

# Kathleen B Hall

## List of Publications by Year in descending order

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31  
papers

1,048  
citations

586496

16  
h-index

488211

31  
g-index

135  
all docs

135  
docs citations

135  
times ranked

1630  
citing authors

#	ARTICLE	IF	CITATIONS
1	The SARS-CoV-2 nucleocapsid protein is dynamic, disordered, and phase separates with RNA. <i>Nature Communications</i> , 2021, 12, 1936.	5.8	334
2	Reweighting of molecular simulations with explicit-solvent SAXS restraints elucidates ion-dependent RNA ensembles. <i>Nucleic Acids Research</i> , 2021, 49, e84-e84.	6.5	25
3	Ribosomal Protein L11 Selectively Stabilizes a Tertiary Structure of the GTPase Center rRNA Domain. <i>Journal of Molecular Biology</i> , 2020, 432, 991-1007.	2.0	7
4	Divalent ions tune the kinetics of a bacterial GTPase center rRNA folding transition from secondary to tertiary structure. <i>Rna</i> , 2018, 24, 1828-1838.	1.6	20
5	Molecular principles underlying dual RNA specificity in the <i>Drosophila</i> SNF protein. <i>Nature Communications</i> , 2018, 9, 2220.	5.8	7
6	Computational Assessment of Potassium and Magnesium Ion Binding to a Buried Pocket in GTPase-Associating Center RNA. <i>Journal of Physical Chemistry B</i> , 2017, 121, 451-462.	1.2	15
7	RNA and Proteins: Mutual Respect. <i>F1000Research</i> , 2017, 6, 345.	0.8	8
8	Nucleobases Undergo Dynamic Rearrangements during RNA Tertiary Folding. <i>Journal of Molecular Biology</i> , 2016, 428, 4490-4502.	2.0	5
9	Divalent Ion Dependent Conformational Changes in an RNA Stem-Loop Observed by Molecular Dynamics. <i>Journal of Chemical Theory and Computation</i> , 2016, 12, 3382-3389.	2.3	48
10	Mighty tiny. <i>Rna</i> , 2015, 21, 630-631.	1.6	21
11	2-Aminopurine Fluorescence as a Probe of Local RNA Structure and Dynamics and Global Folding. <i>Methods in Enzymology</i> , 2015, 558, 99-124.	0.4	7
12	Formation of Tertiary Interactions during rRNA GTPase Center Folding. <i>Journal of Molecular Biology</i> , 2015, 427, 2799-2815.	2.0	6
13	Effect of Loop Composition on the Stability and Folding Kinetics of RNA Hairpins with Large Loops. <i>Biochemistry</i> , 2015, 54, 1886-1896.	1.2	10
14	Stem-Loop V of Varkud Satellite RNA Exhibits Characteristics of the Mg <sup>2+</sup> Bound Structure in the Presence of Monovalent Ions. <i>Journal of Physical Chemistry B</i> , 2015, 119, 12355-12364.	1.2	22
15	Climbing the vertebrate branch of U1A/U2B <sup>ε</sup> protein evolution. <i>Rna</i> , 2014, 20, 1035-1045.	1.6	10
16	Protein binding cannot subdue a lively RNA. <i>Nature</i> , 2014, 506, 303-304.	18.7	3
17	Linkage and Allostery in snRNP Protein/RNA Complexes. <i>Biochemistry</i> , 2014, 53, 3529-3539.	1.2	14
18	RNA does the folding dance of twist, turn, stack. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 16706-16707.	3.3	8

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19	Spectroscopic Probes of RNA Structure and Dynamics. <i>Methods in Molecular Biology</i> , 2012, 875, 67-84.	0.4	6
20	Interactions between PTB RRMs Induce Slow Motions and Increase RNA Binding Affinity. <i>Journal of Molecular Biology</i> , 2010, 397, 260-277.	2.0	18
21	The Domains of Polypyrimidine Tract Binding Protein Have Distinct RNA Structural Preferences. <i>Biochemistry</i> , 2009, 48, 2063-2074.	1.2	36
22	2-Aminopurine as a Probe of RNA Conformational Transitions. <i>Methods in Enzymology</i> , 2009, 469, 269-285.	0.4	37
23	RNA in motion. <i>Current Opinion in Chemical Biology</i> , 2008, 12, 612-618.	2.8	53
24	Dynamics of the IRE RNA hairpin loop probed by 2-aminopurine fluorescence and stochastic dynamics simulations. <i>Rna</i> , 2004, 10, 34-47.	1.6	47
25	RNA-protein interactions. <i>Current Opinion in Structural Biology</i> , 2002, 12, 283-288.	2.6	89
26	Thermodynamics of 2-ribose substitutions in UUCG tetraloops. <i>Rna</i> , 2001, 7, 44-53.	1.6	27
27	Spatial Orientation and Dynamics of the U1A Proteins in the U1A~UTR Complex. <i>Biochemistry</i> , 2000, 39, 7320-7329.	1.2	4
28	Global and local dynamics of the human U1A protein determined by tryptophan fluorescence. <i>Protein Science</i> , 1999, 8, 2110-2120.	3.1	14
29	A Model of the Iron Responsive Element RNA Hairpin Loop Structure Determined from NMR and Thermodynamic Data. <i>Biochemistry</i> , 1996, 35, 13586-13596.	1.2	70
30	Thermodynamic Comparison of the Salt Dependence of Natural RNA Hairpins and RNA Hairpins with Non-Nucleotide Spacers. <i>Biochemistry</i> , 1996, 35, 14665-14670.	1.2	42
31	Contribution of the tyrosines to the structure and function of the human U1A N-terminal RNA binding domain. <i>Protein Science</i> , 1996, 5, 1567-1583.	3.1	34