

Bart W Hoogenboom

List of Publications by Year in descending order

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Version: 2024-02-01

70
papers

2,674
citations

147566

31
h-index

205818

48
g-index

87
all docs

87
docs citations

87
times ranked

3344
citing authors

#	ARTICLE	IF	CITATIONS
1	Crowding-induced phase separation of nuclear transport receptors in FG nucleoporin assemblies. <i>ELife</i> , 2022, 11, .	2.8	10
2	In-situ nanoscale imaging reveals self-concentrating nanomolar antimicrobial pores. <i>Nanoscale</i> , 2022, , .	2.8	0
3	Cooperative amyloid fibre binding and disassembly by the Hsp70 disaggregase. <i>EMBO Journal</i> , 2022, 41, .	3.5	14
4	Lipid specificity of the immune effector perforin. <i>Faraday Discussions</i> , 2021, 232, 236-255.	1.6	7
5	Atomic force microscopy to elucidate how peptides disrupt membranes. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2021, 1863, 183447.	1.4	36
6	AFM imaging of pore forming proteins. <i>Methods in Enzymology</i> , 2021, 649, 149-188.	0.4	3
7	Base-pair resolution analysis of the effect of supercoiling on DNA flexibility and major groove recognition by triplex-forming oligonucleotides. <i>Nature Communications</i> , 2021, 12, 1053.	5.8	73
8	Switching Cytolytic Nanopores into Antimicrobial Fractal Ruptures by a Single Side Chain Mutation. <i>ACS Nano</i> , 2021, 15, 9679-9689.	7.3	17
9	Physical modeling of multivalent interactions in the nuclear pore complex. <i>Biophysical Journal</i> , 2021, 120, 1565-1577.	0.2	14
10	Stretching the resolution limit of atomic force microscopy. <i>Nature Structural and Molecular Biology</i> , 2021, 28, 629-630.	3.6	8
11	Physics of the nuclear pore complex: Theory, modeling and experiment. <i>Physics Reports</i> , 2021, 921, 1-53.	10.3	44
12	Single-molecule measurements reveal that PARP1 condenses DNA by loop stabilization. <i>Science Advances</i> , 2021, 7, .	4.7	23
13	TopoStats â€” A program for automated tracing of biomolecules from AFM images. <i>Methods</i> , 2021, 193, 68-79.	1.9	23
14	Imaging the Effects of Peptide Materials on Phospholipid Membranes by Atomic Force Microscopy. <i>Methods in Molecular Biology</i> , 2021, 2208, 225-235.	0.4	3
15	Phase separation in the outer membrane of <i>Escherichia coli</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	53
16	Engineering Chirally Blind Protein Pseudocapsids into Antibacterial Persisters. <i>ACS Nano</i> , 2020, 14, 1609-1622.	7.3	42
17	Flowering Porationâ€™ A Synergistic Multi-Mode Antibacterial Mechanism by a Bacteriocin Fold. <i>IScience</i> , 2020, 23, 101423.	1.9	16
18	Structural basis for tuning activity and membrane specificity of bacterial cytolysins. <i>Nature Communications</i> , 2020, 11, 5818.	5.8	13

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19	Bacterial killing by complement requires direct anchoring of membrane attack complex precursor C5b-7. PLoS Pathogens, 2020, 16, e1008606.	2.1	28
20	Intrinsically disordered nuclear pore proteins show ideal-polymer morphologies and dynamics. Physical Review E, 2020, 101, 022420.	0.8	18
21	Acoustic Immunosensing of Exosomes Using a Quartz Crystal Microbalance with Dissipation Monitoring. Analytical Chemistry, 2020, 92, 4082-4093.	3.2	55
22	Membrane disrupting peptides: mechanistic elucidation of antimicrobial activity. Amino Acids, Peptides and Proteins, 2020, , 115-139.	0.7	0
23	Title is missing!. , 2020, 16, e1008606.		0
24	Title is missing!. , 2020, 16, e1008606.		0
25	Title is missing!. , 2020, 16, e1008606.		0
26	The cryo-EM structure of the acid activatable pore-forming immune effector Macrophage-expressed gene 1. Nature Communications, 2019, 10, 4288.	5.8	65
27	Quantification of Biomolecular Dynamics Inside Real and Synthetic Nuclear Pore Complexes Using Time-Resolved Atomic Force Microscopy. ACS Nano, 2019, 13, 7949-7956.	7.3	14
28	Helminth Defense Molecules as Design Templates for Membrane Active Antibiotics. ACS Infectious Diseases, 2019, 5, 1471-1479.	1.8	11
29	Single-molecule kinetics of pore assembly by the membrane attack complex. Nature Communications, 2019, 10, 2066.	5.8	74
30	Lipid order and charge protect killer T cells from accidental death. Nature Communications, 2019, 10, 5396.	5.8	56
31	Imaging live bacteria at the nanoscale: comparison of immobilisation strategies. Analyst, The, 2019, 144, 6944-6952.	1.7	21
32	PEGylated surfaces for the study of DNA-protein interactions by atomic force microscopy. Nanoscale, 2019, 11, 20072-20080.	2.8	15
33	Bacterial killing by complement requires membrane attack complex formation via surface-bound C5 convertases. EMBO Journal, 2019, 38, .	3.5	76
34	A Programmable DNA Origami Platform for Organizing Intrinsically Disordered Nucleoporins within Nanopore Confinement. ACS Nano, 2018, 12, 1508-1518.	7.3	84
35	The case for biophysics super-groups in physics departments. Physical Biology, 2018, 15, 060201.	0.8	2
36	Tuneable poration: host defense peptides as sequence probes for antimicrobial mechanisms. Scientific Reports, 2018, 8, 14926.	1.6	24

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37	Atomic force microscopy reveals structural variability amongst nuclear pore complexes. Life Science Alliance, 2018, 1, e201800142.	1.3	28
38	Real-time visualization of perforin nanopore assembly. Nature Nanotechnology, 2017, 12, 467-473.	15.6	88
39	Biomechanics of the transport barrier in the nuclear pore complex. Seminars in Cell and Developmental Biology, 2017, 68, 42-51.	2.3	37
40	Engineering monolayer poration for rapid exfoliation of microbial membranes. Chemical Science, 2017, 8, 1105-1115.	3.7	35
41	Antimicrobial peptide capsids of de novo design. Nature Communications, 2017, 8, 2263.	5.8	63
42	The membrane attack complex, perforin and cholesterol-dependent cytolysin superfamily of pore-forming proteins. Journal of Cell Science, 2016, 129, 2125-33.	1.2	45
43	Studies of G-quadruplexes formed within self-assembled DNA mini-circles. Chemical Communications, 2016, 52, 12454-12457.	2.2	15
44	Atomic force microscopy of membrane pore formation by cholesterol dependent cytolysins. Current Opinion in Structural Biology, 2016, 39, 8-15.	2.6	17
45	Imaging DNA Structure by Atomic Force Microscopy. Methods in Molecular Biology, 2016, 1431, 47-60.	0.4	14
46	Structurally plastic peptide capsules for synthetic antimicrobial viruses. Chemical Science, 2016, 7, 1707-1711.	3.7	43
47	A physical model describing the interaction of nuclear transport receptors with FG nucleoporin domain assemblies. ELife, 2016, 5, .	2.8	69
48	Reversible Dissolution of Microdomains in Detergent-Resistant Membranes at Physiological Temperature. PLoS ONE, 2015, 10, e0132696.	1.1	2
49	Nanoscale stiffness topography reveals structure and mechanics of the transport barrier in intact nuclear pore complexes. Nature Nanotechnology, 2015, 10, 60-64.	15.6	57
50	AFM in Liquids. , 2015, , 1-9.		0
51	Atomic force microscopy on plasma membranes from <i>Xenopus laevis</i> oocytes containing human aquaporin 4. Journal of Molecular Recognition, 2014, 27, 669-675.	1.1	1
52	Single-Molecule Reconstruction of Oligonucleotide Secondary Structure by Atomic Force Microscopy. Small, 2014, 10, 3257-3261.	5.2	96
53	Stepwise visualization of membrane pore formation by sulysin, a bacterial cholesterol-dependent cytolysin. ELife, 2014, 3, e04247.	2.8	145
54	Physical modelling of the nuclear pore complex. Soft Matter, 2013, 9, 10442.	1.2	28

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55	Model Inspired by Nuclear Pore Complex Suggests Possible Roles for Nuclear Transport Receptors in Determining Its Structure. <i>Biophysical Journal</i> , 2013, 105, 2781-2789.	0.2	29
56	Nanoscale imaging reveals laterally expanding antimicrobial pores in lipid bilayers. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 8918-8923.	3.3	112
57	Using Micromechanical Resonators to Measure Rheological Properties and Alcohol Content of Model Solutions and Commercial Beverages. <i>Sensors</i> , 2012, 12, 6497-6507.	2.1	13
58	Bistable collective behavior of polymers tethered in a nanopore. <i>Physical Review E</i> , 2012, 85, 061917.	0.8	35
59	Enhanced quality factors and force sensitivity by attaching magnetic beads to cantilevers for atomic force microscopy in liquid. <i>Journal of Applied Physics</i> , 2012, 112, .	1.1	12
60	Resolving the structure of a model hydrophobic surface: DODAB monolayers on mica. <i>RSC Advances</i> , 2012, 2, 4181.	1.7	10
61	Atomic Force Microscopy with Nanoscale Cantilevers Resolves Different Structural Conformations of the DNA Double Helix. <i>Nano Letters</i> , 2012, 12, 3846-3850.	4.5	83
62	Improved Kelvin probe force microscopy for imaging individual DNA molecules on insulating surfaces. <i>Applied Physics Letters</i> , 2010, 97, .	1.5	36
63	Imaging the Essential Role of Spin Fluctuations in High- T_c Superconductivity. <i>Physical Review Letters</i> , 2009, 103, 227001.	2.9	40
64	Imaging Surface Charges of Individual Biomolecules. <i>Nano Letters</i> , 2009, 9, 2769-2773.	4.5	85
65	The Supramolecular Assemblies of Voltage-dependent Anion Channels in the Native Membrane. <i>Journal of Molecular Biology</i> , 2007, 370, 246-255.	2.0	157
66	Quantitative dynamic-mode scanning force microscopy in liquid. <i>Applied Physics Letters</i> , 2006, 88, 193109.	1.5	88
67	Field Dependent Coherence Length in the Superclean, High- T_c Superconductor $CeCoIn_5$. <i>Physical Review Letters</i> , 2006, 97, 127001.	2.9	37
68	Hexagonal and Square Flux Line Lattices in $CeCoIn_5$. <i>Physical Review Letters</i> , 2003, 90, 187001.	2.9	53
69	Linear and Field-Independent Relation between Vortex Core State Energy and Gap in $Bi_2Sr_2CaCu_2O_{8+\delta}$. <i>Physical Review Letters</i> , 2001, 87, 267001.	2.9	42
70	Charge transfer and doping-dependent hybridization of C_{60} on noble metals. <i>Physical Review B</i> , 1998, 57, 11939-11942.	1.1	104