

Chandra S Verma

List of Publications by Year in Descending Order

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

39
papers

470
citations

11
h-index

20
g-index

45
ext. papers

659
ext. citations

5.9
avg, IF

3.8
L-index

#	Paper	IF	Citations
39	Molecular descriptors suggest stapling as a strategy for optimizing membrane permeability of cyclic peptides.. <i>Journal of Chemical Physics</i> , 2022 , 156, 065101	3.9	
38	How well does molecular simulation reproduce environment-specific conformations of the intrinsically disordered peptides PLP, TP2 and ONEG?. <i>Chemical Science</i> , 2022 , 13, 1957-1971	9.4	2
37	Discovery of cell active macrocyclic peptides with on-target inhibition of KRAS signaling.. <i>Chemical Science</i> , 2021 , 12, 15975-15987	9.4	11
36	Liquid-Liquid Phase Separation of Short Histidine- and Tyrosine-Rich Peptides: Sequence Specificity and Molecular Topology. <i>Journal of Physical Chemistry B</i> , 2021 , 125, 6776-6790	3.4	4
35	Computational Methods and Tools in Antimicrobial Peptide Research. <i>Journal of Chemical Information and Modeling</i> , 2021 , 61, 3172-3196	6.1	11
34	Computational Design of Macrocyclic Binders of S100B(β) Novel Peptide Theranostics. <i>Molecules</i> , 2021 , 26,	4.8	2
33	A chemical biology approach reveals a dependency of glioblastoma on biotin distribution. <i>Science Advances</i> , 2021 , 7, eabf6033	14.3	2
32	Macrocyclization of an all-d linear helical peptide imparts cellular permeability. <i>Chemical Science</i> , 2020 , 11, 5577-5591	9.4	17
31	De-risking Drug Discovery of Intracellular Targeting Peptides: Screening Strategies to Eliminate False-Positive Hits. <i>ACS Medicinal Chemistry Letters</i> , 2020 , 11, 1993-2001	4.3	11
30	DStabilize: A Web Resource to Generate Mirror Images of Biomolecules. <i>Structure</i> , 2020 , 28, 1358-1360.	5.2	
29	Deciphering the mechanistic effects of eIF4E phosphorylation on mRNA-cap recognition. <i>Protein Science</i> , 2020 , 29, 1373-1386	6.3	9
28	DisorderOrder Interplay of a Barnacle Cement Protein Triggered by Interactions with Calcium and Carbonate Ions: A Molecular Dynamics Study. <i>Chemistry of Materials</i> , 2020 , 32, 8845-8859	9.6	5
27	Molecular modeling and interaction between Arabidopsis sulfite oxidase and the GW motif of Turnip crinkle virus coat protein. <i>Virology</i> , 2020 , 551, 64-74	3.6	0
26	Dissecting the Molecular Mechanism of Colistin Resistance in -1 Bacteria. <i>Journal of Chemical Information and Modeling</i> , 2020 , 60, 4975-4984	6.1	4
25	Extending the Martini Coarse-Grained Force Field to -Glycans. <i>Journal of Chemical Information and Modeling</i> , 2020 , 60, 3864-3883	6.1	7
24	Critical Role for Cold Shock Protein YB-1 in Cytokinesis. <i>Cancers</i> , 2020 , 12,	6.6	4
23	Straightforward Incorporation of Multiple Ligand Types into Molecular Dynamics Simulations for Efficient Binding Site Detection and Characterization. <i>Journal of Chemical Theory and Computation</i> , 2020 , 16, 6633-6644	6.4	6

22	Molecular basis of dengue virus serotype 2 morphological switch from 29°C to 37°C. <i>PLoS Pathogens</i> , 2019 , 15, e1007996	7.6	8
21	Structural insights reveal a recognition feature for tailoring hydrocarbon stapled-peptides against the eukaryotic translation initiation factor 4E protein. <i>Chemical Science</i> , 2019 , 10, 2489-2500	9.4	12
20	Targeted covalent inhibitors of MDM2 using electrophile-bearing stapled peptides. <i>Chemical Communications</i> , 2019 , 55, 7914-7917	5.8	12
19	Stereoisomerism of stapled peptide inhibitors of the p53-Mdm2 interaction: an assessment of synthetic strategies and activity profiles. <i>Chemical Science</i> , 2019 , 10, 6457-6466	9.4	12
18	Characterization of Hydration Properties in Structural Ensembles of Biomolecules. <i>Journal of Chemical Information and Modeling</i> , 2019 , 59, 3316-3329	6.1	3
17	Exploring Gatekeeper Mutations in EGFR through Computer Simulations. <i>Journal of Chemical Information and Modeling</i> , 2019 , 59, 2850-2858	6.1	3
16	Inhibiting S100B for Activating Wild-Type p53: Design of Stapled Peptides. <i>ACS Omega</i> , 2019 , 4, 5335-5344	3.4	5
15	Efficient development of stable and highly functionalised peptides targeting the CK2 protein-protein interaction. <i>Chemical Science</i> , 2019 , 10, 5056-5063	9.4	20
14	Discovering Putative Protein Targets of Small Molecules: A Study of the p53 Activator Nutlin. <i>Journal of Chemical Information and Modeling</i> , 2019 , 59, 1529-1546	6.1	9
13	A Novel Determinant of PSMD9 PDZ Binding Guides the Evolution of the First Generation of Super Binding Peptides. <i>Biochemistry</i> , 2019 , 58, 3422-3433	3.2	6
12	Effects of Single Nucleotide Polymorphisms on the Binding of Afatinib to EGFR: A Potential Patient Stratification Factor Revealed by Modeling Studies. <i>Journal of Chemical Information and Modeling</i> , 2019 , 59, 309-315	6.1	2
11	Molecular Insights into the Membrane Affinities of Model Hydrophobes. <i>ACS Omega</i> , 2018 , 3, 2498-2507	3.9	7
10	Optimization of Selective Mitogen-Activated Protein Kinase Interacting Kinases 1 and 2 Inhibitors for the Treatment of Blast Crisis Leukemia. <i>Journal of Medicinal Chemistry</i> , 2018 , 61, 4348-4369	8.3	23
9	Comparison of Charge Derivation Methods Applied to Amino Acid Parameterization. <i>ACS Omega</i> , 2018 , 3, 4664-4673	3.9	4
8	Conformational Transitions of Melittin between Aqueous and Lipid Phases: Comparison of Simulations with Experiments. <i>Journal of Physical Chemistry B</i> , 2018 , 122, 8698-8705	3.4	6
7	A Funneled Conformational Landscape Governs Flavivirus Fusion Peptide Interaction with Lipid Membranes. <i>Journal of Chemical Theory and Computation</i> , 2018 , 14, 3920-3932	6.4	6
6	Modeling the full length HIV-1 Gag polyprotein reveals the role of its p6 subunit in viral maturation and the effect of non-cleavage site mutations in protease drug resistance. <i>Journal of Biomolecular Structure and Dynamics</i> , 2018 , 36, 4366-4377	3.6	22
5	Stapled peptides with improved potency and specificity that activate p53. <i>ACS Chemical Biology</i> , 2013 , 8, 506-12	4.9	169

4	Stabilizing the eIF4G1 α helix increases its binding affinity with eIF4E: implications for peptidomimetic design strategies. <i>Journal of Molecular Biology</i> , 2011 , 405, 736-53	6.5	19
3	C-terminal substitution of MDM2 interacting peptides modulates binding affinity by distinctive mechanisms. <i>PLoS ONE</i> , 2011 , 6, e24122	3.7	24
2	Critical role for cold shock protein YB-1 in cytokinesis		1
1	TRAF4 inhibits bladder cancer progression by promoting BMP/SMAD signalling pathway		1