

# Ravi Radhakrishnan

## List of Publications by Year in descending order

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Version: 2024-02-01

94  
papers

5,658  
citations

172207

29  
h-index

82410

72  
g-index

103  
all docs

103  
docs citations

103  
times ranked

9093  
citing authors

#	ARTICLE	IF	CITATIONS
1	Exosomal PD-L1 contributes to immunosuppression and is associated with anti-PD-1 response. <i>Nature</i> , 2018, 560, 382-386.	13.7	1,836
2	Effects of confinement on freezing and melting. <i>Journal of Physics Condensed Matter</i> , 2006, 18, R15-R68.	0.7	614
3	ErbB3/HER3 intracellular domain is competent to bind ATP and catalyze autophosphorylation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 7692-7697.	3.3	395
4	ALK Mutations Confer Differential Oncogenic Activation and Sensitivity to ALK Inhibition Therapy in Neuroblastoma. <i>Cancer Cell</i> , 2014, 26, 682-694.	7.7	302
5	Erlotinib binds both inactive and active conformations of the EGFR tyrosine kinase domain. <i>Biochemical Journal</i> , 2012, 448, 417-423.	1.7	228
6	Orchestration of cooperative events in DNA synthesis and repair mechanism unraveled by transition path sampling of DNA polymerase $\beta$ 's closing. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 5970-5975.	3.3	146
7	Computational model for nanocarrier binding to endothelium validated using in vivo, in vitro, and atomic force microscopy experiments. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 16530-16535.	3.3	116
8	Reduction of Nanoparticle Avidity Enhances the Selectivity of Vascular Targeting and PET Detection of Pulmonary Inflammation. <i>ACS Nano</i> , 2013, 7, 2461-2469.	7.3	94
9	Exo70 Generates Membrane Curvature for Morphogenesis and Cell Migration. <i>Developmental Cell</i> , 2013, 26, 266-278.	3.1	88
10	Regulation of actin assembly by PI(4,5)P2 and other inositol phospholipids: An update on possible mechanisms. <i>Biochemical and Biophysical Research Communications</i> , 2018, 506, 307-314.	1.0	82
11	Mesoscale computational studies of membrane bilayer remodeling by curvature-inducing proteins. <i>Physics Reports</i> , 2014, 543, 1-60.	10.3	71
12	Fidelity Discrimination in DNA Polymerase $\beta$ : Differing Closing Profiles for a Mismatched (G:A) versus Matched (G:C) Base Pair. <i>Journal of the American Chemical Society</i> , 2005, 127, 13245-13252.	6.6	69
13	Regulation of DNA Repair Fidelity by Molecular Checkpoints: $\beta$ Gates in DNA Polymerase $\beta$ 's Substrate Selection. <i>Biochemistry</i> , 2006, 45, 15142-15156.	1.2	66
14	Biomolecular free energy profiles by a shooting/umbrella sampling protocol, $\beta$ BOLAS. <i>Journal of Chemical Physics</i> , 2004, 121, 2436-2444.	1.2	62
15	Nanoparticle Brownian motion and hydrodynamic interactions in the presence of flow fields. <i>Physics of Fluids</i> , 2011, 23, 73602-7360215.	1.6	60
16	Structural Insights into Pseudokinase Domains of Receptor Tyrosine Kinases. <i>Molecular Cell</i> , 2020, 79, 390-405.e7.	4.5	56
17	Systems biology and physical biology of clathrin-mediated endocytosis. <i>Integrative Biology (United Kingdom)</i> 11, 0784314. <a href="#">DOI: 10.1039/c1ib00055g</a>	0.6	55
18	Correct and incorrect nucleotide incorporation pathways in DNA polymerase $\beta$ . <i>Biochemical and Biophysical Research Communications</i> , 2006, 350, 521-529.	1.0	52

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19	Coarse-Grained Models for Protein-Cell Membrane Interactions. <i>Polymers</i> , 2013, 5, 890-936.	2.0	51
20	Minimal Mesoscale Model for Protein-Mediated Vesiculation in Clathrin-Dependent Endocytosis. <i>PLoS Computational Biology</i> , 2010, 6, e1000926.	1.5	47
21	The Role of Glycocalyx in Nanocarrier-Cell Adhesion Investigated Using a Thermodynamic Model and Monte Carlo Simulations. <i>Journal of Physical Chemistry C</i> , 2007, 111, 15848-15856.	1.5	44
22	Multivalent Binding of Nanocarrier to Endothelial Cells under Shear Flow. <i>Biophysical Journal</i> , 2011, 101, 319-326.	0.2	41
23	Quantum and All-Atom Molecular Dynamics Simulations of Protonation and Divalent Ion Binding to Phosphatidylinositol 4,5-Bisphosphate (PIP <sub>2</sub> ). <i>Journal of Physical Chemistry B</i> , 2013, 117, 8322-8329.	1.2	38
24	Biophysics of membrane curvature remodeling at molecular and mesoscopic lengthscales. <i>Journal of Physics Condensed Matter</i> , 2018, 30, 273001.	0.7	35
25	ULK1 phosphorylates Exo70 to suppress breast cancer metastasis. <i>Nature Communications</i> , 2020, 11, 117.	5.8	35
26	Curvature-undulation coupling as a basis for curvature sensing and generation in bilayer membranes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E5117-24.	3.3	34
27	Mechanisms that determine nanocarrier targeting to healthy versus inflamed lung regions. <i>Nanomedicine: Nanotechnology, Biology, and Medicine</i> , 2017, 13, 1495-1506.	1.7	34
28	Counterion-mediated pattern formation in membranes containing anionic lipids. <i>Advances in Colloid and Interface Science</i> , 2014, 208, 177-188.	7.0	33
29	Understanding and Controlling Food Protein Structure and Function in Foods: Perspectives from Experiments and Computer Simulations. <i>Annual Review of Food Science and Technology</i> , 2020, 11, 365-387.	5.1	33
30	A multiscale modeling approach to investigate molecular mechanisms of pseudokinase activation and drug resistance in the HER3/ErbB3 receptor tyrosine kinase signaling network. <i>Molecular BioSystems</i> , 2011, 7, 2066.	2.9	32
31	Atomistic Insights into Regulatory Mechanisms of the HER2 Tyrosine Kinase Domain: A Molecular Dynamics Study. <i>Biophysical Journal</i> , 2009, 96, 2321-2334.	0.2	31
32	Multiscale Modeling of Functionalized Nanocarriers in Targeted Drug Delivery. <i>Current Nanoscience</i> , 2011, 7, 727-735.	0.7	29
33	Phase separation in confined systems. <i>Reports on Progress in Physics</i> , 2000, 63, 727-727.	8.1	28
34	Molecular dynamics analysis of conserved hydrophobic and hydrophilic bond-interaction networks in ErbB family kinases. <i>Biochemical Journal</i> , 2011, 436, 241-251.	1.7	27
35	Multivalent Binding of a Ligand-Coated Particle: Role of Shape, Size, and Ligand Heterogeneity. <i>Biophysical Journal</i> , 2018, 114, 1830-1846.	0.2	27
36	Nanocarrier Hydrodynamics and Binding in Targeted Drug Delivery: Challenges in Numerical Modeling and Experimental Validation. <i>Journal of Nanotechnology in Engineering and Medicine</i> , 2013, 4, 101011-1010115.	0.8	26

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37	Multiscale computational models in physical systems biology of intracellular trafficking. IET Systems Biology, 2014, 8, 198-213.	0.8	26
38	Biophysically inspired model for functionalized nanocarrier adhesion to cell surface: roles of protein expression and mechanical factors. Royal Society Open Science, 2016, 3, 160260.	1.1	26
39	Molecular systems biology of ErbB1 signaling: bridging the gap through multiscale modeling and high-performance computing. Molecular BioSystems, 2008, 4, 1151.	2.9	25
40	Composite generalized Langevin equation for Brownian motion in different hydrodynamic and adhesion regimes. Physical Review E, 2015, 91, 052303.	0.8	25
41	Role of Network Branching in Eliciting Differential Short-Term Signaling Responses in the Hypersensitive Epidermal Growth Factor Receptor Mutants Implicated in Lung Cancer. Biotechnology Progress, 2008, 24, 540-553.	1.3	22
42	Lateral distribution of phosphatidylinositol 4,5-bisphosphate in membranes regulates formin- and ARP2/3-mediated actin nucleation. Journal of Biological Chemistry, 2019, 294, 4704-4722.	1.6	22
43	Molecular Dynamics Simulations Reveal that Tyr-317 Phosphorylation Reduces Shc Binding Affinity for Phosphotyrosyl Residues of Epidermal Growth Factor Receptor. Biophysical Journal, 2009, 96, 2278-2288.	0.2	21
44	Analysis of Somatic Mutations in Cancer: Molecular Mechanisms of Activation in the ErbB Family of Receptor Tyrosine Kinases. Cancers, 2011, 3, 1195-1231.	1.7	21
45	Coupling of Fast and Slow Modes in the Reaction Pathway of the Minimal Hammerhead Ribozyme Cleavage. Biophysical Journal, 2007, 93, 2391-2399.	0.2	20
46	Defining the free-energy landscape of curvature-inducing proteins on membrane bilayers. Physical Review E, 2014, 90, 022717.	0.8	20
47	Calculation of free energies in fluid membranes subject to heterogeneous curvature fields. Physical Review E, 2009, 80, 011925.	0.8	17
48	High-throughput mutagenesis reveals functional determinants for DNA targeting by activation-induced deaminase. Nucleic Acids Research, 2014, 42, 9964-9975.	6.5	17
49	Divalent cations bind to phosphoinositides to induce ion and isomer specific propensities for nano-cluster initiation in bilayer membranes. Royal Society Open Science, 2020, 7, 192208.	1.1	17
50	Nanoparticle stochastic motion in the inertial regime and hydrodynamic interactions close to a cylindrical wall. Physical Review Fluids, 2016, 1, .	1.0	17
51	Curvature-Driven Migration of Colloids on Tense Lipid Bilayers. Langmuir, 2017, 33, 600-610.	1.6	16
52	Landscape of finite-temperature equilibrium behaviour of curvature-inducing proteins on a bilayer membrane explored using a linearized elastic free energy model. Molecular Physics, 2008, 106, 1913-1923.	0.8	15
53	Application of a free-energy-landscape approach to study tension-dependent bilayer tubulation mediated by curvature-inducing proteins. Physical Review E, 2015, 92, 042715.	0.8	15
54	Excess area dependent scaling behavior of nano-sized membrane tethers. Physical Biology, 2018, 15, 026002.	0.8	15

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55	Stiffness can mediate balance between hydrodynamic forces and avidity to impact the targeting of flexible polymeric nanoparticles in flow. <i>Nanoscale</i> , 2019, 11, 6916-6928.	2.8	15
56	Microstructure of Flow-Driven Suspension of Hardspheres in Cylindrical Confinement: A Dynamical Density Functional Theory and Monte Carlo Study. <i>Langmuir</i> , 2017, 33, 11332-11344.	1.6	14
57	A survey of multiscale modeling: Foundations, historical milestones, current status, and future prospects. <i>AIChE Journal</i> , 2021, 67, e17026.	1.8	14
58	"KMC-TDGL" a coarse-grained methodology for simulating interfacial dynamics in complex fluids: application to protein-mediated membrane processes. <i>Molecular Physics</i> , 2006, 104, 3653-3666.	0.8	13
59	Temporal multiscale approach for nanocarrier motion with simultaneous adhesion and hydrodynamic interactions in targeted drug delivery. <i>Journal of Computational Physics</i> , 2013, 244, 252-263.	1.9	13
60	Molecular modeling of ErbB4/HER4 kinase in the context of the HER4 signaling network helps rationalize the effects of clinically identified HER4 somatic mutations on the cell phenotype. <i>Biotechnology Journal</i> , 2013, 8, 1452-1464.	1.8	13
61	Hydrodynamic interactions of deformable polymeric nanocarriers and the effect of crosslinking. <i>Soft Matter</i> , 2015, 11, 5955-5969.	1.2	13
62	Physical chemistry and membrane properties of two phosphatidylinositol bisphosphate isomers. <i>Physical Chemistry Chemical Physics</i> , 2015, 17, 12608-12615.	1.3	12
63	Motion of a nano-spheroid in a cylindrical vessel flow: Brownian and hydrodynamic interactions. <i>Journal of Fluid Mechanics</i> , 2017, 821, 117-152.	1.4	12
64	Rheology of colloidal suspensions in confined flow: Treatment of hydrodynamic interactions in particle-based simulations inspired by dynamical density functional theory. <i>Physical Review E</i> , 2018, 98, .	0.8	12
65	Computational algorithms for in silico profiling of activating mutations in cancer. <i>Cellular and Molecular Life Sciences</i> , 2019, 76, 2663-2679.	2.4	11
66	A hybrid formalism combining fluctuating hydrodynamics and generalized Langevin dynamics for the simulation of nanoparticle thermal motion in an incompressible fluid medium. <i>Molecular Physics</i> , 2012, 110, 1057-1067.	0.8	10
67	Computational Models for Nanoscale Fluid Dynamics and Transport Inspired by Nonequilibrium Thermodynamics1. <i>Journal of Heat Transfer</i> , 2017, 139, 0330011-330019.	1.2	10
68	Emergent membrane morphologies in relaxed and tense membranes in presence of reversible adhesive pinning interactions. <i>Physical Biology</i> , 2019, 16, 066011.	0.8	10
69	Multiscale modeling of protein membrane interactions for nanoparticle targeting in drug delivery. <i>Current Opinion in Structural Biology</i> , 2020, 64, 104-110.	2.6	9
70	Structural Systems Biology and Multiscale Signaling Models. <i>Annals of Biomedical Engineering</i> , 2012, 40, 2295-2306.	1.3	8
71	Nanoparticle transport phenomena in confined flows. <i>Advances in Heat Transfer</i> , 2019, 51, 55-129.	0.4	8
72	Thermodynamic analysis of multivalent binding of functionalized nanoparticles to membrane surface reveals the importance of membrane entropy and nanoparticle entropy in adhesion of flexible nanoparticles. <i>Soft Matter</i> , 2019, 15, 9271-9286.	1.2	7

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73	A multiscale biophysical model for the recruitment of actin nucleating proteins at the membrane interface. <i>Soft Matter</i> , 2020, 16, 4941-4954.	1.2	7
74	Multiscale Cancer Modeling and In Silico Oncology: Emerging Computational Frontiers in Basic and Translational Cancer Research. <i>Journal of Bioengineering &amp; Biomedical Science</i> , 2013, 03, .	0.2	6
75	Crowding-induced membrane remodeling: Interplay of membrane tension, polymer density, architecture. <i>Biophysical Journal</i> , 2022, 121, 3674-3683.	0.2	6
76	Effect of wall-mediated hydrodynamic fluctuations on the kinetics of a Brownian nanoparticle. <i>Proceedings of the Royal Society A: Mathematical, Physical and Engineering Sciences</i> , 2016, 472, 20160397.	1.0	5
77	Deletion Mutations Keep Kinase Inhibitors in the Loop. <i>Cancer Cell</i> , 2016, 29, 423-425.	7.7	5
78	Thermodynamic free energy methods to investigate shape transitions in bilayer membranes. <i>International Journal of Advances in Engineering Sciences and Applied Mathematics</i> , 2016, 8, 88-100.	0.7	5
79	Time-dependent antagonist-agonist switching in receptor tyrosine kinase-mediated signaling. <i>BMC Bioinformatics</i> , 2019, 20, 242.	1.2	5
80	Nanofluid Dynamics of Flexible Polymeric Nanoparticles Under Wall Confinement. <i>Journal of Heat Transfer</i> , 2019, 141, 0524011-524016.	1.2	5
81	An interdomain helix in IRE1 $\beta$ mediates the conformational change required for the sensor's activation. <i>Journal of Biological Chemistry</i> , 2021, 296, 100781.	1.6	5
82	Phenomenology Based Multiscale Models as Tools to Understand Cell Membrane and Organelle Morphologies. <i>Behavior Research Methods</i> , 2015, 22, 129-175.	2.3	4
83	Probing lipid membrane bending mechanics using gold nanorod tracking. <i>Physical Review Research</i> , 2022, 4, .	1.3	4
84	Computational studies of anaplastic lymphoma kinase mutations reveal common mechanisms of oncogenic activation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, e2019132118.	3.3	3
85	Dimerization and structure formation of colloids via capillarity at curved fluid interfaces. <i>Soft Matter</i> , 2020, 16, 5861-5870.	1.2	2
86	Membrane signalosome: Where biophysics meets systems biology. <i>Current Opinion in Systems Biology</i> , 2021, 25, 34-41.	1.3	2
87	Biophysical Considerations in the Rational Design and Cellular Targeting of Flexible Polymeric Nanoparticles. <i>Advanced Materials Interfaces</i> , 2021, 8, 2101290.	1.9	2
88	Quantification of Curvature Sensing Behavior of Curvature-Inducing Proteins on Model Wavy Substrates. <i>Journal of Membrane Biology</i> , 2022, 255, 175-184.	1.0	2
89	Modeling of a Nanoparticle Motion in a Newtonian Fluid: A Comparison Between Fluctuating Hydrodynamics and Generalized Langevin Procedures. , 2012, 2012, 735-743.		1
90	Machine learning predictions of cancer driver mutations. , 2014, 2014, .		1

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91	Multiphysics pharmacokinetic model for targeted nanoparticles. <i>Frontiers in Medical Technology</i> , 0, 4, .	1.3	1
92	Data driven and biophysical insights into the regulation of trafficking vesicles by extracellular matrix stiffness. <i>IScience</i> , 2022, 25, 104721.	1.9	1
93	Structural Insights into Pseudokinase Domains of Receptor Tyrosine Kinases. <i>FASEB Journal</i> , 2021, 35, .	0.2	0
94	Biophysical Considerations in the Rational Design and Cellular Targeting of Flexible Polymeric Nanoparticles ( <i>Adv. Mater. Interfaces</i> 23/2021). <i>Advanced Materials Interfaces</i> , 2021, 8, .	1.9	0