

Geoffrey A Mueller

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1610491/publications.pdf>

Version: 2024-02-01

112
papers

4,055
citations

126858

33
h-index

133188

59
g-index

114
all docs

114
docs citations

114
times ranked

4286
citing authors

#	ARTICLE	IF	CITATIONS
1	A robust and cost-effective method for the production of Val, Leu, Ile (delta 1) methyl-protonated ¹⁵ N-, ¹³ C-, 2H-labeled proteins. <i>Journal of Biomolecular NMR</i> , 1999, 13, 369-374.	1.6	461
2	Orienting domains in proteins using dipolar couplings measured by liquid-state NMR: differences in solution and crystal forms of maltodextrin binding protein loaded with ¹² I-cyclodextrin. <i>Journal of Molecular Biology</i> , 2000, 295, 1265-1273.	2.0	197
3	Global folds of proteins with low densities of NOEs using residual dipolar couplings: application to the 370-residue maltodextrin-binding protein. <i>Journal of Molecular Biology</i> , 2000, 300, 197-212.	2.0	165
4	WHO/IUIS Allergen Nomenclature: Providing a common language. <i>Molecular Immunology</i> , 2018, 100, 3-13.	1.0	162
5	The Crystal Structure of a Major Dust Mite Allergen Der p 2, and its Biological Implications. <i>Journal of Molecular Biology</i> , 2002, 318, 189-197.	2.0	137
6	A synergistic approach to protein crystallization: Combination of a fixed-arm carrier with surface entropy reduction. <i>Protein Science</i> , 2010, 19, 901-913.	3.1	131
7	ZATT (ZNF451)-mediated resolution of topoisomerase 2 DNA-protein cross-links. <i>Science</i> , 2017, 357, 1412-1416.	6.0	127
8	Title is missing!. <i>Journal of Biomolecular NMR</i> , 1999, 14, 333-343.	1.6	122
9	The structure of the dust mite allergen Der p 7 reveals similarities to innate immune proteins. <i>Journal of Allergy and Clinical Immunology</i> , 2010, 125, 909-917.e4.	1.5	99
10	Ara h 2: crystal structure and IgE binding distinguish two subpopulations of peanut allergic patients by epitope diversity. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , 2011, 66, 878-885.	2.7	86
11	Tertiary Structure of the Major House Dust Mite Allergen Der p 2: Sequential and Structural Homologies. <i>Biochemistry</i> , 1998, 37, 12707-12714.	1.2	84
12	NIAID, NIEHS, NHLBI, and MCAN Workshop Report: The indoor environment and childhood asthma implications for home environmental intervention in asthma prevention and management. <i>Journal of Allergy and Clinical Immunology</i> , 2017, 140, 933-949.	1.5	75
13	Dependence of Amino Acid Side Chain ¹³ C Shifts on Dihedral Angle: Application to Conformational Analysis. <i>Journal of the American Chemical Society</i> , 2008, 130, 11097-11105.	6.6	71
14	Serological, genomic and structural analyses of the major mite allergen Der p 23. <i>Clinical and Experimental Allergy</i> , 2016, 46, 365-376.	1.4	69
15	The novel structure of the cockroach allergen Bla g 1 has implications for allergenicity and exposure assessment. <i>Journal of Allergy and Clinical Immunology</i> , 2013, 132, 1420-1426.e9.	1.5	64
16	Identification of Maillard reaction products on peanut allergens that influence binding to the receptor for advanced glycation end products. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , 2013, 68, 1546-1554.	2.7	63
17	New Insights into Cockroach Allergens. <i>Current Allergy and Asthma Reports</i> , 2017, 17, 25.	2.4	63
18	Crystal Structure of Calmodulin Binding Domain of Orai1 in Complex with Ca ²⁺ : Calmodulin Displays a Unique Binding Mode. <i>Journal of Biological Chemistry</i> , 2012, 287, 43030-43041.	1.6	58

#	ARTICLE	IF	CITATIONS
19	The Molecular Basis of Peanut Allergy. <i>Current Allergy and Asthma Reports</i> , 2014, 14, 429.	2.4	58
20	Hydrogen Exchange Nuclear Magnetic Resonance Spectroscopy Mapping of Antibody Epitopes on the House Dust Mite Allergen Der p 2. <i>Journal of Biological Chemistry</i> , 2001, 276, 9359-9365.	1.6	54
21	Solution Structure of the RNase H Domain of the HIV-1 Reverse Transcriptase in the Presence of Magnesium ²⁺ . <i>Biochemistry</i> , 2003, 42, 639-650.	1.2	53
22	Dynamic Characterization of a DNA Repair Enzyme: NMR Studies of [methyl- ¹³ C]Methionine-Labeled DNA Polymerase β . <i>Biochemistry</i> , 2004, 43, 8911-8922.	1.2	53
23	Der p 5 Crystal Structure Provides Insight into the Group 5 Dust Mite Allergens. <i>Journal of Biological Chemistry</i> , 2010, 285, 25394-25401.	1.6	52
24	Analysis of glutathione S-transferase allergen cross-reactivity in a North American population: Relevance for molecular diagnosis. <i>Journal of Allergy and Clinical Immunology</i> , 2015, 136, 1369-1377.	1.5	52
25	Future directions for allergen immunotherapy. <i>Journal of Allergy and Clinical Immunology</i> , 1998, 102, 335-343.	1.5	51
26	Multiple roles of Bet v 1 ligands in allergen stabilization and modulation of endosomal protease activity. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , 2019, 74, 2382-2393.	2.7	51
27	Direct structure refinement of high molecular weight proteins against residual dipolar couplings and carbonyl chemical shift changes upon alignment: an application to maltose binding protein. <i>Journal of Biomolecular NMR</i> , 2001, 21, 31-40.	1.6	50
28	APE2 Zf-GRF facilitates 3'–5' resection of DNA damage following oxidative stress. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 304-309.	3.3	50
29	Expression and Secondary Structure Determination by NMR Methods of the Major House Dust Mite Allergen Der p 2. <i>Journal of Biological Chemistry</i> , 1997, 272, 26893-26898.	1.6	49
30	Revised structure of the AbrB N-terminal domain unifies a diverse superfamily of putative DNA-binding proteins. <i>FEBS Letters</i> , 2005, 579, 5669-5674.	1.3	45
31	Comparison of a Neural Net-Based QSAR Algorithm (PCANN) with Hologram- and Multiple Linear Regression-Based QSAR Approaches: Application to 1,4-Dihydropyridine-Based Calcium Channel Antagonists. <i>Journal of Chemical Information and Computer Sciences</i> , 2001, 41, 505-511.	2.8	43
32	Structural studies of the PARP-1 BRCT domain. <i>BMC Structural Biology</i> , 2011, 11, 37.	2.3	41
33	The Focal Adhesion Targeting Domain of Focal Adhesion Kinase Contains a Hinge Region that Modulates Tyrosine 926 Phosphorylation. <i>Structure</i> , 2004, 12, 881-891.	1.6	37
34	Allergens and their associated small molecule ligands—their dual role in sensitization. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , 2021, 76, 2367-2382.	2.7	36
35	Conformational dependence of ¹³ C shielding and coupling constants for methionine methyl groups. <i>Journal of Biomolecular NMR</i> , 2010, 48, 31-47.	1.6	35
36	A comparison of BRCT domains involved in nonhomologous end-joining: Introducing the solution structure of the BRCT domain of polymerase lambda. <i>DNA Repair</i> , 2008, 7, 1340-1351.	1.3	33

#	ARTICLE	IF	CITATIONS
37	100 Years later: Celebrating the contributions of x-ray crystallography to allergy and clinical immunology. <i>Journal of Allergy and Clinical Immunology</i> , 2015, 136, 29-37.e10.	1.5	33
38	Context matters: TH2 polarization resulting from pollen composition and not from protein-intrinsic allergenicity. <i>Journal of Allergy and Clinical Immunology</i> , 2018, 142, 984-987.e6.	1.5	33
39	Abundance and Stability as Common Properties of Allergens. <i>Frontiers in Allergy</i> , 2021, 2, 769728.	1.2	33
40	Primary Identification, Biochemical Characterization, and Immunologic Properties of the Allergenic Pollen Cyclophilin Cat r 1. <i>Journal of Biological Chemistry</i> , 2014, 289, 21374-21385.	1.6	31
41	Metal-induced DNA translocation leads to DNA polymerase conformational activation. <i>Nucleic Acids Research</i> , 2012, 40, 2974-2983.	6.5	30
42	Structural Aspects of the Allergen-Antibody Interaction. <i>Frontiers in Immunology</i> , 2020, 11, 2067.	2.2	29
43	Solution Structure of the Lyase Domain of Human DNA Polymerase β . <i>Biochemistry</i> , 2003, 42, 9564-9574.	1.2	27
44	Solution structure of the Drosha double-stranded RNA-binding domain. <i>Silence: A Journal of RNA Regulation</i> , 2010, 1, 2.	8.0	26
45	Solution Structure of Polymerase β 's BRCT Domain Reveals an Element Essential for Its Role in Nonhomologous End Joining. <i>Biochemistry</i> , 2007, 46, 12100-12110.	1.2	25
46	Selective unfolding of one Ribonuclease H domain of HIV reverse transcriptase is linked to homodimer formation. <i>Nucleic Acids Research</i> , 2014, 42, 5361-5377.	6.5	25
47	Backbone Dynamics of the RNase H Domain of HIV-1 Reverse Transcriptase. <i>Biochemistry</i> , 2004, 43, 9332-9342.	1.2	24
48	NMR analysis of [methyl- ^{13}C]methionine UvrB from <i>Bacillus caldotenax</i> reveals UvrB domain 4 heterodimer formation in solution. <i>Journal of Molecular Biology</i> , 2007, 373, 282-295.	2.0	24
49	NMR Studies of the Interaction of a Type II Dihydrofolate Reductase with Pyridine Nucleotides Reveal Unexpected Phosphatase and Reductase Activity. <i>Biochemistry</i> , 2003, 42, 11150-11160.	1.2	22
50	NvAssign: protein NMR spectral assignment with NMRView. <i>Bioinformatics</i> , 2004, 20, 1201-1203.	1.8	21
51	Mapping Human Monoclonal IgE Epitopes on the Major Dust Mite Allergen Der p 2. <i>Journal of Immunology</i> , 2020, 205, 1999-2007.	0.4	21
52	Mouse/human chimeric IgG1 and IgG4 antibodies directed to the house dust mite allergen Der p 2: use in quantification of allergen specific IgG. <i>Clinical and Experimental Allergy</i> , 1997, 27, 1095-1102.	1.4	20
53	The Nuclease A Inhibitor Represents a New Variation of the Rare PR-1 Fold. <i>Journal of Molecular Biology</i> , 2002, 320, 771-782.	2.0	20
54	Nuclear Magnetic Resonance Solution Structure of the <i>Escherichia coli</i> DNA Polymerase III β Subunit. <i>Journal of Bacteriology</i> , 2005, 187, 7081-7089.	1.0	19

#	ARTICLE	IF	CITATIONS
55	NMR Structure of AbhN and Comparison with AbrBN. <i>Journal of Biological Chemistry</i> , 2006, 281, 21399-21409.	1.6	19
56	Solution characterization of [methyl-13C]methionine HIV-1 reverse transcriptase by NMR spectroscopy. <i>Antiviral Research</i> , 2009, 84, 205-214.	1.9	19
57	Homodimerization of the p51 Subunit of HIV-1 Reverse Transcriptase. <i>Biochemistry</i> , 2010, 49, 2821-2833.	1.2	19
58	Contributions and Future Directions for Structural Biology in the Study of Allergens. <i>International Archives of Allergy and Immunology</i> , 2017, 174, 57-66.	0.9	19
59	A Human IgE Antibody Binding Site on Der p 2 for the Design of a Recombinant Allergen for Immunotherapy. <i>Journal of Immunology</i> , 2019, 203, 2545-2556.	0.4	19
60	Identification of the amino-terminal fragment of Ara h 1 as a major target of the IgE-binding activity in the basic peanut protein fraction. <i>Clinical and Experimental Allergy</i> , 2020, 50, 401-405.	1.4	19
61	Asymmetric conformational maturation of HIV-1 reverse transcriptase. <i>ELife</i> , 2015, 4, .	2.8	19
62	Complementation of Der p 2-induced histamine release from human basophils sensitized with monoclonal IgE: Not only by IgE, but also by IgG antibodies directed to a nonoverlapping epitope of Der p 2. <i>Journal of Allergy and Clinical Immunology</i> , 1998, 101, 404-409.	1.5	18
63	The N-Terminal Domain of the Drosophila Histone mRNA Binding Protein, SLBP, Is Intrinsically Disordered with Nascent Helical Structure,. <i>Biochemistry</i> , 2004, 43, 9390-9400.	1.2	18
64	Genomic, RNAseq, and Molecular Modeling Evidence Suggests That the Major Allergen Domain in Insects Evolved from a Homodimeric Origin. <i>Genome Biology and Evolution</i> , 2013, 5, 2344-2358.	1.1	18
65	Leucine-rich repeats and calponin homology containing 4 (Lrch4) regulates the innate immune response. <i>Journal of Biological Chemistry</i> , 2019, 294, 1997-2008.	1.6	16
66	Keeping Allergen Names Clear and Defined. <i>Frontiers in Immunology</i> , 2019, 10, 2600.	2.2	16
67	A method for incorporating dipolar couplings into structure calculations in cases of (near) axial symmetry of alignment. <i>Journal of Biomolecular NMR</i> , 2000, 18, 183-188.	1.6	15
68	Are dust mite allergens more abundant and/or more stable than other Dermatophagoides pteronyssinus proteins?. <i>Journal of Allergy and Clinical Immunology</i> , 2017, 139, 1030-1032.e1.	1.5	15
69	Proteases of Dermatophagoides pteronyssinus. <i>International Journal of Molecular Sciences</i> , 2017, 18, 1204.	1.8	14
70	The Draft Genome Assembly of <i>Dermatophagoides pteronyssinus</i> Supports Identification of Novel Allergen Isoforms in <i>Dermatophagoides</i> Species. <i>International Archives of Allergy and Immunology</i> , 2018, 175, 136-146.	0.9	14
71	Hydrophobic ligands influence the structure, stability, and processing of the major cockroach allergen Bla g 1. <i>Scientific Reports</i> , 2019, 9, 18294.	1.6	14
72	Comparison of phytochemical composition of Ginkgo biloba extracts using a combination of non-targeted and targeted analytical approaches. <i>Analytical and Bioanalytical Chemistry</i> , 2020, 412, 6789-6809.	1.9	14

#	ARTICLE	IF	CITATIONS
73	Solution structure and function of YndB, an AHSA1 protein from <i>Bacillus subtilis</i> . <i>Proteins: Structure, Function and Bioinformatics</i> , 2010, 78, 3328-3340.	1.5	13
74	Mechanism of APTX nicked DNA sensing and pleiotropic inactivation in neurodegenerative disease. <i>EMBO Journal</i> , 2018, 37, .	3.5	13
75	The mosquito protein AEG12 displays both cytolytic and antiviral properties via a common lipid transfer mechanism. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	13
76	Phage Like It HOT. <i>Structure</i> , 2004, 12, 2221-2231.	1.6	12
77	Analytical solution to the coupled evolution of multidimensional NMR data. <i>Journal of Biomolecular NMR</i> , 2009, 44, 13-23.	1.6	12
78	Enhanced Approaches for Identifying Amadori Products: Application to Peanut Allergens. <i>Journal of Agricultural and Food Chemistry</i> , 2016, 64, 1406-1413.	2.4	11
79	Human IgE monoclonal antibody recognition of mite allergen Der p 2 defines structural basis of an epitope for IgE cross-linking and anaphylaxis <i>in vivo</i> . , 2022, 1, .		11
80	Metal and ligand binding to the HIV-RNase H active site are remotely monitored by Ile556. <i>Nucleic Acids Research</i> , 2012, 40, 10543-10553.	6.5	10
81	A metabolomic, geographic, and seasonal analysis of the contribution of pollen-derived adenosine to allergic sensitization. <i>Metabolomics</i> , 2016, 12, 1.	1.4	10
82	Unfolding the HIV-1 reverse transcriptase RNase H domain “how to lose a molecular tug-of-war. <i>Nucleic Acids Research</i> , 2016, 44, 1776-1788.	6.5	10
83	Structure, Immunogenicity, and IgE Cross-Reactivity among Walnut and Peanut Vicilin-Buried Peptides. <i>Journal of Agricultural and Food Chemistry</i> , 2022, 70, 2389-2400.	2.4	9
84	Identification of drivers for the metamorphic transition of HIV-1 reverse transcriptase. <i>Biochemical Journal</i> , 2017, 474, 3321-3338.	1.7	7
85	Are allergens more abundant and/or more stable than other proteins in pollens and dust?. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , 2020, 75, 1267-1269.	2.7	7
86	Mixture analyses of air-sampled pollen extracts can accurately differentiate pollen taxa. <i>Atmospheric Environment</i> , 2020, 243, 117746.	1.9	7
87	Removal and Replacement of Endogenous Ligands from Lipid-Bound Proteins and Allergens. <i>Journal of Visualized Experiments</i> , 2021, , .	0.2	7
88	Characterization of the Redox Transition of the XRCC1 N-terminal Domain. <i>Structure</i> , 2014, 22, 1754-1763.	1.6	6
89	Characterization of an anti-Bla g 1 scFv: Epitope mapping and cross-reactivity. <i>Molecular Immunology</i> , 2014, 59, 200-207.	1.0	6
90	Structural Analysis of Recent Allergen-Antibody Complexes and Future Directions. <i>Current Allergy and Asthma Reports</i> , 2019, 19, 17.	2.4	6

#	ARTICLE	IF	CITATIONS
91	Nanobody Paratope Ensembles in Solution Characterized by MD Simulations and NMR. <i>International Journal of Molecular Sciences</i> , 2022, 23, 5419.	1.8	6
92	Antigenic and Molecular Structure of the Mite Allergen Der p 2. <i>International Archives of Allergy and Immunology</i> , 1997, 113, 99-101.	0.9	5
93	NMR assignment of polymerase \hat{I}^2 labeled with 2H , ^{13}C , and ^{15}N in complex with substrate DNA. <i>Biomolecular NMR Assignments</i> , 2007, 1, 33-35.	0.4	5
94	Protein-Mediated Antagonism between HIV Reverse Transcriptase Ligands Nevirapine and MgATP. <i>Biophysical Journal</i> , 2013, 104, 2695-2705.	0.2	5
95	1H , ^{13}C , and ^{15}N NMR assignments for the <i>Bacillus subtilis</i> yndB START domain. <i>Biomolecular NMR Assignments</i> , 2009, 3, 191-194.	0.4	4
96	Transitions in DNA polymerase \hat{I}^2 $\hat{I}^{1/4}$ s-ms dynamics related to substrate binding and catalysis. <i>Nucleic Acids Research</i> , 2018, 46, 7309-7322.	6.5	3
97	NMR assignment of protein side chains using residue-correlated labeling and NOE spectra. <i>Journal of Magnetic Resonance</i> , 2003, 165, 237-247.	1.2	2
98	Evaluation of the allergenic activity of the Glutathione Transferase from <i>Blomia tropicalis</i> (Blo t 8) in a mouse model of airway inflammation. <i>Journal of Allergy and Clinical Immunology</i> , 2019, 143, AB187.	1.5	2
99	Influence of Hydrophobic Cargo Binding on the Structure, Stability, and Allergenicity of the Cockroach Allergen Bla g 1. <i>Journal of Allergy and Clinical Immunology</i> , 2019, 143, AB213.	1.5	2
100	The Der p 7 Crystal Structure Reveals Similarities to Innate Immune Proteins. <i>Journal of Allergy and Clinical Immunology</i> , 2010, 125, AB188.	1.5	1
101	Analysis of GST Allergen Cross-Reactivity in a North American Population for Molecular Diagnosis. <i>Journal of Allergy and Clinical Immunology</i> , 2015, 135, AB187.	1.5	1
102	Structural, Serological, and Genomic Analyses of the Major Mite Allergen Der p 23. <i>Journal of Allergy and Clinical Immunology</i> , 2016, 137, AB267.	1.5	1
103	Mapping human monoclonal IgE epitopes on Der p 2. <i>Journal of Allergy and Clinical Immunology</i> , 2019, 143, AB183.	1.5	1
104	The Cockroach Allergen Bla g 1 Forms Alpha Helical Capsules with an Internal Lipid Binding Cavity: Implications for Allergenicity. <i>Journal of Allergy and Clinical Immunology</i> , 2013, 131, AB16.	1.5	0
105	Epitope Mapping Of An Anti-Bla g 1 ScFv Used For Cockroach Allergen Quantitation. <i>Journal of Allergy and Clinical Immunology</i> , 2014, 133, AB100.	1.5	0
106	Antigenic Analysis Of The Major Cockroach Allergen Bla g 5 and Its Dust Mite Homolog Der p 8. <i>Journal of Allergy and Clinical Immunology</i> , 2014, 133, AB100.	1.5	0
107	Are Dust Mite Allergens More Abundant or More Stable Than Other Dermatophagoides Pteronyssinus Proteins?. <i>Journal of Allergy and Clinical Immunology</i> , 2016, 137, AB268.	1.5	0
108	The NMR structure and IgE Epitopes of Ara h 1 Leader Sequence. <i>Journal of Allergy and Clinical Immunology</i> , 2019, 143, AB240.	1.5	0

#	ARTICLE	IF	CITATIONS
109	Analysis Of IgE Antigenic Determinants On Der p 2 For Design Of Immunotherapy. Journal of Allergy and Clinical Immunology, 2019, 143, AB185.	1.5	0
110	Response to Letter to the Editor regarding "Comparison of phytochemical composition of Ginkgo biloba extracts using a combination of non-targeted and targeted analytical approaches". Analytical and Bioanalytical Chemistry, 2021, 413, 7627-7629.	1.9	0
111	Editorial: Activation of Innate Immunity by Allergens and Allergenic Sources. Frontiers in Allergy, 2021, 2, 800929.	1.2	0
112	Bet v 1 and other birch allergens are more resistant to proteolysis and more abundant than other birch pollen proteins. Allergy: European Journal of Allergy and Clinical Immunology, 2022, 77, 1307-1309.	2.7	0