## Anirban Bhunia

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6583718/publications.pdf

Version: 2024-02-01

		101384	138251
134	4,337	36	58
papers	citations	h-index	g-index
1.40	1.40	1.40	F.(17
140	140	140	5617
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	A rationally designed synthetic antimicrobial peptide against Pseudomonas-associated corneal keratitis: Structure-function correlation. Biophysical Chemistry, 2022, 286, 106802.	1.5	5
2	Atomic-Resolution Structures and Mode of Action of Clinically Relevant Antimicrobial Peptides. International Journal of Molecular Sciences, 2022, 23, 4558.	1.8	11
3	Structural insights into the interaction of antifungal peptides and ergosterol containing fungal membrane. Biochimica Et Biophysica Acta - Biomembranes, 2022, 1864, 183996.	1.4	10
4	Effect of Secondary Structure and Side Chain Length of Hydrophobic Amino Acid Residues on the Antimicrobial Activity and Toxicity of 14â€Residue‣ong de novo AMPs. ChemMedChem, 2021, 16, 355-367.	1.6	23
5	Gut-Brain axis in Parkinson's disease etiology: The role of lipopolysaccharide. Chemistry and Physics of Lipids, 2021, 235, 105029.	1.5	20
6	An explicitly designed paratope of amyloid- $\hat{l}^2$ prevents neuronal apoptosis <i>in vitro</i> and hippocampal damage in rat brain. Chemical Science, 2021, 12, 2853-2862.	3.7	7
7	Hanudatta S. Atreya (1974–2020). Magnetic Resonance in Chemistry, 2021, 59, 201-212.	1.1	O
8	Targeting C-terminal Helical bundle of NCOVID19 Envelope (E) protein. International Journal of Biological Macromolecules, 2021, 175, 131-139.	3.6	7
9	Attenuation of Human Lysozyme Amyloid Fibrillation by ACE Inhibitor Captopril: A Combined Spectroscopy, Microscopy, Cytotoxicity, and Docking Study. Biomacromolecules, 2021, 22, 1910-1920.	2.6	21
10	Inhibition behavior of Sennoside A and Sennoside C on amyloid fibrillation of human lysozyme and its possible mechanism. International Journal of Biological Macromolecules, 2021, 178, 424-433.	3 <b>.</b> 6	26
11	Conformational distortion in a fibril-forming oligomer arrests alpha-Synuclein fibrillation and minimizes its toxic effects. Communications Biology, 2021, 4, 518.	2.0	8
12	Effect of PEGylation on Host Defense Peptide Complexation with Bacterial Lipopolysaccharide. Bioconjugate Chemistry, 2021, 32, 1729-1741.	1.8	8
13	Synthesis and antibacterial study of cell-penetrating peptide conjugated trifluoroacetyl and thioacetyl lysine modified peptides. European Journal of Medicinal Chemistry, 2021, 219, 113447.	2.6	7
14	Solvent Relaxation NMR: A Tool for Real-Time Monitoring Water Dynamics in Protein Aggregation Landscape. ACS Chemical Neuroscience, 2021, 12, 2903-2916.	1.7	8
15	Deciphering the Role of Ion Channels in Early Defense Signaling against Herbivorous Insects. Cells, 2021, 10, 2219.	1.8	9
16	Editorial: Secondary Metabolites and Peptides as Unique Natural Reservoirs of New Therapeutic Leads for Treatment of Cancer and Microbial Infections. Frontiers in Chemistry, 2021, 9, 748180.	1.8	1
17	Zinc oxide nanoparticle interface moderation with tyrosine and tryptophan reverses the pro-amyloidogenic property of the particle. Biochimie, 2021, , .	1.3	1
18	Selfâ $\in$ Assembly and Neurotoxicity of βâ $\in$ Amyloid (21â $\in$ "40) Peptide Fragment: The Regulatory Role of GxxxG Motifs. ChemMedChem, 2020, 15, 293-301.	1.6	16

#	Article	IF	CITATIONS
19	Salt Dependence Conformational Stability of the Dimeric SAM Domain of MAPKKK Ste11 from Budding Yeast: A Native-State H/D Exchange NMR Study. Biochemistry, 2020, 59, 2849-2858.	1.2	3
20	Nonthermal Atmospheric Plasma-Induced Cellular Envelope Damage of <i>Staphylococcus aureus</i> and <i>Candida albicans</i> Biofilms: Spectroscopic and Biochemical Investigations. IEEE Transactions on Plasma Science, 2020, 48, 2768-2776.	0.6	2
21	Highâ€resolution structure of a partially folded insulin aggregation intermediate. Proteins: Structure, Function and Bioinformatics, 2020, 88, 1648-1659.	1.5	13
22	Investigating the inhibitory effects of entacapone on amyloid fibril formation of human lysozyme. International Journal of Biological Macromolecules, 2020, 161, 1393-1404.	3.6	25
23	Host-membrane interacting interface of the SARS coronavirus envelope protein: Immense functional potential of C-terminal domain. Biophysical Chemistry, 2020, 266, 106452.	1.5	41
24	Comparison of Synthetic Neuronal Model Membrane Mimics in Amyloid Aggregation at Atomic Resolution. ACS Chemical Neuroscience, 2020, 11, 1965-1977.	1.7	18
25	7-Methoxytacrine and 2-Aminobenzothiazole Heterodimers: Structure–Mechanism Relationship of Amyloid Inhibitors Based on Rational Design. ACS Chemical Neuroscience, 2020, 11, 715-729.	1.7	10
26	Targeted inhibition of amyloidogenesis using a non-toxic, serum stable strategically designed cyclic peptide with therapeutic implications. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2020, 1868, 140378.	1.1	12
27	Rationally designed antimicrobial peptides: Insight into the mechanism of eleven residue peptides against microbial infections. Biochimica Et Biophysica Acta - Biomembranes, 2020, 1862, 183177.	1.4	21
28	Interaction with zinc oxide nanoparticle kinetically traps $\hat{l}_{\pm}$ -synuclein fibrillation into off-pathway non-toxic intermediates. International Journal of Biological Macromolecules, 2020, 150, 68-79.	3.6	11
29	Molecular Details of a Salt Bridge and Its Role in Insulin Fibrillation by NMR and Raman Spectroscopic Analysis. Journal of Physical Chemistry B, 2020, 124, 1125-1136.	1.2	10
30	Characterization of Antimicrobial Peptide–Membrane Interaction Using All-Atom Molecular Dynamic Simulation. Springer Protocols, 2020, , 163-176.	0.1	2
31	NMR Assisted Antimicrobial Peptide Designing: Structure Based Modifications and Functional Correlation of a Designed Peptide VG16KRKP. Current Medicinal Chemistry, 2020, 27, 1387-1404.	1.2	6
32	Combining Antimicrobial Peptides with Nanotechnology: An Emerging Field in Theranostics. Current Protein and Peptide Science, 2020, 21, 413-428.	0.7	17
33	Structural insights into the combinatorial effects of antimicrobial peptides reveal a role of aromatic–aromatic interactions in antibacterial synergism. Journal of Biological Chemistry, 2019, 294, 14615-14633.	1.6	20
34	Do Catechins (ECG and EGCG) Bind to the Same Site as Thioflavin T (ThT) in Amyloid Fibril? Answer From Saturation Transfer Difference NMR. Natural Product Communications, 2019, 14, 1934578X1984979.	0.2	5
35	Role of non-electrostatic forces in antimicrobial potency of a dengue-virus derived fusion peptide VG16KRKP: Mechanistic insight into the interfacial peptide-lipid interactions. Biochimica Et Biophysica Acta - Biomembranes, 2019, 1861, 798-809.	1.4	13
36	Mitochondrial-membrane association of $\hat{l}_{\pm}$ -synuclein: Pros and cons in consequence of Parkinson's disease pathophysiology. Gene Reports, 2019, 16, 100423.	0.4	4

#	Article	IF	Citations
37	Design, Synthesis, Antibacterial Potential, and Structural Characterization of N-Acylated Derivatives of the Human Autophagy 16 Polypeptide. Bioconjugate Chemistry, 2019, 30, 1998-2010.	1.8	13
38	Enhanced Silkworm Cecropin B Antimicrobial Activity against <i>Pseudomonas aeruginosa</i> from Single Amino Acid Variation. ACS Infectious Diseases, 2019, 5, 1200-1213.	1.8	31
39	Probing transient non-native states in amyloid beta fiber elongation by NMR. Chemical Communications, 2019, 55, 4483-4486.	2.2	46
40	Lipopolysaccharide from Gut Microbiota Modulates α-Synuclein Aggregation and Alters Its Biological Function. ACS Chemical Neuroscience, 2019, 10, 2229-2236.	1.7	73
41	A Peptide-Nanoparticle System with Improved Efficacy against Multidrug Resistant Bacteria. Scientific Reports, 2019, 9, 4485.	1.6	80
42	Nanomedical Relevance of the Intermolecular Interaction Dynamics—Examples from Lysozymes and Insulins. ACS Omega, 2019, 4, 4206-4220.	1.6	11
43	Structural characterization of VapB46 antitoxin from Mycobacterium tuberculosis : insights into VapB46– DNA binding. FEBS Journal, 2019, 286, 1174-1190.	2.2	4
44	Microgels as carriers of antimicrobial peptides â€" Effects of peptide PEGylation. Colloids and Surfaces A: Physicochemical and Engineering Aspects, 2019, 565, 8-15.	2.3	26
45	Cell-Penetrating Peptides as Theranostics Against Impaired Blood-Brain Barrier Permeability: Implications for Pathogenesis and Therapeutic Treatment of Neurodegenerative Disease. Neuromethods, 2019, , 115-136.	0.2	0
46	Sequence specificity of amylin-insulin interaction: a fragment-based insulin fibrillation inhibition study. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2019, 1867, 405-415.	1.1	13
47	Application of tungsten disulfide quantum dot-conjugated antimicrobial peptides in bio-imaging and antimicrobial therapy. Colloids and Surfaces B: Biointerfaces, 2019, 176, 360-370.	2.5	40
48	Multitude NMR studies of $\hat{l}\pm$ -synuclein familial mutants: probing their differential aggregation propensities. Chemical Communications, 2018, 54, 3605-3608.	2.2	33
49	Insulin–eukaryotic model membrane interaction: Mechanistic insight of insulin fibrillation and membrane disruption. Biochimica Et Biophysica Acta - Biomembranes, 2018, 1860, 1917-1926.	1.4	17
50	Structural insights of a self-assembling 9-residue peptide from the C-terminal tail of the SARS corona virus E-protein in DPC and SDS micelles: A combined high and low resolution spectroscopic study. Biochimica Et Biophysica Acta - Biomembranes, 2018, 1860, 335-346.	1.4	17
51	Insights into the Mechanism of Antimicrobial Activity of Seven-Residue Peptides. Journal of Medicinal Chemistry, 2018, 61, 7614-7629.	2.9	19
52	Binding Moiety Mapping by Saturation Transfer Difference NMR. Methods in Molecular Biology, 2018, 1824, 49-65.	0.4	5
53	Nonproductive Binding Modes as a Prominent Feature of $A\hat{l}^2$ (sub>40 Fiber Elongation: Insights from Molecular Dynamics Simulation. Journal of Chemical Information and Modeling, 2018, 58, 1576-1586.	2.5	11
54	Evidences for zinc (II) and copper (II) ion interactions with Mycobacterium leprae HSP18: Effect on its structure and chaperone function. Journal of Inorganic Biochemistry, 2018, 188, 62-75.	1.5	6

#	Article	IF	Citations
55	Inhibition and Degradation of Amyloid Beta (A $\hat{1}^2$ 40) Fibrillation by Designed Small Peptide: A Combined Spectroscopy, Microscopy, and Cell Toxicity Study. ACS Chemical Neuroscience, 2017, 8, 718-722.	1.7	44
56	Reduced Lipid Bilayer Thickness Regulates the Aggregation and Cytotoxicity of Amyloid- $\hat{l}^2$ . Journal of Biological Chemistry, 2017, 292, 4638-4650.	1.6	145
57	Structural and Dynamic Insights into a Glycine-Mediated Short Analogue of a Designed Peptide in Lipopolysaccharide Micelles: Correlation Between Compact Structure and Anti-Endotoxin Activity. Biochemistry, 2017, 56, 1348-1362.	1.2	15
58	Tryptophan end-tagging for promoted lipopolysaccharide interactions and anti-inflammatory effects. Scientific Reports, 2017, 7, 212.	1.6	13
59	Synthesis of novel muramic acid derivatives and their interaction with lysozyme: Action of lysozyme revisited. Journal of Colloid and Interface Science, 2017, 498, 395-404.	5.0	12
60	Conformational Aspects of High Content Packing of Antimicrobial Peptides in Polymer Microgels. ACS Applied Materials & Samp; Interfaces, 2017, 9, 40094-40106.	4.0	17
61	Lysozyme's lectin-like characteristics facilitates its immune defense function. Quarterly Reviews of Biophysics, 2017, 50, e9.	2.4	29
62	Multivalent gold nanoparticle–peptide conjugates for targeting intracellular bacterial infections. Nanoscale, 2017, 9, 14074-14093.	2.8	60
63	Membrane perturbing activities and structural properties of the frog-skin derived peptide Esculentin-1a(1-21)NH2 and its Diastereomer Esc(1-21)-1c: Correlation with their antipseudomonal and cytotoxic activity. Biochimica Et Biophysica Acta - Biomembranes, 2017, 1859, 2327-2339.	1.4	27
64	Accelerated molecular dynamics simulation analysis of MSI-594 in a lipid bilayer. Physical Chemistry Chemical Physics, 2017, 19, 19289-19299.	1.3	46
65	An Approach Towards Structure Based Antimicrobial Peptide Design for Use in Development of Transgenic Plants: A Strategy for Plant Disease Management. Current Medicinal Chemistry, 2017, 24, 1350-1364.	1.2	13
66	Ribavirin suppresses bacterial virulence by targeting LysR-type transcriptional regulators. Scientific Reports, 2016, 6, 39454.	1.6	23
67	Identification of modes of interactions between 9-aminoacridine hydrochloride hydrate and serum proteins by low and high resolution spectroscopy and molecular modeling. RSC Advances, 2016, 6, 53454-53468.	1.7	21
68	Evidence for Inhibition of Lysozyme Amyloid Fibrillization by Peptide Fragments from Human Lysozyme: A Combined Spectroscopy, Microscopy, and Docking Study. Biomacromolecules, 2016, 17, 1998-2009.	2.6	35
69	Inhibition of Insulin Amyloid Fibrillation by a Novel Amphipathic Heptapeptide. Journal of Biological Chemistry, 2016, 291, 23545-23556.	1.6	62
70	Deciphering the role of the AT-rich interaction domain and the HMG-box domain of ARID-HMG proteins of Arabidopsis thaliana. Plant Molecular Biology, 2016, 92, 371-388.	2.0	16
71	Enhanced stability and activity of an antimicrobial peptide in conjugation with silver nanoparticle. Journal of Colloid and Interface Science, 2016, 483, 385-393.	5.0	97
72	Structural Elucidation of the Cell-Penetrating Penetratin Peptide in Model Membranes at the Atomic Level: Probing Hydrophobic Interactions in the Blood–Brain Barrier. Biochemistry, 2016, 55, 4982-4996.	1.2	24

#	Article	IF	Citations
73	Mode of Action of a Designed Antimicrobial Peptide: High Potency against Cryptococcus neoformans. Biophysical Journal, 2016, 111, 1724-1737.	0.2	37
74	Expedient synthesis of the pentasaccharide repeating unit of the O-antigen of Escherichia coli O86 and its conformational analysis. Glycoconjugate Journal, 2016, 33, 887-896.	1.4	5
75	Biophysical insights into the membrane interaction of the core amyloid-forming Aβ <sub>40</sub> fragment K16–K28 and its role in the pathogenesis of Alzheimer's disease. Physical Chemistry Chemical Physics, 2016, 18, 16890-16901.	1.3	16
76	Role of Aromatic Amino Acids in Lipopolysaccharide and Membrane Interactions of Antimicrobial Peptides for Use in Plant Disease Control. Journal of Biological Chemistry, 2016, 291, 13301-13317.	1.6	46
77	Structure and Dynamics of Antifreeze Protein–Model Membrane Interactions: A Combined Spectroscopic and Molecular Dynamics Study. Journal of Physical Chemistry B, 2016, 120, 902-914.	1.2	20
78	C -cinnamoyl glycosides as a new class of anti-filarial agents. European Journal of Medicinal Chemistry, 2016, 114, 308-317.	2.6	14
79	NMR structure and binding of esculentin-1a $(1\hat{a}\in 21)$ NH 2 and its diastereomer to lipopolysaccharide: Correlation with biological functions. Biochimica Et Biophysica Acta - Biomembranes, 2016, 1858, 800-812.	1.4	16
80	Amyloid- $\hat{l}^2$ adopts a conserved, partially folded structure upon binding to zwitterionic lipid bilayers prior to amyloid formation. Chemical Communications, 2016, 52, 882-885.	2.2	66
81	Designing potent antimicrobial peptides by disulphide linked dimerization and N-terminal lipidation to increase antimicrobial activity and membrane perturbation: Structural insights into lipopolysaccharide binding. Journal of Colloid and Interface Science, 2016, 461, 335-345.	5.0	41
82	Influence of a curcumin derivative on hIAPP aggregation in the absence and presence of lipid membranes. Chemical Communications, 2016, 52, 942-945.	2.2	63
83	Reactivity of Metal-Free and Metal-Associated Amyloid $\hat{l}^2$ with Glycosylated Polyphenols and Their Esterified Derivatives. Scientific Reports, 2015, 5, 17842.	1.6	44
84	An Alternative Phosphorylation Switch in Integrin β2 (CD18) Tail for Dok1 Binding. Scientific Reports, 2015, 5, 11630.	1.6	15
85	Antimicrobial Peptides and their Pore/Ion Channel Properties in Neutralization of Pathogenic Microbes. Current Topics in Medicinal Chemistry, 2015, 16, 46-53.	1.0	39
86	Biophysical Characterization of Essential Phosphorylation at the Flexible C-Terminal Region of C-Raf with 14-3-3ζ Protein. PLoS ONE, 2015, 10, e0135976.	1.1	9
87	Editorial (Thematic Issue: Antimicrobial Peptides in Medicinal Chemistry: Advances and Applications). Current Topics in Medicinal Chemistry, 2015, 16, 2-3.	1.0	4
88	Interaction of ATP with a Small Heat Shock Protein from Mycobacterium leprae: Effect on Its Structure and Function. PLoS Neglected Tropical Diseases, 2015, 9, e0003661.	1.3	13
89	Antimicrobial Peptides: Insights into Membrane Permeabilization, Lipopolysaccharide Fragmentation and Application in Plant Disease Control. Scientific Reports, 2015, 5, 11951.	1.6	70
90	Probing the role of Proline in the antimicrobial activity and lipopolysaccharide binding of indolicidin. Journal of Colloid and Interface Science, 2015, 452, 148-159.	5.0	22

#	Article	IF	Citations
91	Self-Assembly of a Nine-Residue Amyloid-Forming Peptide Fragment of SARS Corona Virus E-Protein: Mechanism of Self Aggregation and Amyloid-Inhibition of hIAPP. Biochemistry, 2015, 54, 2249-2261.	1.2	50
92	Detergent-Type Membrane Fragmentation by MSI-78, MSI-367, MSI-594, and MSI-843 Antimicrobial Peptides and Inhibition by Cholesterol: A Solid-State Nuclear Magnetic Resonance Study. Biochemistry, 2015, 54, 1897-1907.	1.2	55
93	Will It Be Beneficial To Simulate the Antifreeze Proteins at Ice Freezing Condition or at Lower Temperature?. Journal of Physical Chemistry B, 2015, 119, 11485-11495.	1.2	12
94	Biophysical and biochemical aspects of antifreeze proteins: Using computational tools to extract atomistic information. Progress in Biophysics and Molecular Biology, 2015, 119, 194-204.	1.4	15
95	Membrane disruptive antimicrobial activities of human $\hat{l}^2$ -defensin-3 analogs. European Journal of Medicinal Chemistry, 2015, 91, 91-99.	2.6	44
96	Double GC:GC Mismatch in dsDNA Enhances Local Dynamics Retaining the DNA Footprint: A Highâ€Resolution NMR Study. ChemMedChem, 2014, 9, 2059-2064.	1.6	14
97	Linear synthesis and conformational analysis of the pentasaccharide repeating unit of the cell wall O-antigen of Escherichia coli O13. Carbohydrate Research, 2014, 391, 9-15.	1.1	3
98	Potent Î <sup>3</sup> -secretase inhibitors/modulators interact with amyloid-Î <sup>2</sup> fibrils but do not inhibit fibrillation: A high-resolution NMR study. Biochemical and Biophysical Research Communications, 2014, 447, 590-595.	1.0	17
99	Acetylation of Gly1 and Lys2 Promotes Aggregation of Human $\hat{I}^3$ D-Crystallin. Biochemistry, 2014, 53, 7269-7282.	1.2	26
100	Synthesis of the pentasaccharide repeating unit of the O-antigen of Escherichia coli O175 using one-pot glycosylations and its conformational analysis. Tetrahedron, 2014, 70, 9262-9267.	1.0	2
101	Sequence context induced antimicrobial activity: insight into lipopolysaccharide permeabilization. Molecular BioSystems, 2014, 10, 1596-1612.	2.9	30
102	Synthesis of the tetrasaccharide repeating unit of the O-antigen of the Escherichia coli O69 strain and its conformational analysis. RSC Advances, 2014, 4, 37079-37084.	1.7	3
103	Indolicidin Targets Duplex DNA: Structural and Mechanistic Insight through a Combination of Spectroscopy and Microscopy. ChemMedChem, 2014, 9, 2052-2058.	1.6	<b>7</b> 5
104	Convergent Synthesis and Conformational Analysis of the Hexasaccharide Repeating Unit of the ⟨i⟩O⟨ i⟩â€Antigen of ⟨i⟩Shigella flexneri⟨ i⟩ Serotype 1d. European Journal of Organic Chemistry, 2014, 2014, 4577-4584.	1.2	20
105	Human cathelicidin peptide LL37 binds telomeric G-quadruplex. Molecular BioSystems, 2013, 9, 1833.	2.9	25
106	Novel G-quadruplex stabilizing agents: in-silico approach and dynamics. Journal of Biomolecular Structure and Dynamics, 2013, 31, 1497-1518.	2.0	15
107	Use of a Small Peptide Fragment as an Inhibitor of Insulin Fibrillation Process: A Study by High and Low Resolution Spectroscopy. PLoS ONE, 2013, 8, e72318.	1.1	64
108	NMR Structure of Temporin-1 Ta in Lipopolysaccharide Micelles: Mechanistic Insight into Inactivation by Outer Membrane. PLoS ONE, 2013, 8, e72718.	1.1	31

7

#	Article	IF	Citations
109	Structural determinants of the specificity of a membrane binding domain of the scaffold protein Ste5 of budding yeast: Implications in signaling by the scaffold protein in MAPK pathway. Biochimica Et Biophysica Acta - Biomembranes, 2012, 1818, 1250-1260.	1.4	8
110	NMR Structure, Localization, and Vesicle Fusion of Chikungunya Virus Fusion Peptide. Biochemistry, 2012, 51, 7863-7872.	1.2	16
111	Solution Structures, Dynamics, and Ice Growth Inhibitory Activity of Peptide Fragments Derived from an Antarctic Yeast Protein. PLoS ONE, 2012, 7, e49788.	1.1	21
112	Applications of saturation transfer difference NMR in biological systems. Drug Discovery Today, 2012, 17, 505-513.	3.2	126
113	Mapping residue-specific contacts of polymyxin B with lipopolysaccharide by saturation transfer difference NMR: Insights into outer-membrane disruption and endotoxin neutralization. Biopolymers, 2011, 96, 273-287.	1.2	29
114	NMR Structures and Interactions of Temporin-1Tl and Temporin-1Tb with Lipopolysaccharide Micelles. Journal of Biological Chemistry, 2011, 286, 24394-24406.	1.6	84
115	NMR Structure of Pardaxin, a Pore-forming Antimicrobial Peptide, in Lipopolysaccharide Micelles. Journal of Biological Chemistry, 2010, 285, 3883-3895.	1.6	123
116	Structure, Interactions, and Antibacterial Activities of MSI-594 Derived Mutant Peptide MSI-594F5A in Lipopolysaccharide Micelles: Role of the Helical Hairpin Conformation in Outer-Membrane Permeabilization. Journal of the American Chemical Society, 2010, 132, 18417-18428.	6.6	104
117	Why Structurally Different Cyclic Peptides Can Be Glycomimetics of the HNK-1 Carbohydrate Antigen. Journal of the American Chemical Society, 2010, 132, 96-105.	6.6	32
118	Micelle-bound structures and dynamics of the hinge deleted analog of melittin and its diastereomer: Implications in cell selective lysis by d-amino acid containing antimicrobial peptides. Biochimica Et Biophysica Acta - Biomembranes, 2010, 1798, 128-139.	1.4	31
119	Functional and structural characterization of the talin FOF1 domain. Biochemical and Biophysical Research Communications, 2010, 391, 159-165.	1.0	3
120	Designed $\hat{l}^2$ -Boomerang Antiendotoxic and Antimicrobial Peptides. Journal of Biological Chemistry, 2009, 284, 21991-22004.	1.6	94
121	NMR Solution Conformations and Interactions of Integrin $\hat{l}\pm L\hat{l}^22$ Cytoplasmic Tails. Journal of Biological Chemistry, 2009, 284, 3873-3884.	1.6	31
122	Helical Hairpin Structure of a Potent Antimicrobial Peptide MSIâ€594 in Lipopolysaccharide Micelles by NMR Spectroscopy. Chemistry - A European Journal, 2009, 15, 2036-2040.	1.7	89
123	Lipopolysaccharide bound structures of the active fragments of fowlicidinâ€1, a cathelicidin family of antimicrobial and antiendotoxic peptide from chicken, determined by transferred nuclear overhauser effect spectroscopy. Biopolymers, 2009, 92, 9-22.	1.2	56
124	NMR structural studies of the Ste11 SAM domain in the dodecyl phosphocholine micelle. Proteins: Structure, Function and Bioinformatics, 2009, 74, 328-343.	1.5	24
125	Consistent Bioactive Conformation of the Neu5Acα(2â†'3)Gal Epitope Upon Lectin Binding. ChemBioChem, 2008, 9, 2941-2945.	1.3	20
126	Berberine Targets Assembly of Escherichia coli Cell Division Protein FtsZ. Biochemistry, 2008, 47, 3225-3234.	1.2	209

#	ARTICLE	IF	CITATION
127	Interactions of a designed peptide with lipopolysaccharide: Bound conformation and anti-endotoxic activity. Biochemical and Biophysical Research Communications, 2008, 369, 853-857.	1.0	22
128	Equilibrium Unfolding of the Dimeric SAM Domain of MAPKKK Ste11 from the Budding Yeast:  Role of the Interfacial Residues in Structural Stability and Binding. Biochemistry, 2008, 47, 651-659.	1.2	8
129	Structural and thermodynamic analyses of the interaction between melittin and lipopolysaccharide. Biochimica Et Biophysica Acta - Biomembranes, 2007, 1768, 3282-3291.	1.4	58
130	High-Resolution Solution Structure of a Designed Peptide Bound to Lipopolysaccharide:  Transferred Nuclear Overhauser Effects, Micelle Selectivity, and Anti-Endotoxic Activity, Biochemistry, 2007, 46, 5864-5874.	1.2	49
131	Inhibition of bacterial cell division protein FtsZ by cinnamaldehyde. Biochemical Pharmacology, 2007, 74, 831-840.	2.0	213
132	Chelerythrine and Sanguinarine Dock at Distinct Sites on BclXL that are Not the Classic BH3 Binding Cleft. Journal of Molecular Biology, 2006, 364, 536-549.	2.0	58
133	Saturation transfer difference NMR and computational modeling of a sialoadhesin–sialyl lactose complex. Carbohydrate Research, 2004, 339, 259-267.	1.1	37
134	Horseradish peroxidase catalyzed degradation of industrially important dyes. Biotechnology and Bioengineering, 2001, 72, 562-567.	1.7	184