

# Lalith Perera

## List of Publications by Year in descending order

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125  
papers

24,371  
citations

71061

41  
h-index

19169

118  
g-index

131  
all docs

131  
docs citations

131  
times ranked

25582  
citing authors

#	ARTICLE	IF	CITATIONS
1	A smooth particle mesh Ewald method. <i>Journal of Chemical Physics</i> , 1995, 103, 8577-8593.	1.2	18,266
2	New tricks for modelers from the crystallography toolkit: the particle mesh Ewald algorithm and its use in nucleic acid simulations. <i>Structure</i> , 1999, 7, R55-R60.	1.6	571
3	Molecular Dynamics Simulation of Sodium Dodecyl Sulfate Micelle in Water: Micellar Structural Characteristics and Counterion Distribution. <i>Journal of Physical Chemistry B</i> , 2002, 106, 3788-3793.	1.2	334
4	Many-body effects in molecular dynamics simulations of $\text{Na}^+(\text{H}_2\text{O})_n$ and $\text{Cl}^-(\text{H}_2\text{O})_n$ clusters. <i>Journal of Chemical Physics</i> , 1991, 95, 1954-1963.	1.2	322
5	Molecular Dynamics Simulations of Sodium Dodecyl Sulfate Micelle in Water: The Behavior of Water. <i>Journal of Physical Chemistry B</i> , 2002, 106, 10902-10907.	1.2	173
6	Towards accurate solvation dynamics of divalent cations in water using the polarizable amoeba force field: From energetics to structure. <i>Journal of Chemical Physics</i> , 2006, 125, 054511.	1.2	169
7	Phenobarbital Indirectly Activates the Constitutive Active Androstane Receptor (CAR) by Inhibition of Epidermal Growth Factor Receptor Signaling. <i>Science Signaling</i> , 2013, 6, ra31.	1.6	163
8	Structures of $\text{Cl}^-(\text{H}_2\text{O})_n$ and $\text{F}^-(\text{H}_2\text{O})_n$ ( $n=2,3,\dots,15$ ) clusters. <i>Molecular dynamics computer simulations. Journal of Chemical Physics</i> , 1994, 100, 3085-3093.	1.2	154
9	Effect of the treatment of long-range forces on the dynamics of ions in aqueous solutions. <i>Journal of Chemical Physics</i> , 1995, 102, 450-456.	1.2	148
10	Plasminogen Alleles Influence Susceptibility to Invasive Aspergillosis. <i>PLoS Genetics</i> , 2008, 4, e1000101.	1.5	145
11	Uncovering the polymerase-induced cytotoxicity of an oxidized nucleotide. <i>Nature</i> , 2015, 517, 635-639.	13.7	133
12	Potential SARS-CoV-2 main protease inhibitors. <i>Drug Discovery Today</i> , 2021, 26, 804-816.	3.2	128
13	GATA3-dependent cellular reprogramming requires activation-domain dependent recruitment of a chromatin remodeler. <i>Genome Biology</i> , 2016, 17, 36.	3.8	121
14	Structure and dynamics of $\text{Cl}^-(\text{H}_2\text{O})_{20}$ clusters: The effect of the polarizability and the charge of the ion. <i>Journal of Chemical Physics</i> , 1992, 96, 8288-8294.	1.2	118
15	The Origin of the Hydration Interaction of Lipid Bilayers from MD Simulation of Dipalmitoylphosphatidylcholine Membranes in Gel and Liquid Crystalline Phases. <i>Langmuir</i> , 1995, 11, 4519-4531.	1.6	117
16	Dephosphorylation of Threonine 38 Is Required for Nuclear Translocation and Activation of Human Xenobiotic Receptor CAR (NR1I3). <i>Journal of Biological Chemistry</i> , 2009, 284, 34785-34792.	1.6	117
17	Dynamics of ion solvation in a Stockmayer fluid. <i>Journal of Chemical Physics</i> , 1992, 96, 3092-3101.	1.2	104
18	The solvation of $\text{Cl}^-$ , $\text{Br}^-$ , and $\text{I}^-$ in acetonitrile clusters: Photoelectron spectroscopy and molecular dynamics simulations. <i>Journal of Chemical Physics</i> , 1996, 105, 2675-2685.	1.2	103

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19	Stabilization energies of Cl <sup>-</sup> , Br <sup>-</sup> , and I <sup>-</sup> ions in water clusters. <i>Journal of Chemical Physics</i> , 1993, 99, 4222-4224.	1.2	100
20	Spectral shifts and structural classes in microsolutions of rare gas clusters containing a molecular chromophore. <i>Journal of Chemical Physics</i> , 1990, 93, 4884-4897.	1.2	95
21	Cryo-EM structures of the SARS-CoV-2 endoribonuclease Nsp15 reveal insight into nuclease specificity and dynamics. <i>Nature Communications</i> , 2021, 12, 636.	5.8	80
22	Charge localization in negative ion dynamics: Effect on caging of Br <sup>-</sup> 2 in Ar <sub>n</sub> and (CO <sub>2</sub> ) <sub>n</sub> clusters. <i>Journal of Chemical Physics</i> , 1989, 90, 7354-7368.	1.2	79
23	Surface solvation for an ion in a water cluster. <i>Journal of Chemical Physics</i> , 2005, 122, 024513.	1.2	78
24	Characterization of Estrogenic and Androgenic Activities for Bisphenol A-like Chemicals (BPs): In Vitro Estrogen and Androgen Receptors Transcriptional Activation, Gene Regulation, and Binding Profiles. <i>Toxicological Sciences</i> , 2019, 172, 23-37.	1.4	76
25	Differential <i>in Vitro</i> Biological Action, Coregulator Interactions, and Molecular Dynamic Analysis of Bisphenol A (BPA), BPAF, and BPS Ligand-ER± Complexes. <i>Environmental Health Perspectives</i> , 2018, 126, 017012.	2.8	74
26	Structure and Dynamics of Zymogen Human Blood Coagulation Factor X. <i>Biophysical Journal</i> , 2002, 82, 1190-1206.	0.2	73
27	GATA3 zinc finger 2 mutations reprogram the breast cancer transcriptional network. <i>Nature Communications</i> , 2018, 9, 1059.	5.8	72
28	Role of Water in the Hydration Force Acting between Lipid Bilayers. <i>Langmuir</i> , 1996, 12, 2625-2629.	1.6	71
29	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
30	Enthalpies of formation and stabilization energies of Br <sup>-</sup> (H <sub>2</sub> O) <sub>n</sub> (n=1,2, ..., 15) clusters. Comparisons between molecular dynamics computer simulations and experiment. <i>Chemical Physics Letters</i> , 1994, 218, 377-382.	1.2	65
31	Reaction Mechanism of the $\beta$ Subunit of E. coli DNA Polymerase III: Insights into Active Site Metal Coordination and Catalytically Significant Residues. <i>Journal of the American Chemical Society</i> , 2009, 131, 1550-1556.	6.6	64
32	Cube to cage transitions in (H <sub>2</sub> O) <sub>n</sub> (n=12, 16, and 20). <i>Journal of Chemical Physics</i> , 1996, 105, 3715-3721.	1.2	63
33	Solution Structure of the Dickerson DNA Dodecamer Containing a Single Ribonucleotide. <i>Biochemistry</i> , 2012, 51, 2407-2416.	1.2	56
34	An Ancient Family of RNA-Binding Proteins: Still Important!. <i>Trends in Biochemical Sciences</i> , 2017, 42, 285-296.	3.7	55
35	The structure of water at platinum/water interfaces Molecular dynamics computer simulations. <i>Surface Science</i> , 1995, 335, 401-415.	0.8	54
36	Simple formulas for improved point-charge electrostatics in classical force fields and hybrid quantum mechanical/molecular mechanical embedding. <i>International Journal of Quantum Chemistry</i> , 2008, 108, 1905-1912.	1.0	53

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37	Catalytic mechanism of human DNA polymerase $\beta$ with Mg <sup>2+</sup> and Mn <sup>2+</sup> from ab initio quantum mechanical/molecular mechanical studies. <i>DNA Repair</i> , 2008, 7, 1824-1834.	1.3	52
38	Template strand scrunching during DNA gap repair synthesis by human polymerase $\beta$ . <i>Nature Structural and Molecular Biology</i> , 2009, 16, 967-972.	3.6	49
39	Requirement for transient metal ions revealed through computational analysis for DNA polymerase going in reverse. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E5228-36.	3.3	49
40	Molecular Insights into DNA Polymerase Deterrents for Ribonucleotide Insertion. <i>Journal of Biological Chemistry</i> , 2011, 286, 31650-31660.	1.6	45
41	Binding of 5-phospho-D-arabinonohydroxamate and 5-phospho-D-arabinonate inhibitors to zinc phosphomannose isomerase from <i>Candida albicans</i> studied by polarizable molecular mechanics and quantum mechanics. <i>Journal of Computational Chemistry</i> , 2007, 28, 938-957.	1.5	44
42	Binding of bisphenol A, bisphenol AF, and bisphenol S on the androgen receptor: Coregulator recruitment and stimulation of potential interaction sites. <i>Toxicology in Vitro</i> , 2017, 44, 287-302.	1.1	44
43	Characterization of SARS2 Nsp15 nuclease activity reveals it's mad about U. <i>Nucleic Acids Research</i> , 2021, 49, 10136-10149.	6.5	44
44	Amino Acid Substitution in the Active Site of DNA Polymerase $\beta$ Explains the Energy Barrier of the Nucleotidyl Transfer Reaction. <i>Journal of the American Chemical Society</i> , 2013, 135, 8078-8088.	6.6	40
45	Thermally Induced Structural Changes in F-(H <sub>2</sub> O) <sub>11</sub> and Cl-(H <sub>2</sub> O) <sub>11</sub> Clusters: $\hat{A}$ Molecular Dynamics Computer Simulations. <i>The Journal of Physical Chemistry</i> , 1996, 100, 1350-1356.	2.9	39
46	Molecular Mechanism of Substrate Specificity for Heparan Sulfate 2-O-Sulfotransferase. <i>Journal of Biological Chemistry</i> , 2014, 289, 13407-13418.	1.6	39
47	Phylogenetic Distribution and Evolution of the Linked RNA-Binding and NOT1-Binding Domains in the Tristetraprolin Family of Tandem CCCH Zinc Finger Proteins. <i>Journal of Interferon and Cytokine Research</i> , 2014, 34, 297-306.	0.5	38
48	Energetics and structure in I <sub>2</sub> (CO <sub>2</sub> ) <sub>n</sub> clusters. <i>Zeitschrift für Physik D-Atoms Molecules and Clusters</i> , 1991, 20, 173-175.	1.0	37
49	Ultrafast solvation dynamics in a Stockmayer fluid. <i>Journal of Chemical Physics</i> , 1992, 97, 5253-5254.	1.2	35
50	Conformational dependence of <sup>13</sup> C shielding and coupling constants for methionine methyl groups. <i>Journal of Biomolecular NMR</i> , 2010, 48, 31-47.	1.6	35
51	Free energy profiles for lithium(1+) and iodide ions approaching the platinum(100) surface: a molecular dynamics study. <i>The Journal of Physical Chemistry</i> , 1993, 97, 13803-13806.	2.9	33
52	Mitochondrial single-stranded DNA binding protein novel de novo SSBP1 mutation in a child with single large-scale mtDNA deletion (SLSMD) clinically manifesting as Pearson, Kearns-Sayre, and Leigh syndromes. <i>PLoS ONE</i> , 2019, 14, e0221829.	1.1	32
53	Modeling Zymogen Protein C. <i>Biophysical Journal</i> , 2000, 79, 2925-2943.	0.2	31
54	Phosphorylated Nuclear Receptor CAR Forms a Homodimer To Repress Its Constitutive Activity for Ligand Activation. <i>Molecular and Cellular Biology</i> , 2017, 37, .	1.1	31

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55	Cooperative damage recognition by UvrA and UvrB: Identification of UvrA residues that mediate DNA binding. <i>DNA Repair</i> , 2008, 7, 392-404.	1.3	29
56	Molecular mechanisms for the regulation of histone mRNA stem-loop binding protein by phosphorylation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E2937-46.	3.3	29
57	Largazole Analogues Embodying Radical Changes in the Depsipeptide Ring: Development of a More Selective and Highly Potent Analogue. <i>Journal of Medicinal Chemistry</i> , 2016, 59, 10642-10660.	2.9	29
58	Engineering of betabellin-15d: A 64 residue beta sheet protein that forms long narrow multimeric fibrils. <i>Protein Science</i> , 1998, 7, 1545-1554.	3.1	28
59	An all-atom solution-equilibrated model for human extrinsic blood coagulation complex (sTF-VIIa-Xa): a protein-protein docking and molecular dynamics refinement study. <i>Journal of Thrombosis and Haemostasis</i> , 2003, 1, 2577-2588.	1.9	27
60	Proposed structural models of human factor Va and prothrombinase. <i>Journal of Thrombosis and Haemostasis</i> , 2008, 6, 83-89.	1.9	27
61	Revealing the role of the product metal in DNA polymerase $\beta$ catalysis. <i>Nucleic Acids Research</i> , 2017, 45, gkw1363.	6.5	27
62	Mobility of stretched water. <i>Journal of Chemical Physics</i> , 1993, 98, 9859-9862.	1.2	25
63	A finite field method for calculating molecular polarizability tensors for arbitrary multipole rank. <i>Journal of Computational Chemistry</i> , 2011, 32, 3283-3295.	1.5	24
64	Reversal of DNA damage induced Topoisomerase 2 DNA-protein crosslinks by Tdp2. <i>Nucleic Acids Research</i> , 2016, 44, 3829-3844.	6.5	23
65	Identification of the effector domain of biglycan that facilitates BMP-2 osteogenic function. <i>Scientific Reports</i> , 2018, 8, 7022.	1.6	23
66	Ion solvation in water clusters. <i>Zeitschrift für Physik D-Atoms Molecules and Clusters</i> , 1993, 26, 166-168.	1.0	22
67	Modeling Human Zymogen Factor IX. <i>Thrombosis and Haemostasis</i> , 2001, 85, 596-603.	1.8	22
68	Determining the endocrine disruption potential of industrial chemicals using an integrative approach: Public databases, in vitro exposure, and modeling receptor interactions. <i>Environment International</i> , 2019, 131, 104969.	4.8	22
69	Heparan Sulfate Biosynthesis: A Theoretical Study of the Initial Sulfation Step by N-Deacetylase/N-Sulfotransferase. <i>Biophysical Journal</i> , 2000, 79, 2909-2917.	0.2	21
70	Four loops of the catalytic domain of factor VIIa mediate the effect of the first EGF-like domain substitution on factor VIIa catalytic activity. Edited by R. Huber. <i>Journal of Molecular Biology</i> , 2001, 307, 1503-1517.	2.0	20
71	The discovery of new coding alleles of human CYP26A1 that are potentially defective in the metabolism of all-trans retinoic acid and their assessment in a recombinant cDNA expression system. <i>Pharmacogenetics and Genomics</i> , 2007, 17, 169-180.	0.7	20
72	Trans-Cis Isomerization of Proline 22 in Bovine Prothrombin Fragment 1: A Surprising Result of Structural Characterization. <i>Biochemistry</i> , 1998, 37, 10920-10927.	1.2	19

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73	Reversal of drug resistance by JS-K and nitric oxide in ABCB1- and ABCG2-expressing multi-drug resistant human tumor cells. <i>Biomedicine and Pharmacotherapy</i> , 2019, 120, 109468.	2.5	19
74	Asymmetric conformational maturation of HIV-1 reverse transcriptase. <i>ELife</i> , 2015, 4, .	2.8	19
75	Atomic forces for geometry-dependent point multipole and Gaussian multipole models. <i>Journal of Computational Chemistry</i> , 2010, 31, 2702-2713.	1.5	18
76	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
77	Mutational and Structural Analysis of the Tandem Zinc Finger Domain of Tristetraprolin. <i>Journal of Biological Chemistry</i> , 2014, 289, 565-580.	1.6	18
78	The tandem zinc finger RNA binding domain of members of the tristetraprolin protein family. <i>Wiley Interdisciplinary Reviews RNA</i> , 2019, 10, e1531.	3.2	17
79	Unusual Fragmentation Pathways in Collagen Glycopeptides. <i>Journal of the American Society for Mass Spectrometry</i> , 2013, 24, 1072-1081.	1.2	16
80	The Drosophila Tis11 Protein and Its Effects on mRNA Expression in Flies. <i>Journal of Biological Chemistry</i> , 2014, 289, 35042-35060.	1.6	16
81	Synthesis and biological evaluation of largazole analogues with modified surface recognition cap groups. <i>European Journal of Medicinal Chemistry</i> , 2014, 86, 528-541.	2.6	16
82	Predicted solution structure of zymogen human coagulation FVII. <i>Journal of Computational Chemistry</i> , 2002, 23, 35-47.	1.5	15
83	HPAM: Hirshfeld partitioned atomic multipoles. <i>Computer Physics Communications</i> , 2012, 183, 390-397.	3.0	15
84	Functional Equivalence of an Evolutionarily Conserved RNA Binding Module. <i>Journal of Biological Chemistry</i> , 2015, 290, 24413-24423.	1.6	15
85	Explicit Water Near the Catalytic I Helix Thr in the Predicted Solution Structure of CYP2A4. <i>Biophysical Journal</i> , 2003, 84, 57-68.	0.2	14
86	Binuclear manganese(II) complexes in biological systems. <i>Molecular Physics</i> , 2007, 105, 2893-2898.	0.8	14
87	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
88	Probing the Structural Changes in the Light Chain of Human Coagulation Factor VIIa Due to Tissue Factor Association. <i>Biophysical Journal</i> , 1999, 77, 99-113.	0.2	13
89	Total Synthesis and Selective Activity of a New Class of Conformationally Restrained Epothilones. <i>Chemistry - A European Journal</i> , 2008, 14, 570-581.	1.7	13
90	Transmembrane Domain Interactions and Residue Proline 378 Are Essential for Proper Structure, Especially Disulfide Bond Formation, in the Human Vitamin K-Dependent $\Gamma^3$ -Glutamyl Carboxylase. <i>Biochemistry</i> , 2008, 47, 6301-6310.	1.2	13

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91	Modeling of the DNA-binding site of yeast Pms1 by mass spectrometry. <i>DNA Repair</i> , 2011, 10, 454-465.	1.3	13
92	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
93	What causes the enhancement of activity of factor VIIa by tissue factor?. <i>Journal of Thrombosis and Haemostasis</i> , 2006, 4, 2726-2729.	1.9	11
94	Ligand induced dissociation of the AR homodimer precedes AR monomer translocation to the nucleus. <i>Scientific Reports</i> , 2019, 9, 16734.	1.6	11
95	A proposed structural model of human protein Z. <i>Journal of Thrombosis and Haemostasis</i> , 2007, 5, 1558-1561.	1.9	10
96	Hiding in Plain Sight: The Bimetallic Magnesium Covalent Bond in Enzyme Active Sites. <i>Inorganic Chemistry</i> , 2017, 56, 313-320.	1.9	10
97	Structural basis for proficient oxidized ribonucleotide insertion in double strand break repair. <i>Nature Communications</i> , 2021, 12, 5055.	5.8	10
98	NCX-4040, a Unique Nitric Oxide Donor, Induces Reversal of Drug-Resistance in Both ABCB1- and ABCG2-Expressing Multidrug Human Cancer Cells. <i>Cancers</i> , 2021, 13, 1680.	1.7	9
99	Structural Characterization of the Conformational Change in Calbindin-D28k upon Calcium Binding Using Differential Surface Modification Analyzed by Mass Spectrometry. <i>Biochemistry</i> , 2009, 48, 8603-8614.	1.2	8
100	Probing Dominant Negative Behavior of Glucocorticoid Receptor $\alpha$ through a Hybrid Structural and Biochemical Approach. <i>Molecular and Cellular Biology</i> , 2018, 38, .	1.1	8
101	A ubiquitin-like domain is required for stabilizing the N-terminal ATPase module of human SMCHD1. <i>Communications Biology</i> , 2019, 2, 255.	2.0	8
102	Identification of drivers for the metamorphic transition of HIV-1 reverse transcriptase. <i>Biochemical Journal</i> , 2017, 474, 3321-3338.	1.7	7
103	A Bioactive Resveratrol Trimer from the Stem Bark of the Sri Lankan Endemic Plant <i>Vateria copallifera</i> . <i>Journal of Natural Products</i> , 2018, 81, 1693-1700.	1.5	7
104	ESR1 Mutations Associated With Estrogen Insensitivity Syndrome Change Conformation of Ligand-Receptor Complex and Altered Transcriptome Profile. <i>Endocrinology</i> , 2020, 161, .	1.4	7
105	Phosphopeptide interactions of the Nbs1 N-terminal FHA-BRCT1/2 domains. <i>Scientific Reports</i> , 2021, 11, 9046.	1.6	7
106	Probing the mechanisms of two exonuclease domain mutators of DNA polymerase $\beta$ . <i>Nucleic Acids Research</i> , 2022, 50, 962-974.	6.5	7
107	A structural exposé of noncanonical molecular reactivity within the protein tyrosine phosphatase WPD loop. <i>Nature Communications</i> , 2022, 13, 2231.	5.8	7
108	Computational study of the putative active form of protein Z (PZa): Sequence design and structural modeling. <i>Protein Science</i> , 2008, 17, 1354-1361.	3.1	6

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109	Characterization of an anti-Bla g 1 scFv: Epitope mapping and cross-reactivity. <i>Molecular Immunology</i> , 2014, 59, 200-207.	1.0	6
110	A new class of cytotoxic agents targets tubulin and disrupts microtubule dynamics. <i>Bioorganic Chemistry</i> , 2021, 116, 105297.	2.0	6
111	A reconsideration of the evidence for structural reorganization in FVII zymogen. <i>Journal of Thrombosis and Haemostasis</i> , 2005, 3, 1543-1545.	1.9	5
112	Applications of Quantum Mechanical/Molecular Mechanical Methods to the Chemical Insertion Step of DNA and RNA Polymerization. <i>Advances in Protein Chemistry and Structural Biology</i> , 2014, 97, 83-113.	1.0	5
113	Mechanisms of SSBP1 variants in mitochondrial disease: Molecular dynamics simulations reveal stable tetramers with altered DNA binding surfaces. <i>DNA Repair</i> , 2021, 107, 103212.	1.3	4
114	A post-transcriptional regulon controlled by TtpA, the single tristetraprolin family member expressed in <i>Dictyostelium discoideum</i> . <i>Nucleic Acids Research</i> , 2021, 49, 11920-11937.	6.5	3
115	Pharmacophore optimization of imidazole chalcones to modulate microtubule dynamics. <i>Bioorganic Chemistry</i> , 2022, 122, 105700.	2.0	3
116	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
117	Molecular Dynamics Computer Simulations of Aqueous Solution/Platinum Interface. , 1994, , 101-118.		1
118	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
119	A smooth particle mesh Ewald method. , 0, .		1
120	Early Unfolding Response of a Stable Protein Domain to Environmental Changes. <i>Journal of Physical Chemistry A</i> , 2004, 108, 9834-9840.	1.1	0
121	Preferential DNA Polymerase $\beta$ Reverse Reaction with Imidodiphosphate. <i>ACS Omega</i> , 2020, 5, 15317-15324.	1.6	0
122	Solvation Dynamics in a Stockmayer Fluid. , 1993, , 461-483.		0
123	Abstract 964: GATA3 modulates chromatin structure to establish active enhancers in breast cancer cells. , 2015, , .		0
124	SAT-205 ESR1 Q375H and R394H Mutants Associated with Estrogen Insensitivity Syndrome Mediate Genome-Wide Genetic and Epigenetic Aberrances. <i>Journal of the Endocrine Society</i> , 2019, 3, .	0.1	0
125	Structural and functional consequences of SMCHD1 mutations associated with arhinia and muscular dystrophy. <i>FASEB Journal</i> , 2019, 33, 493.5.	0.2	0