

William M Westler

List of Publications by Citations

Source: <https://exaly.com/author-pdf/8521632/william-m-westler-publications-by-citations.pdf>

Version: 2024-04-23

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

22
papers

329
citations

12
h-index

17
g-index

22
ext. papers

407
ext. citations

7.7
avg, IF

2.98
L-index

#	Paper	IF	Citations
22	Integrative NMR for biomolecular research. <i>Journal of Biomolecular NMR</i> , 2016 , 64, 307-32	3	36
21	Comparison of the accuracy of protein solution structures derived from conventional and network-edited NOESY data. <i>Protein Science</i> , 1995 , 4, 2289-99	6.3	35
20	Evidence for a strong hydrogen bond in the catalytic dyad of transition-state analogue inhibitor complexes of chymotrypsin from proton-triton NMR isotope shifts. <i>Journal of the American Chemical Society</i> , 2002 , 124, 4196-7	16.4	27
19	I-PINE web server: an integrative probabilistic NMR assignment system for proteins. <i>Journal of Biomolecular NMR</i> , 2019 , 73, 213-222	3	26
18	Quantum chemical calculations on structural models of the catalytic site of chymotrypsin: comparison of calculated results with experimental data from NMR spectroscopy. <i>Journal of the American Chemical Society</i> , 2002 , 124, 14373-81	16.4	23
17	Amplification of One-Bond ¹ H/ ² H Isotope Effects on ¹⁵ N Chemical Shifts in <i>Clostridium pasteurianum</i> Rubredoxin by Fermi-Contact Effects through Hydrogen Bonds. <i>Journal of the American Chemical Society</i> , 1998 , 120, 4893-4894	16.4	23
16	Forazoline A: Marine-Derived Polyketide with Antifungal In Vivo Efficacy. <i>Angewandte Chemie</i> , 2014 , 126, 11767-11770	3.6	20
15	Unique identifiers for small molecules enable rigorous labeling of their atoms. <i>Scientific Data</i> , 2017 , 4, 170073	8.2	19
14	Spin System Modeling of Nuclear Magnetic Resonance Spectra for Applications in Metabolomics and Small Molecule Screening. <i>Analytical Chemistry</i> , 2017 , 89, 12201-12208	7.8	19
13	Increasing rigor in NMR-based metabolomics through validated and open source tools. <i>Current Opinion in Biotechnology</i> , 2017 , 43, 56-61	11.4	17
12	Usb1 controls U6 snRNP assembly through evolutionarily divergent cyclic phosphodiesterase activities. <i>Nature Communications</i> , 2017 , 8, 497	17.4	16
11	Uncovering a membrane-distal conformation of KRAS available to recruit RAF to the plasma membrane. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 24258-24268	11.5	12
10	Two-dimensional spectra of intact tissue: homonuclear Hartmann-Hahn spectroscopy provides increased sensitivity and information content as compared to COSY. <i>Magnetic Resonance in Medicine</i> , 1990 , 15, 142-51	4.4	11
9	Probabilistic validation of protein NMR chemical shift assignments. <i>Journal of Biomolecular NMR</i> , 2016 , 64, 17-25	3	8
8	NMR studies of retinoid-protein interactions: the conformation of [¹³ C]-beta-ionones bound to beta-lactoglobulin B. <i>Pharmaceutical Research</i> , 1999 , 16, 651-9	4.5	8
7	One- and two-dimensional NMR spectral analysis of the consequences of single amino acid replacements in proteins. <i>Journal of Cellular Biochemistry</i> , 1986 , 30, 291-309	4.7	8
6	Automated evaluation of consistency within the PubChem Compound database. <i>Scientific Data</i> , 2019 , 6, 190023	8.2	7

5	NMRFAM-SDF: a protein structure determination framework. <i>Journal of Biomolecular NMR</i> , 2015 , 62, 481-95	3	4
4	Probabilistic identification of saccharide moieties in biomolecules and their protein complexes. <i>Scientific Data</i> , 2020 , 7, 210	8.2	4
3	Tools for Enhanced NMR-Based Metabolomics Analysis. <i>Methods in Molecular Biology</i> , 2019 , 2037, 413-427	4	3
2	Solution Structural Studies of GTP:Adenosylcobinamide-Phosphateguanylyl Transferase (CobY) from <i>Methanocaldococcus jannaschii</i> . <i>PLoS ONE</i> , 2015 , 10, e0141297	3.7	2
1	Fragment screening targeting Ebola virus nucleoprotein C-terminal domain identifies lead candidates. <i>Antiviral Research</i> , 2020 , 180, 104822	10.8	1