

Viswanathan V Krishnan

List of Publications by Year in descending order

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Version: 2024-02-01

54
papers

1,390
citations

471509

17
h-index

361022

35
g-index

55
all docs

55
docs citations

55
times ranked

2017
citing authors

#	ARTICLE	IF	CITATIONS
1	Metabolomics Study at the Postharvest Conditions of Cold Storage and Fungicide (Imazalil Sulfate) Treatment in Navel Oranges and Clementine Mandarins. <i>ACS Agricultural Science and Technology</i> , 2022, 2, 79-89.	2.3	2
2	Role of solvent dielectric constant on the enthalpy-entropy compensation in the hindered amide bond rotation. <i>Chemical Physics Letters</i> , 2022, 792, 139412.	2.6	2
3	Evaluation of thrombotic thrombocytopenic purpura and other thrombotic microangiopathies: Lessons learned from a 14-year retrospective study. <i>Therapeutic Apheresis and Dialysis</i> , 2022, , .	0.9	0
4	On the Differential Roles of Mg ²⁺ , Zn ²⁺ , and Cu ²⁺ in the Equilibrium of ¹⁵ N-Methyl-Amino-L-Alanine (BMAA) and its Carbamates. <i>Neurotoxicity Research</i> , 2021, 39, 6-16.	2.7	6
5	NMR based real-time enzyme kinetics on estimating the inhibitory effect of sucralose in the enzymatic conversion of sucrose. <i>Biophysical Chemistry</i> , 2021, 268, 106495.	2.8	2
6	Conformational Ensembles by NMR and MD Simulations in Model Heptapeptides with Select Tri-Peptide Motifs. <i>International Journal of Molecular Sciences</i> , 2021, 22, 1364.	4.1	3
7	Identification of ligand binding sites in intrinsically disordered proteins with a differential binding score. <i>Scientific Reports</i> , 2021, 11, 22583.	3.3	4
8	Role of glycosylation on the ensemble of conformations in the MUC1 immunodominant epitope. <i>Journal of Peptide Science</i> , 2020, 26, e3229.	1.4	3
9	Keto-Enol Tautomerization of Acetylacetone in Mixed Solvents by NMR Spectroscopy. A Physical Chemistry Experiment on the Application of the Onsager-Kirkwood Model for Solvation Thermodynamics. <i>Journal of Chemical Education</i> , 2020, 97, 825-830.	2.3	6
10	Validation of Enthalpy-Entropy Compensation Mechanism in Partial Amide Bond Rotation. <i>ACS Omega</i> , 2020, 5, 9348-9355.	3.5	5
11	NMR spectroscopy analysis reveals differential metabolic responses in arabidopsis roots and leaves treated with a cytokinesis inhibitor. <i>PLoS ONE</i> , 2020, 15, e0241627.	2.5	2
12	Unexpected change in NOE with increasing temperature: Crosstalk between chemical exchange and cross relaxation in a NiN2S2 complex. <i>Chemical Physics Letters</i> , 2019, 715, 160-165.	2.6	1
13	The Ensemble of Conformations of Antifreeze Glycoproteins (AFGP8): A Study Using Nuclear Magnetic Resonance Spectroscopy. <i>Biomolecules</i> , 2019, 9, 235.	4.0	9
14	Molecular Thermodynamics Using Nuclear Magnetic Resonance (NMR) Spectroscopy. <i>Inventions</i> , 2019, 4, 13.	2.5	17
15	Enzymatic conversion of sucrose to glucose and its anomerization by quantitative NMR spectroscopy: Application of a simple consecutive reaction rates approach. <i>Journal of Molecular Structure</i> , 2018, 1153, 187-191.	3.6	3
16	Chemistry and Chemical Equilibrium Dynamics of BMAA and Its Carbamate Adducts. <i>Neurotoxicity Research</i> , 2018, 33, 76-86.	2.7	24
17	Effect of sucralose on the enzyme kinetics of invertase using real-time NMR spectroscopy and progress curve analysis. <i>Carbohydrate Research</i> , 2018, 455, 5-9.	2.3	3
18	Proteomic profiles by multiplex microsphere suspension array. <i>Journal of Immunological Methods</i> , 2018, 461, 1-14.	1.4	12

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19	Metabolomic Analysis of HER2 ⁺ positive Breast Cancer Cells. <i>FASEB Journal</i> , 2018, 32, 658.10.	0.5	0
20	Gender Differences in Bile Acids and Microbiota in Relationship with Gender Dissimilarity in Steatosis Induced by Diet and FXR Inactivation. <i>Scientific Reports</i> , 2017, 7, 1748.	3.3	103
21	Kinetics and thermodynamics of oxidation mediated reaction in L-cysteine and its methyl and ethyl esters in dimethyl sulfoxide-d ₆ by NMR spectroscopy. <i>Journal of Molecular Structure</i> , 2017, 1131, 196-200.	3.6	12
22	Restricted amide rotation with steric hindrance induced multiple conformations. <i>Chemical Physics Letters</i> , 2017, 689, 148-151.	2.6	12
23	Western Diet ⁺ Induced Dysbiosis in Farnesoid X Receptor Knockout Mice Causes Persistent Hepatic Inflammation after Antibiotic Treatment. <i>American Journal of Pathology</i> , 2017, 187, 1800-1813.	3.8	90
24	Upregulation of cystathione β -synthase and p70S6K/S6 in neonatal hypoxic ischemic brain injury. <i>Brain Pathology</i> , 2017, 27, 449-458.	4.1	16
25	Ensemble characterization of an intrinsically disordered FG ⁺ Nup peptide and its F&A mutant in DMSO-d ₆ . <i>Biopolymers</i> , 2017, 108, e23036.	2.4	1
26	Equilibrium Dynamics of β -N-Methylamino-L-Alanine (BMAA) and Its Carbamate Adducts at Physiological Conditions. <i>PLoS ONE</i> , 2016, 11, e0160491.	2.5	11
27	Application of data mining tools for classification of protein structural class from residue based averaged NMR chemical shifts. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2015, 1854, 1545-1552.	2.3	9
28	Real-Time Enzyme Kinetics by Quantitative NMR Spectroscopy and Determination of the Michaelis ⁺ Menten Constant Using the Lambert-W Function. <i>Journal of Chemical Education</i> , 2015, 92, 1943-1948.	2.3	28
29	Data Mining Strategies to Improve Multiplex Microbead Immunoassay Tolerance in a Mouse Model of Infectious Diseases. <i>PLoS ONE</i> , 2015, 10, e0116262.	2.5	10
30	NMR based solvent exchange experiments to understand the conformational preference of intrinsically disordered proteins using FG ⁺ nucleoporin peptide as a model. <i>Biopolymers</i> , 2014, 102, 69-77.	2.4	5
31	Multiplexed measurements of immunomodulator levels in peripheral blood of healthy subjects: Effects of analytical variables based on anticoagulants, age, and gender. , 2014, 86, 426-435.		22
32	Mapping the amino acid properties of constituent nucleoporins onto the yeast nuclear pore complex. <i>Bioinformatics</i> , 2014, 10, 94-97.	0.5	0
33	Radiation damping in modern NMR experiments: Progress and challenges. <i>Progress in Nuclear Magnetic Resonance Spectroscopy</i> , 2013, 68, 41-57.	7.5	91
34	Exploratory Study on Plasma Immunomodulator and Antibody Profiles in Tuberculosis Patients. <i>Vaccine Journal</i> , 2013, 20, 1283-1290.	3.1	19
35	Multiplexed measurements of immunomodulator levels in peripheral blood of healthy subjects: Effects of analytical variables based on anticoagulants, age and gender. , 2013, , n/a-n/a.		15
36	Modulations in restricted amide rotation by steric induced conformational trapping. <i>Chemical Physics Letters</i> , 2012, 523, 124-127.	2.6	11

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37	Plasma Antibody Profiles as Diagnostic Biomarkers for Tuberculosis. <i>Vaccine Journal</i> , 2011, 18, 2148-2153.	3.1	32
38	A Bimodal Distribution of Two Distinct Categories of Intrinsically Disordered Structures with Separate Functions in FG Nucleoporins. <i>Molecular and Cellular Proteomics</i> , 2010, 9, 2205-2224.	3.8	289
39	Characterization of protein secondary structure from NMR chemical shifts. <i>Progress in Nuclear Magnetic Resonance Spectroscopy</i> , 2009, 54, 141-165.	7.5	95
40	Intramolecular Cohesion of Coils Mediated by Phenylalanine-Glycine Motifs in the Natively Unfolded Domain of a Nucleoporin. <i>PLoS Computational Biology</i> , 2008, 4, e1000145.	3.2	48
41	Temperature dependence of protein-hydration hydrodynamics by molecular dynamics simulations. <i>Biophysical Chemistry</i> , 2007, 130, 55-64.	2.8	3
42	Radiation damping in microcoil NMR probes. <i>Journal of Magnetic Resonance</i> , 2006, 179, 294-298.	2.1	10
43	Estimation of protein secondary structure content directly from NMR spectra using an improved empirical correlation with averaged chemical shift. <i>Journal of Structural and Functional Genomics</i> , 2006, 6, 281-285.	1.2	4
44	An evaluation of chemical shift index-based secondary structure determination in proteins: Influence of random coil chemical shifts*. <i>Journal of Biomolecular NMR</i> , 2004, 30, 143-153.	2.8	30
45	Translational dynamics of antifreeze glycoprotein in supercooled water. <i>Biophysical Chemistry</i> , 2004, 110, 223-230.	2.8	12
46	A primer for nuclear magnetic relaxation in liquids. <i>Concepts in Magnetic Resonance</i> , 2003, 17A, 86-116.	1.3	13
47	An Empirical Correlation between Secondary Structure Content and Averaged Chemical Shifts in Proteins. <i>Biophysical Journal</i> , 2003, 84, 1223-1227.	0.5	30
48	Protein structural class identification directly from NMR spectra using averaged chemical shifts. <i>Bioinformatics</i> , 2003, 19, 2054-2064.	4.1	30
49	Solution Structure and Backbone Dynamics of the Human DNA Ligase III \pm BRCT Domain. <i>Biochemistry</i> , 2001, 40, 13158-13166.	2.5	41
50	Novel relaxation compensated method to measure proton exchange rates in biomolecules based on decorrelation of heteronuclear two-spin order. <i>Magnetic Resonance in Chemistry</i> , 2000, 38, 789-794.	1.9	1
51	Dynamics of Cellular Retinoic Acid Binding Protein I on Multiple Time Scales with Implications for Ligand Binding. <i>Biochemistry</i> , 2000, 39, 9119-9129.	2.5	45
52	An improved experimental scheme to measure self-diffusion coefficients of biomolecules with an advantageous use of radiation damping. <i>Chemical Physics Letters</i> , 1999, 302, 317-323.	2.6	19
53	An empirical relationship between rotational correlation time and solvent accessible surface area. <i>J. Biomol. NMR</i> , 1998, 12, 177-182.		44
54	Determination of Oligomeric State of Proteins in Solution from Pulsed-Field-Gradient Self-Diffusion Coefficient Measurements. A Comparison of Experimental, Theoretical, and Hard-Sphere Approximated Values. <i>Journal of Magnetic Resonance</i> , 1997, 124, 468-473.	2.1	72