,. Molecular Basis of Broad Spectrum <i>N</i> -Clycan Specificity and Processing of Therapeutic IgG Monoclonal Antibodies by Endoglycosidase S2. ACS Central Science, 2019, 5, 524-538. Antigen-Induced Allosteric Changes in a Human IgG1 Fc Increase Low-Affinity Fcî³ Receptor Binding.	3.2	28
12 13 14	Acquisition. ACS Infectious Diseases, 2020, 6, 2073-2085. Structure and Dynamics of FosA-Mediated Fosfomycin Resistance in Klebsiella pneumoniae and Escherichia coli. Antimicrobial Agents and Chemotherapy, 2017, 61, . Molecular Basis of Broad Spectrum <i>N</i> -Glycan Specificity and Processing of Therapeutic IgG Monoclonal Antibodies by Endoglycosidase S2. ACS Central Science, 2019, 5, 524-538. Antigen-Induced Allosteric Changes in a Human IgG1 Fc Increase Low-Affinity Fcî³ Receptor Binding. Structure, 2020, 28, 516-527.e5. Hydrogen/Deuterium Exchange Kinetics Demonstrate Long Range Allosteric Effects of Thumb Site 2 Inhibitors of Hepatitis C Viral RNA-dependent RNA Polymerase. Journal of Biological Chemistry, 2016,	3.2 11.3 3.3	28 27 23
12 13 14	Acquisition. ACS Infectious Diseases, 2020, 6, 2073-2085. Structure and Dynamics of FosA-Mediated Fosfomycin Resistance in Klebsiella pneumoniae and Escherichia coli. Antimicrobial Agents and Chemotherapy, 2017, 61,. Molecular Basis of Broad Spectrum <i>N</i> Clycan Specificity and Processing of Therapeutic IgG Monoclonal Antibodies by Endoglycosidase S2. ACS Central Science, 2019, 5, 524-538. Antigen-Induced Allosteric Changes in a Human IgG1 Fc Increase Low-Affinity FcÎ3 Receptor Binding. Structure, 2020, 28, 516-527.e5. Hydrogen/Deuterium Exchange Kinetics Demonstrate Long Range Allosteric Effects of Thumb Site 2 Inhibitors of Hepatitis C Viral RNA-dependent RNA Polymerase. Journal of Biological Chemistry, 2016, 291, 10078-10088. Ligand-induced allostery in the interaction of the <i>Pseudomonas aeruginosa</i> Ji>heme binding protein with heme oxygenase. Proceedings of the National Academy of Sciences of the United States of	3.2 11.3 3.3	28 27 23 22

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19	Monitoring dendrimer conformational transition using 19 F and 1 H 2 O NMR. Magnetic Resonance in Chemistry, 2019, 57, 861-872.	1.9	10
20	A temperature-dependent conformational shift in p38α MAPK substrate–binding region associated with changes in substrate phosphorylation profile. Journal of Biological Chemistry, 2019, 294, 12624-12637.	3.4	9
21	Conformational transition of a non-associative fluorinated amphiphile in aqueous solution. II. Conformational transition <i>vs.</i> supramolecular assembly. RSC Advances, 2019, 9, 1956-1966.	3.6	9
22	Molecular Basis of Selective Cytokine Signaling Inhibition by Antibodies Targeting a Shared Receptor. Frontiers in Immunology, 2021, 12, 779100.	4.8	9
23	Modeling the native ensemble of PhuS using enhanced sampling MD and HDX-ensemble reweighting. Biophysical Journal, 2021, 120, 5141-5157.	0.5	7
24	A fusion of the Bacteroides fragilis ferrous iron import proteins reveals a role for FeoA in stabilizing GTP-bound FeoB. Journal of Biological Chemistry, 2022, 298, 101808.	3.4	6
25	Insights into substrate recognition and specificity for IgG by Endoglycosidase S2. PLoS Computational Biology, 2021, 17, e1009103.	3.2	5
26	Hydrogen-Deuterium Exchange Mass Spectrometry (HDX-MS) Centroid Data Measured between 3.6 \hat{A}° C and 25.4 \hat{A}° C for the Fab Fragment of NISTmAb. Journal of Research of the National Institute of Standards and Technology, 2019, 124, 1-7.	1.2	3
27	Interpreting hydrogen-deuterium exchange experiments with molecular simulations: Tutorials and applications of the HDXer ensemble reweighting software [Article v1.0]. Living Journal of Computational Molecular Science, 2022, 3, .	6.4	3
28	Characterization of interaction between blood coagulation factor <scp>VIII</scp> and <scp>LRP1</scp> suggests dynamic binding by alternating complex contacts. Journal of Thrombosis and Haemostasis, 0, , .	3.8	2
29	A structure and function relationship study to identify the impact of the R721G mutation in the human mitochondrial lon protease. Archives of Biochemistry and Biophysics, 2021, 710, 108983.	3.0	1
30	Remodeling KRAS. Structure, 2017, 25, 1323-1324.	3.3	0