

Daniel J Muller

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279
papers

19,380
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78
h-index

128
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286
ext. papers

21,987
ext. citations

9.6
avg, IF

6.93
L-index

#	Paper	IF	Citations
279	Atomic force microscopy as a multifunctional molecular toolbox in nanobiotechnology. <i>Nature Nanotechnology</i> , 2008 , 3, 261-9	28.7	568
278	GSDMD membrane pore formation constitutes the mechanism of pyroptotic cell death. <i>EMBO Journal</i> , 2016 , 35, 1766-78	13	521
277	Imaging modes of atomic force microscopy for application in molecular and cell biology. <i>Nature Nanotechnology</i> , 2017 , 12, 295-307	28.7	494
276	Hydrostatic pressure and the actomyosin cortex drive mitotic cell rounding. <i>Nature</i> , 2011 , 469, 226-30	50.4	453
275	Structural biology. Proton-powered turbine of a plant motor. <i>Nature</i> , 2000 , 405, 418-9	50.4	439
274	Observing single biomolecules at work with the atomic force microscope. <i>Nature Structural Biology</i> , 2000 , 7, 715-8		434
273	Single-cell force spectroscopy. <i>Journal of Cell Science</i> , 2008 , 121, 1785-91	5.3	380
272	Force probing surfaces of living cells to molecular resolution. <i>Nature Chemical Biology</i> , 2009 , 5, 383-90	11.7	371
271	Imaging and manipulation of biological structures with the AFM. <i>Micron</i> , 2002 , 33, 385-97	2.3	324
270	Electrostatically balanced subnanometer imaging of biological specimens by atomic force microscope. <i>Biophysical Journal</i> , 1999 , 76, 1101-11	2.9	318
269	Multiparametric imaging of biological systems by force-distance curve-based AFM. <i>Nature Methods</i> , 2013 , 10, 847-54	21.6	317
268	Adsorption of biological molecules to a solid support for scanning probe microscopy. <i>Journal of Structural Biology</i> , 1997 , 119, 172-88	3.4	282
267	Atomic force microscopy: a nanoscopic window on the cell surface. <i>Trends in Cell Biology</i> , 2011 , 21, 461-9	18.3	279
266	Atomic force microscopy-based mechanobiology. <i>Nature Reviews Physics</i> , 2019 , 1, 41-57	23.6	274
265	Structure of the rhodopsin dimer: a working model for G-protein-coupled receptors. <i>Current Opinion in Structural Biology</i> , 2006 , 16, 252-9	8.1	229
264	Tapping-mode atomic force microscopy produces faithful high-resolution images of protein surfaces. <i>Biophysical Journal</i> , 1999 , 77, 1150-8	2.9	223
263	Conformational changes in surface structures of isolated connexin 26 gap junctions. <i>EMBO Journal</i> , 2002 , 21, 3598-607	13	209

262	Voltage and pH-induced channel closure of porin OmpF visualized by atomic force microscopy. <i>Journal of Molecular Biology</i> , 1999 , 285, 1347-51	6.5	205
261	Neuronal uptake and propagation of a rare phosphorylated high-molecular-weight tau derived from Alzheimer's disease brain. <i>Nature Communications</i> , 2015 , 6, 8490	17.4	204
260	A new technical approach to quantify cell-cell adhesion forces by AFM. <i>Ultramicroscopy</i> , 2006 , 106, 637-41	4.1	195
259	Assembly of collagen into microribbons: effects of pH and electrolytes. <i>Journal of Structural Biology</i> , 2004 , 148, 268-78	3.4	191
258	AFM: a nanotool in membrane biology. <i>Biochemistry</i> , 2008 , 47, 7986-98	3.2	186
257	Control of directed cell migration in vivo by membrane-to-cortex attachment. <i>PLoS Biology</i> , 2010 , 8, e1000544	10.4	185
256	Atomic force microscopy and spectroscopy of native membrane proteins. <i>Nature Protocols</i> , 2007 , 2, 2191-8	17.8	178
255	Force-induced conformational change of bacteriorhodopsin. <i>Journal of Molecular Biology</i> , 1995 , 249, 239-43	6.5	174
254	Bacterial Na(+)-ATP synthase has an undecameric rotor. <i>EMBO Reports</i> , 2001 , 2, 229-33	6.5	170
253	Revealing early steps of alpha2beta1 integrin-mediated adhesion to collagen type I by using single-cell force spectroscopy. <i>Molecular Biology of the Cell</i> , 2007 , 18, 1634-44	3.5	165
252	High resolution imaging of native biological sample surfaces using scanning probe microscopy. <i>Current Opinion in Structural Biology</i> , 1997 , 7, 279-84	8.1	156
251	Kindlin-2 cooperates with talin to activate integrins and induces cell spreading by directly binding paxillin. <i>ELife</i> , 2016 , 5, e10130	8.9	155
250	The c15 ring of the <i>Spirulina platensis</i> F-ATP synthase: F1/F0 symmetry mismatch is not obligatory. <i>EMBO Reports</i> , 2005 , 6, 1040-4	6.5	150
249	Stability of bacteriorhodopsin alpha-helices and loops analyzed by single-molecule force spectroscopy. <i>Biophysical Journal</i> , 2002 , 83, 3578-88	2.9	150
248	Straight GDP-tubulin protofilaments form in the presence of taxol. <i>Current Biology</i> , 2007 , 17, 1765-70	6.3	147
247	Measuring cell adhesion forces of primary gastrulating cells from zebrafish using atomic force microscopy. <i>Journal of Cell Science</i> , 2005 , 118, 4199-206	5.3	143
246	Surface tongue-and-groove contours on lens MIP facilitate cell-to-cell adherence. <i>Journal of Molecular Biology</i> , 2000 , 300, 779-89	6.5	139
245	Observing structure, function and assembly of single proteins by AFM. <i>Progress in Biophysics and Molecular Biology</i> , 2002 , 79, 1-43	4.7	138

244	Quantifying cellular adhesion to extracellular matrix components by single-cell force spectroscopy. <i>Nature Protocols</i> , 2010 , 5, 1353-61	18.8	137
243	Five challenges to bringing single-molecule force spectroscopy into living cells. <i>Nature Methods</i> , 2011 , 8, 123-7	21.6	136
242	From images to interactions: high-resolution phase imaging in tapping-mode atomic force microscopy. <i>Biophysical Journal</i> , 2001 , 80, 3009-18	2.9	132
241	Observing growth steps of collagen self-assembly by time-lapse high-resolution atomic force microscopy. <i>Journal of Structural Biology</i> , 2006 , 154, 232-45	3.4	128
240	A practical guide to quantify cell adhesion using single-cell force spectroscopy. <i>Methods</i> , 2013 , 60, 169-78	7.6	127
239	Nanomechanical mapping of first binding steps of a virus to animal cells. <i>Nature Nanotechnology</i> , 2017 , 12, 177-183	28.7	127
238	Surface structures of native bacteriorhodopsin depend on the molecular packing arrangement in the membrane. <i>Journal of Molecular Biology</i> , 1999 , 285, 1903-9	6.5	127
237	Cholesterol increases kinetic, energetic, and mechanical stability of the human β -adrenergic receptor. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, E3463-72	11.5	121
236	Mechanism of membrane pore formation by human gasdermin-D. <i>EMBO Journal</i> , 2018 , 37,	13	114
235	Deciphering molecular interactions of native membrane proteins by single-molecule force spectroscopy. <i>Annual Review of Biophysics and Biomolecular Structure</i> , 2007 , 36, 233-60		113
234	Analyzing focal adhesion structure by atomic force microscopy. <i>Journal of Cell Science</i> , 2005 , 118, 5315-23	23	110
233	The fuzzy coat of pathological human Tau fibrils is a two-layered polyelectrolyte brush. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, E313-21	11.5	109
232	Electrostatic cell-surface repulsion initiates lumen formation in developing blood vessels. <i>Current Biology</i> , 2010 , 20, 2003-9	6.3	108
231	Quantification of surface tension and internal pressure generated by single mitotic cells. <i>Scientific Reports</i> , 2014 , 4, 6213	4.9	105
230	Atomic force microscopy of native purple membrane. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2000 , 1460, 27-38	4.6	104
229	Hydrodynamic effects in fast AFM single-molecule force measurements. <i>European Biophysics Journal</i> , 2005 , 34, 91-6	1.9	103
228	Cdk1-dependent mitotic enrichment of cortical myosin II promotes cell rounding against confinement. <i>Nature Cell Biology</i> , 2015 , 17, 148-59	23.4	102
227	High-resolution atomic force microscopy and spectroscopy of native membrane proteins. <i>Reports on Progress in Physics</i> , 2011 , 74, 086601	14.4	102

226	Unfolding pathways of native bacteriorhodopsin depend on temperature. <i>EMBO Journal</i> , 2003 , 22, 5220-9	100
225	Oligomer formation of tau protein hyperphosphorylated in cells. <i>Journal of Biological Chemistry</i> , 2014 , 289, 34389-407	5.4 99
224	The effect of unlocking RGD-motifs in collagen I on pre-osteoblast adhesion and differentiation. <i>Biomaterials</i> , 2010 , 31, 2827-35	15.6 98
223	Mapping flexible protein domains at subnanometer resolution with the atomic force microscope. <i>FEBS Letters</i> , 1998 , 430, 105-11	3.8 97
222	Single-molecule studies of membrane proteins. <i>Current Opinion in Structural Biology</i> , 2006 , 16, 489-95	8.1 96
221	Atomic force microscopy-based characterization and design of biointerfaces. <i>Nature Reviews Materials</i> , 2017 , 2,	73.3 95
220	A glucose-starvation response regulates the diffusion of macromolecules. <i>ELife</i> , 2016 , 5,	8.9 93
219	The oligomeric state of c rings from cyanobacterial F-ATP synthases varies from 13 to 15. <i>Journal of Bacteriology</i> , 2007 , 189, 5895-902	3.5 92
218	Cellular remodelling of individual collagen fibrils visualized by time-lapse AFM. <i>Journal of Molecular Biology</i> , 2007 , 372, 594-607	6.5 92
217	Impact of holdase chaperones Skp and SurA on the folding of β barrel outer-membrane proteins. <i>Nature Structural and Molecular Biology</i> , 2015 , 22, 795-802	17.6 90
216	Controlled unfolding and refolding of a single sodium-proton antiporter using atomic force microscopy. <i>Journal of Molecular Biology</i> , 2004 , 340, 1143-52	6.5 90
215	Bacteriorhodopsin folds into the membrane against an external force. <i>Journal of Molecular Biology</i> , 2006 , 357, 644-54	6.5 89
214	Surface topographies at subnanometer-resolution reveal asymmetry and sidedness of aquaporin-1. <i>Journal of Molecular Biology</i> , 1996 , 264, 907-18	6.5 88
213	Characterizing molecular interactions in different bacteriorhodopsin assemblies by single-molecule force spectroscopy. <i>Journal of Molecular Biology</i> , 2006 , 355, 640-50	6.5 87
212	Movement directionality in collective migration of germ layer progenitors. <i>Current Biology</i> , 2010 , 20, 161-9	6.3 85
211	Imaging G protein-coupled receptors while quantifying their ligand-binding free-energy landscape. <i>Nature Methods</i> , 2015 , 12, 845-851	21.6 84
210	Multiparametric high-resolution imaging of native proteins by force-distance curve-based AFM. <i>Nature Protocols</i> , 2014 , 9, 1113-30	18.8 83
209	Human Tau isoforms assemble into ribbon-like fibrils that display polymorphic structure and stability. <i>Journal of Biological Chemistry</i> , 2010 , 285, 27302-27313	5.4 83

208	Vertebrate membrane proteins: structure, function, and insights from biophysical approaches. <i>Pharmacological Reviews</i> , 2008 , 60, 43-78	22.5	83
207	Folding and assembly of proteorhodopsin. <i>Journal of Molecular Biology</i> , 2008 , 376, 35-41	6.5	82
206	Fourteen protomers compose the oligomer III of the proton-rotor in spinach chloroplast ATP synthase. <i>Journal of Molecular Biology</i> , 2003 , 333, 337-44	6.5	82
205	Imaging and quantifying chemical and physical properties of native proteins at molecular resolution by force-volume AFM. <i>Angewandte Chemie - International Edition</i> , 2011 , 50, 12103-8	16.4	80
204	Atomic force microscopy: a forceful way with single molecules. <i>Current Biology</i> , 1999 , 9, R133-6	6.3	80
203	Ligand-specific interactions modulate kinetic, energetic, and mechanical properties of the human β adrenergic receptor. <i>Structure</i> , 2012 , 20, 1391-402	5.2	79
202	Imaging the electrostatic potential of transmembrane channels: atomic probe microscopy of OmpF porin. <i>Biophysical Journal</i> , 2002 , 82, 1667-76	2.9	79
201	Mechanism of allosteric regulation of β adrenergic receptor by cholesterol. <i>ELife</i> , 2016 , 5,	8.9	78
200	Locating ligand binding and activation of a single antiporter. <i>EMBO Reports</i> , 2005 , 6, 668-74	6.5	77
199	Rheology of the Active Cell Cortex in Mitosis. <i>Biophysical Journal</i> , 2016 , 111, 589-600	2.9	76
198	Structural changes in native membrane proteins monitored at subnanometer resolution with the atomic force microscope: a review. <i>Journal of Structural Biology</i> , 1997 , 119, 149-57	3.4	74
197	Probing the energy landscape of the membrane protein bacteriorhodopsin. <i>Structure</i> , 2004 , 12, 871-9	5.2	74
196	Fibronectin-bound β 1 integrins sense load and signal to reinforce adhesion in less than a second. <i>Nature Materials</i> , 2017 , 16, 1262-1270	27	72
195	Galectin-3 regulates integrin α 2 β 1-mediated adhesion to collagen-I and -IV. <i>Journal of Biological Chemistry</i> , 2008 , 283, 32264-72	5.4	72
194	Contributions of galectin-3 and -9 to epithelial cell adhesion analyzed by single cell force spectroscopy. <i>Journal of Biological Chemistry</i> , 2007 , 282, 29375-83	5.4	72
193	Engineering rotor ring stoichiometries in the ATP synthase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, E1599-608	11.5	70
192	The central plug in the reconstituted undecameric c cylinder of a bacterial ATP synthase consists of phospholipids. <i>FEBS Letters</i> , 2001 , 505, 353-6	3.8	69
191	α -class integrins exert dual roles on β 1 integrins to strengthen adhesion to fibronectin. <i>Nature Communications</i> , 2017 , 8, 14348	17.4	68

190	Observing membrane protein diffusion at subnanometer resolution. <i>Journal of Molecular Biology</i> , 2003 , 327, 925-30	6.5	67
189	A bond for a lifetime: employing membrane nanotubes from living cells to determine receptor-ligand kinetics. <i>Angewandte Chemie - International Edition</i> , 2008 , 47, 9775-7	16.4	64
188	Aminosulfonate modulated pH-induced conformational changes in connexin26 hemichannels. <i>Journal of Biological Chemistry</i> , 2007 , 282, 8895-904	5.4	64
187	Single-cell force spectroscopy, an emerging tool to quantify cell adhesion to biomaterials. <i>Tissue Engineering - Part B: Reviews</i> , 2014 , 20, 40-55	7.9	63
186	Force nanoscopy of living cells. <i>Current Biology</i> , 2011 , 21, R212-6	6.3	63
185	Studying integrin-mediated cell adhesion at the single-molecule level using AFM force spectroscopy. <i>Science & STKE: Signal Transduction Knowledge Environment</i> , 2007 , 2007, p15		63
184	Sampling the conformational space of membrane protein surfaces with the AFM. <i>European Biophysics Journal</i> , 2002 , 31, 172-8	1.9	63
183	ATP synthase: constrained stoichiometry of the transmembrane rotor. <i>FEBS Letters</i> , 2001 , 504, 219-22	3.8	63
182	Inertial picobalance reveals fast mass fluctuations in mammalian cells. <i>Nature</i> , 2017 , 550, 500-505	50.4	62
181	Surface analysis of the photosystem I complex by electron and atomic force microscopy. <i>Journal of Molecular Biology</i> , 1998 , 283, 83-94	6.5	62
180	New frontiers in atomic force microscopy: analyzing interactions from single-molecules to cells. <i>Current Opinion in Biotechnology</i> , 2009 , 20, 4-13	11.4	61
179	Detecting molecular interactions that stabilize native bovine rhodopsin. <i>Journal of Molecular Biology</i> , 2006 , 358, 255-69	6.5	61
178	Identification and structure of a putative Ca ²⁺ -binding domain at the C terminus of AQP1. <i>Journal of Molecular Biology</i> , 2002 , 318, 1381-94	6.5	61
177	Single Proteins Observed by Atomic Force Microscopy. <i>Single Molecules</i> , 2001 , 2, 59-67		60
176	Directly Observing the Lipid-Dependent Self-Assembly and Pore-Forming Mechanism of the Cytolytic Toxin Listeriolysin O. <i>Nano Letters</i> , 2015 , 15, 6965-73	11.5	59
175	The c13 ring from a thermoalkaliphilic ATP synthase reveals an extended diameter due to a special structural region. <i>Journal of Molecular Biology</i> , 2009 , 388, 611-8	6.5	59
174	Creating ultrathin nanoscopic collagen matrices for biological and biotechnological applications. <i>Small</i> , 2007 , 3, 956-63	11	59
173	Molecular force modulation spectroscopy revealing the dynamic response of single bacteriorhodopsins. <i>Biophysical Journal</i> , 2005 , 88, 1423-31	2.9	57

172	Stages and conformations of the Tau repeat domain during aggregation and its effect on neuronal toxicity. <i>Journal of Biological Chemistry</i> , 2014 , 289, 20318-32	5.4	56
171	Determining molecular forces that stabilize human aquaporin-1. <i>Journal of Structural Biology</i> , 2003 , 142, 369-78	3.4	56
170	SAS-6 engineering reveals interdependence between cartwheel and microtubules in determining centriole architecture. <i>Nature Cell Biology</i> , 2016 , 18, 393-403	23.4	55
169	Charting the surfaces of the purple membrane. <i>Journal of Structural Biology</i> , 1999 , 128, 243-9	3.4	55
168	Mechanical Stimulation of Piezo1 Receptors Depends on Extracellular Matrix Proteins and Directionality of Force. <i>Nano Letters</i> , 2017 , 17, 2064-2072	11.5	54
167	Mitotic cells contract actomyosin cortex and generate pressure to round against or escape epithelial confinement. <i>Nature Communications</i> , 2015 , 6, 8872	17.4	54
166	Mechanical control of mitotic progression in single animal cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015 , 112, 11258-63	11.5	53
165	Identifying and quantifying two ligand-binding sites while imaging native human membrane receptors by AFM. <i>Nature Communications</i> , 2015 , 6, 8857	17.4	53
164	Stabilizing effect of Zn ²⁺ in native bovine rhodopsin. <i>Journal of Biological Chemistry</i> , 2007 , 282, 11377-85	5.4	53
163	Deciphering teneurin domains that facilitate cellular recognition, cell-cell adhesion, and neurite outgrowth using atomic force microscopy-based single-cell force spectroscopy. <i>Nano Letters</i> , 2013 , 13, 2937-46	11.5	52
162	Biomolecular imaging using atomic force microscopy. <i>Trends in Biotechnology</i> , 2002 , 20, S45-S49	15.1	52
161	YidC assists the stepwise and stochastic folding of membrane proteins. <i>Nature Chemical Biology</i> , 2016 , 12, 911-917	11.7	52
160	Wedge AFM-cantilevers for parallel plate cell mechanics. <i>Methods</i> , 2013 , 60, 186-94	4.6	51
159	Multiparametric Atomic Force Microscopy Imaging of Biomolecular and Cellular Systems. <i>Accounts of Chemical Research</i> , 2017 , 50, 924-931	24.3	50
158	Conformational adaptability of Redbeta during DNA annealing and implications for its structural relationship with Rad52. <i>Journal of Molecular Biology</i> , 2009 , 391, 586-98	6.5	50
157	Probing origins of molecular interactions stabilizing the membrane proteins halorhodopsin and bacteriorhodopsin. <i>Structure</i> , 2005 , 13, 235-42	5.2	50
156	Membrane perforation by the pore-forming toxin pneumolysin. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019 , 116, 13352-13357	11.5	49
155	Point mutations in membrane proteins reshape energy landscape and populate different unfolding pathways. <i>Journal of Molecular Biology</i> , 2008 , 376, 1076-90	6.5	48

154	An intermediate step in the evolution of ATPases: a hybrid F(0)-V(0) rotor in a bacterial Na(+) F(1)F(0) ATP synthase. <i>FEBS Journal</i> , 2008 , 275, 1999-2007	5.7	48
153	Transmembrane helices have rough energy surfaces. <i>Journal of the American Chemical Society</i> , 2007 , 129, 246-7	16.4	47
152	Action of the Hsp70 chaperone system observed with single proteins. <i>Nature Communications</i> , 2015 , 6, 6307	17.4	46
151	pH-induced conformational change of the beta-barrel-forming protein OmpG reconstituted into native E. coli lipids. <i>Journal of Molecular Biology</i> , 2010 , 396, 610-6	6.5	46
150	Observing folding pathways and kinetics of a single sodium-proton antiporter from Escherichia coli. <i>Journal of Molecular Biology</i> , 2006 , 355, 2-8	6.5	46
149	Nanomechanical properties of proteins and membranes depend on loading rate and electrostatic interactions. <i>ACS Nano</i> , 2013 , 7, 2642-50	16.7	45
148	Gating of the MlotiK1 potassium channel involves large rearrangements of the cyclic nucleotide-binding domains. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 20802-7	11.5	45
147	Single-molecule force spectroscopy from nanodiscs: an assay to quantify folding, stability, and interactions of native membrane proteins. <i>ACS Nano</i> , 2012 , 6, 961-71	16.7	44
146	Strategies to prepare and characterize native membrane proteins and protein membranes by AFM. <i>Current Opinion in Colloid and Interface Science</i> , 2008 , 13, 338-350	7.6	44
145	The fibronectin synergy site re-enforces cell adhesion and mediates a crosstalk between integrin classes. <i>ELife</i> , 2017 , 6,	8.9	42
144	Preparation techniques for the observation of native biological systems with the atomic force microscope. <i>Biosensors and Bioelectronics</i> , 1997 , 12, 867-877	11.8	41
143	Free energy of membrane protein unfolding derived from single-molecule force measurements. <i>Biophysical Journal</i> , 2007 , 93, 930-7	2.9	41
142	Out but not in: the large transmembrane β barrel protein FhuA unfolds but cannot refold via β hairpins. <i>Structure</i> , 2012 , 20, 2185-90	5.2	40
141	Tracking mechanics and volume of globular cells with atomic force microscopy using a constant-height clamp. <i>Nature Protocols</i> , 2012 , 7, 143-54	18.8	40
140	Differentiating ligand and inhibitor interactions of a single antiporter. <i>Journal of Molecular Biology</i> , 2006 , 362, 925-32	6.5	40
139	Structural evidence for a constant c11 ring stoichiometry in the sodium F-ATP synthase. <i>FEBS Journal</i> , 2005 , 272, 5474-83	5.7	40
138	Isolation and characterization of gap junctions from tissue culture cells. <i>Journal of Molecular Biology</i> , 2002 , 315, 587-600	6.5	40
137	Combining confocal and atomic force microscopy to quantify single-virus binding to mammalian cell surfaces. <i>Nature Protocols</i> , 2017 , 12, 2275-2292	18.8	39

136	Localizing chemical groups while imaging single native proteins by high-resolution atomic force microscopy. <i>Nano Letters</i> , 2014 , 14, 2957-64	11.5	39
135	Mechanical properties of bovine rhodopsin and bacteriorhodopsin: possible roles in folding and function. <i>Langmuir</i> , 2008 , 24, 1330-7	4	38
134	Imaging and detecting molecular interactions of single transmembrane proteins. <i>Neurobiology of Aging</i> , 2006 , 27, 546-61	5.6	38
133	Kinetic, energetic, and mechanical differences between dark-state rhodopsin and opsin. <i>Structure</i> , 2013 , 21, 426-37	5.2	37
132	From valleys to ridges: exploring the dynamic energy landscape of single membrane proteins. <i>ChemPhysChem</i> , 2008 , 9, 954-66	3.2	37
131	One beta hairpin after the other: exploring mechanical unfolding pathways of the transmembrane beta-barrel protein OmpG. <i>Angewandte Chemie - International Edition</i> , 2009 , 48, 8306-8	16.4	36
130	BCR/ABL expression of myeloid progenitors increases beta1-integrin mediated adhesion to stromal cells. <i>Journal of Molecular Biology</i> , 2008 , 377, 1082-93	6.5	36
129	Creating nanoscopic collagen matrices using atomic force microscopy. <i>Microscopy Research and Technique</i> , 2004 , 64, 435-40	2.8	36
128	Reversible loss of crystallinity on photobleaching purple membrane in the presence of hydroxylamine. <i>Journal of Molecular Biology</i> , 2000 , 301, 869-79	6.5	36
127	Products of the Parkinson's disease-related glyoxalase DJ-1, D-lactate and glycolate, support mitochondrial membrane potential and neuronal survival. <i>Biology Open</i> , 2014 , 3, 777-84	2.2	35
126	Competing interactions stabilize pro- and anti-aggregant conformations of human Tau. <i>Journal of Biological Chemistry</i> , 2011 , 286, 20512-24	5.4	35
125	pH-dependent interactions guide the folding and gate the transmembrane pore of the beta-barrel membrane protein OmpG. <i>Journal of Molecular Biology</i> , 2010 , 397, 878-82	6.5	34
124	Protein-enriched outer membrane vesicles as a native platform for outer membrane protein studies. <i>Communications Biology</i> , 2018 , 1, 23	6.7	33
123	The transmembrane protein KpOmpA anchoring the outer membrane of <i>Klebsiella pneumoniae</i> unfolds and refolds in response to tensile load. <i>Structure</i> , 2012 , 20, 121-7	5.2	33
122	Dynamic coupling of ALCAM to the actin cortex strengthens cell adhesion to CD6. <i>Journal of Cell Science</i> , 2014 , 127, 1595-606	5.3	32
121	Probing the interactions of carboxy-atractyloside and atractyloside with the yeast mitochondrial ADP/ATP carrier. <i>Structure</i> , 2010 , 18, 39-46	5.2	32
120	Stimulated single-cell force spectroscopy to quantify cell adhesion receptor crosstalk. <i>Proteomics</i> , 2010 , 10, 1455-62	4.8	32
119	Actin microridges characterized by laser scanning confocal and atomic force microscopy. <i>FEBS Letters</i> , 2005 , 579, 2001-8	3.8	32

118	Virus stamping for targeted single-cell infection in vitro and in vivo. <i>Nature Biotechnology</i> , 2018 , 36, 81-88	14.5	31
117	Quantitative imaging of the electrostatic field and potential generated by a transmembrane protein pore at subnanometer resolution. <i>Nano Letters</i> , 2013 , 13, 5585-93	11.5	31
116	Scanning probe microscopy. <i>Nature Reviews Methods Primers</i> , 2021 , 1,		31
115	One hairpin follows the other: exploring refolding pathways and kinetics of the transmembrane β -barrel protein OmpG. <i>Angewandte Chemie - International Edition</i> , 2011 , 50, 7422-4	16.4	30
114	Electron and atomic force microscopy of membrane proteins. <i>Current Opinion in Structural Biology</i> , 1997 , 7, 543-9	8.1	30
113	Substrate binding tunes conformational flexibility and kinetic stability of an amino acid antiporter. <i>Journal of Biological Chemistry</i> , 2009 , 284, 18651-63	5.4	29
112	Modulation of molecular interactions and function by rhodopsin palmitylation. <i>Biochemistry</i> , 2009 , 48, 4294-304	3.2	29
111	Genome-scale single-cell mechanical phenotyping reveals disease-related genes involved in mitotic rounding. <i>Nature Communications</i> , 2017 , 8, 1266	17.4	28
110	Substrate-induced changes in the structural properties of LacY. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, E1571-80	11.5	28
109	Transducer binding establishes localized interactions to tune sensory rhodopsin II. <i>Structure</i> , 2008 , 16, 1206-13	5.2	28
108	Examining the dynamic energy landscape of an antiporter upon inhibitor binding. <i>Journal of Molecular Biology</i> , 2008 , 375, 1258-66	6.5	28
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