

Kalina Hristova

List of Publications by Year in descending order

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147
papers

7,339
citations

57631

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161
all docs

161
docs citations

161
times ranked

6832
citing authors

#	ARTICLE	IF	CITATIONS
1	Mechanical disruption of E-cadherin complexes with epidermal growth factor receptor actuates growth factor-dependent signaling. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	3.3	23
2	Ligands with different dimeric configurations potently activate the EphA2 receptor and reveal its potential for biased signaling. Science, 2022, 25, 103870.	1.9	8
3	Direct Quantification of Ligand-Induced Lipid and Protein Microdomains with Distinctive Signaling Properties**. ChemSystemsChem, 2022, 4, .	1.1	1
4	Effect of osmotic stress on live cell plasma membranes, probed via Laurdan general polarization measurements. Biophysical Journal, 2022, 121, 2411-2418.	0.2	0
5	The Biased Ligands NGF and NT-3 Differentially Stabilize Trk-A Dimers. Biophysical Journal, 2021, 120, 55-63.	0.2	16
6	Methods Ligand Binding to Receptor Tyrosine Kinases: Thermodynamic Cycles and Experimental Approaches. , 2021, , 766-779.		0
7	Probing Membrane Protein Association Using Concentration-Dependent Number and Brightness. Angewandte Chemie, 2021, 133, 6577-6582.	1.6	2
8	pH-triggered pore-forming peptides with strong composition-dependent membrane selectivity. Biophysical Journal, 2021, 120, 618-630.	0.2	11
9	Probing Membrane Protein Association Using Concentration-Dependent Number and Brightness. Angewandte Chemie - International Edition, 2021, 60, 6503-6508.	7.2	11
10	Interactions between Ligand-Bound EGFR and VEGFR2. Journal of Molecular Biology, 2021, 433, 167006.	2.0	3
11	Neural network strategies for plasma membrane selection in fluorescence microscopy images. Biophysical Journal, 2021, 120, 2374-2385.	0.2	1
12	How Can We Fully Realize the Potential of Mathematical and Biological Models to Reintegrate Biology?. Integrative and Comparative Biology, 2021, , .	0.9	1
13	A cancer mutation promotes EphA4 oligomerization and signaling by altering the conformation of the SAM domain. Journal of Biological Chemistry, 2021, 297, 100876.	1.6	9
14	P120 catenin potentiates constitutive E-cadherin dimerization at the plasma membrane and regulates trans binding. Current Biology, 2021, 31, 3017-3027.e7.	1.8	22
15	Reversible blood-brain barrier opening utilizing the membrane active peptide melittin in vitro and in vivo. Biomaterials, 2021, 275, 120942.	5.7	24
16	Interaction between the transmembrane domains of neurotrophin receptors p75 and TrkA mediates their reciprocal activation. Journal of Biological Chemistry, 2021, 297, 100926.	1.6	8
17	Human herpesvirus 8 molecular mimicry of ephrin ligands facilitates cell entry and triggers EphA2 signaling. PLoS Biology, 2021, 19, e3001392.	2.6	7
18	Quantitative characterization of tetraspanin 8 homointeractions in the plasma membrane. Biochemical Journal, 2021, 478, 3643-3654.	1.7	1

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19	Applications and evolution of melittin, the quintessential membrane active peptide. <i>Biochemical Pharmacology</i> , 2021, 193, 114769.	2.0	45
20	Pondering the mechanism of receptor tyrosine kinase activation: The case for ligand-specific dimer microstate ensembles. <i>Current Opinion in Structural Biology</i> , 2021, 71, 193-199.	2.6	11
21	Membrane-selective Nanoscale Pores in Liposomes by a Synthetically Evolved Peptide: Implications for Triggered Release. <i>Nanoscale</i> , 2021, 13, 12185-12197.	2.8	11
22	Regulation of the EphA2 receptor intracellular region by phosphomimetic negative charges in the kinase-SAM linker. <i>Nature Communications</i> , 2021, 12, 7047.	5.8	11
23	Receptor Tyrosine Kinases. , 2021, , .		1
24	Revisiting a controversy: The effect of EGF on EGFR dimer stability. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2020, 1862, 183015.	1.4	14
25	EGFR forms ligand-independent oligomers that are distinct from the active state. <i>Journal of Biological Chemistry</i> , 2020, 295, 13353-13362.	1.6	28
26	Ligand bias in receptor tyrosine kinase signaling. <i>Journal of Biological Chemistry</i> , 2020, 295, 18494-18507.	1.6	28
27	Rational Modulation of pH-Triggered Macromolecular Poration by Peptide Acylation and Dimerization. <i>Journal of Physical Chemistry B</i> , 2020, 124, 8835-8843.	1.2	3
28	A peptide for transcellular cargo delivery: Structure-function relationship and mechanism of action. <i>Journal of Controlled Release</i> , 2020, 324, 633-643.	4.8	14
29	Quantifying the strength of heterointeractions among receptor tyrosine kinases from different subfamilies: Implications for cell signaling. <i>Journal of Biological Chemistry</i> , 2020, 295, 9917-9933.	1.6	23
30	Enhancing the membrane activity of Piscidin 1 through peptide metallation and the presence of oxidized lipid species: Implications for the unification of host defense mechanisms at lipid membranes. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2020, 1862, 183236.	1.4	11
31	The Mechanism of Membrane Permeabilization by Peptides: Still an Enigma. <i>Australian Journal of Chemistry</i> , 2020, 73, 96.	0.5	34
32	The biophysical basis of receptor tyrosine kinase ligand functional selectivity: Trk-B case study. <i>Biochemical Journal</i> , 2020, 477, 4515-4526.	1.7	11
33	Engineering nanomolar peptide ligands that differentially modulate EphA2 receptor signaling. <i>Journal of Biological Chemistry</i> , 2019, 294, 8791-8805.	1.6	31
34	Direct measurements of VEGFâ€™VEGFR2 binding affinities reveal the coupling between ligand binding and receptor dimerization. <i>Journal of Biological Chemistry</i> , 2019, 294, 9064-9075.	1.6	28
35	Mechanism of Action of Peptides That Cause the pH-Triggered Macromolecular Poration of Lipid Bilayers. <i>Journal of the American Chemical Society</i> , 2019, 141, 6706-6718.	6.6	30
36	The transition model of RTK activation: A quantitative framework for understanding RTK signaling and RTK modulator activity. <i>Cytokine and Growth Factor Reviews</i> , 2019, 49, 23-31.	3.2	31

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37	The RTK Interactome: Overview and Perspective on RTK Heterointeractions. <i>Chemical Reviews</i> , 2019, 119, 5881-5921.	23.0	59
38	The EphA2 receptor is activated through induction of distinct, ligand-dependent oligomeric structures. <i>Communications Biology</i> , 2018, 1, 15.	2.0	62
39	Potent Macromolecule-Sized Poration of Lipid Bilayers by the Macrolittins, A Synthetically Evolved Family of Pore-Forming Peptides. <i>Journal of the American Chemical Society</i> , 2018, 140, 6441-6447.	6.6	41
40	Interactions between NRP1 and VEGFR2 molecules in the plasma membrane. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2018, 1860, 2118-2125.	1.4	23
41	Dimerization of the Trk receptors in the plasma membrane: effects of their cognate ligands. <i>Biochemical Journal</i> , 2018, 475, 3669-3685.	1.7	28
42	Single Proteoliposome High-Content Analysis Reveals Differences in the Homo-Oligomerization of GPCRs. <i>Biophysical Journal</i> , 2018, 115, 300-312.	0.2	19
43	Intracellular Domain Contacts Contribute to Ecadherin Constitutive Dimerization in the Plasma Membrane. <i>Journal of Molecular Biology</i> , 2017, 429, 2231-2245.	2.0	28
44	Interactions between membrane receptors in cellular membranes. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2017, 1859, 1397.	1.4	0
45	Ebola Virus Delta Peptide Is a Viroporin. <i>Journal of Virology</i> , 2017, 91, .	1.5	26
46	Understanding the FRET Signatures of Interacting Membrane Proteins. <i>Journal of Biological Chemistry</i> , 2017, 292, 5291-5310.	1.6	62
47	A New Method to Study Heterodimerization of Membrane Proteins and Its Application to Fibroblast Growth Factor Receptors. <i>Journal of Biological Chemistry</i> , 2017, 292, 1288-1301.	1.6	30
48	pH-Triggered, Macromolecule-Sized Poration of Lipid Bilayers by Synthetically Evolved Peptides. <i>Journal of the American Chemical Society</i> , 2017, 139, 937-945.	6.6	61
49	Cooperative interactions between VEGFR2 extracellular Ig-like subdomains ensure VEGFR2 dimerization. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2017, 1861, 2559-2567.	1.1	7
50	Quantifying the Interaction between EGFR Dimers and Grb2 in Live Cells. <i>Biophysical Journal</i> , 2017, 113, 1353-1364.	0.2	23
51	The SAM domain inhibits EphA2 interactions in the plasma membrane. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2017, 1864, 31-38.	1.9	43
52	VEGFR-2 conformational switch in response to ligand binding. <i>ELife</i> , 2016, 5, e13876.	2.8	94
53	Effect of the achondroplasia mutation on FGFR3 dimerization and FGFR3 structural response to fgf1 and fgf2: A quantitative FRET study in osmotically derived plasma membrane vesicles. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2016, 1858, 1436-1442.	1.4	15
54	Pathogenic Cysteine Removal Mutations in FGFR Extracellular Domains Stabilize Receptor Dimers and Perturb the TM Dimer Structure. <i>Journal of Molecular Biology</i> , 2016, 428, 3903-3910.	2.0	12

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55	Mechanism of FGF receptor dimerization and activation. <i>Nature Communications</i> , 2016, 7, 10262.	5.8	192
56	A small peptide promotes EphA2 kinase-dependent signaling by stabilizing EphA2 dimers. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2016, 1860, 1922-1928.	1.1	28
57	Fully quantified spectral imaging reveals <i>in vivo</i> membrane protein interactions. <i>Integrative Biology (United Kingdom)</i> , 2016, 8, 216-229.	0.6	82
58	Unliganded EphA3 dimerization promoted by the SAM domain. <i>Biochemical Journal</i> , 2015, 471, 101-109.	1.7	45
59	EphA2 Receptor Unliganded Dimers Suppress EphA2 Pro-tumorigenic Signaling. <i>Journal of Biological Chemistry</i> , 2015, 290, 27271-27279.	1.6	58
60	Effect of Thanatophoric Dysplasia Type I Mutations on FGFR3 Dimerization. <i>Biophysical Journal</i> , 2015, 108, 272-278.	0.2	29
61	Testing the limits of rational design by engineering pH sensitivity into membrane-active peptides. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2015, 1848, 951-957.	1.4	27
62	Characterization of Membrane Protein Interactions in Plasma Membrane Derived Vesicles with Quantitative Imaging FRET Resonance Energy Transfer. <i>Accounts of Chemical Research</i> , 2015, 48, 2262-2269.	7.6	45
63	Analytical characterization of plasma membrane-derived vesicles produced via osmotic and chemical vesiculation. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2015, 1848, 1591-1598.	1.4	22
64	Of Rafts and Lipid Chain Lengths. <i>Biophysical Journal</i> , 2015, 108, 2096.	0.2	0
65	FGFR3 Unliganded Dimer Stabilization by the Juxtamembrane Domain. <i>Journal of Molecular Biology</i> , 2015, 427, 1705-1714.	2.0	35
66	Quantification of the Effects of Mutations on Receptor Tyrosine Kinase (RTK) Activation in Mammalian Cells. <i>Methods in Molecular Biology</i> , 2015, 1233, 81-87.	0.4	1
67	Strong dimerization of wild-type ErbB2/Neu transmembrane domain and the oncogenic Val664Glu mutant in mammalian plasma membranes. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2014, 1838, 2326-2330.	1.4	15
68	The FRET Signatures of Noninteracting Proteins in Membranes: Simulations and Experiments. <i>Biophysical Journal</i> , 2014, 106, 1309-1317.	0.2	80
69	Highly Efficient Macromolecule-Sized Poration of Lipid Bilayers by a Synthetically Evolved Peptide. <i>Journal of the American Chemical Society</i> , 2014, 136, 4724-4731.	6.6	59
70	Interfacially active peptides and proteins. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2014, 1838, 2139.	1.4	1
71	Uninduced high-yield bacterial expression of fluorescent proteins. <i>Analytical Biochemistry</i> , 2014, 449, 155-157.	1.1	31
72	How IGF-1 activates its receptor. <i>ELife</i> , 2014, 3, .	2.8	154

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73	FGFR3 Transmembrane Domain Interactions Persist in the Presence of Its Extracellular Domain. <i>Biophysical Journal</i> , 2013, 105, 165-171.	0.2	15
74	Structure of FGFR3 Transmembrane Domain Dimer: Implications for Signaling and Human Pathologies. <i>Structure</i> , 2013, 21, 2087-2093.	1.6	69
75	The electrical response of bilayers to the bee venom toxin melittin: Evidence for transient bilayer permeabilization. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2013, 1828, 1357-1364.	1.4	50
76	A Membrane-Translocating Peptide Penetrates into Bilayers without Significant Bilayer Perturbations. <i>Biophysical Journal</i> , 2013, 104, 2419-2428.	0.2	42
77	Glycophorin A transmembrane domain dimerization in plasma membrane vesicles derived from CHO, HEK 293T, and A431 cells. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2013, 1828, 1829-1833.	1.4	31
78	Direct Cytosolic Delivery of Polar Cargo to Cells by Spontaneous Membrane-translocating Peptides. <i>Journal of Biological Chemistry</i> , 2013, 288, 29974-29986.	1.6	52
79	Multiple Consequences of a Single Amino Acid Pathogenic RTK Mutation: The A391E Mutation in FGFR3. <i>PLoS ONE</i> , 2013, 8, e56521.	1.1	11
80	Consequences of replacing EGFR juxtamembrane domain with an unstructured sequence. <i>Scientific Reports</i> , 2012, 2, 854.	1.6	28
81	Interactions of Membrane Active Peptides with Planar Supported Bilayers: An Impedance Spectroscopy Study. <i>Langmuir</i> , 2012, 28, 6088-6096.	1.6	24
82	Production of Plasma Membrane Vesicles with Chloride Salts and Their Utility as a Cell Membrane Mimetic for Biophysical Characterization of Membrane Protein Interactions. <i>Analytical Chemistry</i> , 2012, 84, 8650-8655.	3.2	68
83	Physical-chemical principles underlying RTK activation, and their implications for human disease. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2012, 1818, 995-1005.	1.4	49
84	Transmembrane helix dimerization: Beyond the search for sequence motifs. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2012, 1818, 183-193.	1.4	143
85	Effect of the G375C and G346E Achondroplasia Mutations on FGFR3 Activation. <i>PLoS ONE</i> , 2012, 7, e34808.	1.1	10
86	A Highly Charged Voltage-Sensor Helix Spontaneously Translocates across Membranes. <i>Angewandte Chemie - International Edition</i> , 2012, 51, 7150-7153.	7.2	28
87	On-the-resin N-terminal modification of long synthetic peptides. <i>Analytical Biochemistry</i> , 2012, 424, 137-139.	1.1	18
88	Direct Assessment of the Effect of the Gly380Arg Achondroplasia Mutation on FGFR3 Dimerization Using Quantitative Imaging FRET. <i>PLoS ONE</i> , 2012, 7, e46678.	1.1	45
89	The Physical Basis of FGFR3 Response to <i>fgf1</i> and <i>fgf2</i> . <i>Biochemistry</i> , 2011, 50, 8576-8582.	1.2	24
90	Spontaneous Membrane-Translocating Peptides by Orthogonal High-Throughput Screening. <i>Journal of the American Chemical Society</i> , 2011, 133, 8995-9004.	6.6	173

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91	Specific inhibition of a pathogenic receptor tyrosine kinase by its transmembrane domain. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2011, 1808, 253-259.	1.4	21
92	The A391E mutation enhances FGFR3 activation in the absence of ligand. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2011, 1808, 2045-2050.	1.4	29
93	High-Throughput Selection of Transmembrane Sequences That Enhance Receptor Tyrosine Kinase Activation. <i>Journal of Molecular Biology</i> , 2011, 412, 43-54.	2.0	26
94	A Look at Arginine in Membranes. <i>Journal of Membrane Biology</i> , 2011, 239, 49-56.	1.0	107
95	Antimicrobial Peptides: Successes, Challenges and Unanswered Questions. <i>Journal of Membrane Biology</i> , 2011, 239, 27-34.	1.0	406
96	FGFR3 Heterodimerization in Achondroplasia, the Most Common Form of Human Dwarfism. <i>Journal of Biological Chemistry</i> , 2011, 286, 13272-13281.	1.6	38
97	Electrically Addressable, Biologically Relevant Surface-Supported Bilayers. <i>Langmuir</i> , 2010, 26, 12054-12059.	1.6	18
98	Physical Basis behind Achondroplasia, the Most Common Form of Human Dwarfism. <i>Journal of Biological Chemistry</i> , 2010, 285, 30103-30114.	1.6	38
99	Receptor tyrosine kinase transmembrane domains. <i>Cell Adhesion and Migration</i> , 2010, 4, 249-254.	1.1	89
100	Assembly of the M2 Tetramer Is Strongly Modulated by Lipid Chain Length. <i>Biophysical Journal</i> , 2010, 99, 1810-1817.	0.2	28
101	Effect of a Polymer Cushion on the Electrical Properties and Stability of Surface-Supported Lipid Bilayers. <i>Langmuir</i> , 2010, 26, 3544-3548.	1.6	32
102	Measuring the Energetics of Membrane Protein Dimerization in Mammalian Membranes. <i>Journal of the American Chemical Society</i> , 2010, 132, 3628-3635.	6.6	121
103	The Extracellular Domain of Fibroblast Growth Factor Receptor 3 Inhibits Ligand-Independent Dimerization. <i>Science Signaling</i> , 2010, 3, ra86.	1.6	51
104	Viewing the Bilayer Hydrocarbon Core Using Neutron Diffraction. <i>Journal of Membrane Biology</i> , 2009, 227, 123-131.	1.0	5
105	Hill Coefficient Analysis of Transmembrane Helix Dimerization. <i>Journal of Membrane Biology</i> , 2009, 230, 49-55.	1.0	9
106	MPEX: A tool for exploring membrane proteins. <i>Protein Science</i> , 2009, 18, 2624-2628.	3.1	238
107	Polar residues in transmembrane helices can decrease electrophoretic mobility in polyacrylamide gels without causing helix dimerization. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2009, 1788, 1321-1331.	1.4	33
108	Increased expression of the integral membrane protein ErbB2 in Chinese hamster ovary cells expressing the anti-apoptotic gene Bcl-xL. <i>Protein Expression and Purification</i> , 2009, 67, 41-47.	0.6	13

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109	Utility of surface-supported bilayers in studies of transmembrane helix dimerization. <i>Journal of Structural Biology</i> , 2009, 168, 53-60.	1.3	17
110	Energetics of ErbB1 Transmembrane Domain Dimerization in Lipid Bilayers. <i>Biophysical Journal</i> , 2009, 96, 4622-4630.	0.2	59
111	Effect of the Ala391Glu mutation on FGFR3 activation in cellular membranes. <i>FASEB Journal</i> , 2009, 23, LB284.	0.2	0
112	Effect of short transmembrane peptides on the activation and dimerization of a FGFR3 pathogenic mutant. <i>FASEB Journal</i> , 2009, 23, LB234.	0.2	0
113	A simple "proximity" correction for Förster resonance energy transfer efficiency determination in membranes using lifetime measurements. <i>Analytical Biochemistry</i> , 2008, 380, 134-136.	1.1	20
114	Protein Folding in Membranes: Insights from Neutron Diffraction Studies of a Membrane β -Sheet Oligomer. <i>Biophysical Journal</i> , 2008, 94, 492-505.	0.2	15
115	Impedance spectroscopy of bilayer membranes on single crystal silicon. <i>Biointerphases</i> , 2008, 3, FA33-FA40.	0.6	24
116	Surface supported bilayer platform for studies of lateral association of proteins in membranes (Mini) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	0.6	6
117	Pathogenic Activation of Receptor Tyrosine Kinases in Mammalian Membranes. <i>Journal of Molecular Biology</i> , 2008, 384, 1130-1142.	2.0	42
118	Characterization of antimicrobial peptide activity by electrochemical impedance spectroscopy. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2008, 1778, 2430-2436.	1.4	46
119	Quantitative Measurements of Protein Interactions in a Crowded Cellular Environment. <i>Analytical Chemistry</i> , 2008, 80, 5976-5985.	3.2	38
120	Chapter 6 Förster Resonance Energy Transfer Measurements of Transmembrane Helix Dimerization Energetics. <i>Methods in Enzymology</i> , 2008, 450, 107-127.	0.4	29
121	Electrical Measurements of Bilayer Membranes Formed by Langmuir-Blodgett Deposition on Single-Crystal Silicon. <i>Langmuir</i> , 2007, 23, 13040-13045.	1.6	28
122	Influence of Applied Potential on the Impedance of Alkanethiol SAMs. <i>Langmuir</i> , 2007, 23, 9681-9685.	1.6	38
123	Effect of Pathogenic Cysteine Mutations on FGFR3 Transmembrane Domain Dimerization in Detergents and Lipid Bilayers. <i>Biochemistry</i> , 2007, 46, 11039-11046.	1.2	31
124	Studies of Receptor Tyrosine Kinase Transmembrane Domain Interactions: The EmEx-FRET Method. <i>Journal of Membrane Biology</i> , 2007, 215, 93-103.	1.0	44
125	Neutron Diffraction Studies of Fluid Bilayers with Transmembrane Proteins: Structural Consequences of the Achondroplasia Mutation. <i>Biophysical Journal</i> , 2006, 91, 3736-3747.	0.2	30
126	Role of Receptor Tyrosine Kinase Transmembrane Domains in Cell Signaling and Human Pathologies. <i>Biochemistry</i> , 2006, 45, 6241-6251.	1.2	212

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127	Bias-Dependent Admittance in Hybrid Bilayer Membranes. <i>Langmuir</i> , 2006, 22, 7156-7158.	1.6	8
128	The Achondroplasia Mutation Does Not Alter the Dimerization Energetics of the Fibroblast Growth Factor Receptor 3 Transmembrane Domain. <i>Biochemistry</i> , 2006, 45, 5551-5556.	1.2	44
129	Spectral Förster Resonance Energy Transfer Detection of Protein Interactions in Surface-Supported Bilayers. <i>Langmuir</i> , 2006, 22, 6986-6992.	1.6	25
130	FGFR3 Dimer Stabilization Due to a Single Amino Acid Pathogenic Mutation. <i>Journal of Molecular Biology</i> , 2006, 356, 600-612.	2.0	95
131	Transmembrane Helix Heterodimerization in Lipid Bilayers: Probing the Energetics behind Autosomal Dominant Growth Disorders. <i>Journal of Molecular Biology</i> , 2006, 358, 1-7.	2.0	36
132	Förster resonance energy transfer in liposomes: Measurements of transmembrane helix dimerization in the native bilayer environment. <i>Analytical Biochemistry</i> , 2005, 340, 154-164.	1.1	105
133	Synthesis and initial characterization of FGFR3 transmembrane domain: consequences of sequence modifications. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2005, 1668, 240-247.	1.4	28
134	Sodium Dodecyl Sulfate~Polyacrylamide Gel Electrophoresis and Förster Resonance Energy Transfer Suggest Weak Interactions between Fibroblast Growth Factor Receptor 3 (FGFR3) Transmembrane Domains in the Absence of Extracellular Domains and Ligands. <i>Biochemistry</i> , 2005, 44, 352-360.	1.2	72
135	An Experiment-Based Algorithm for Predicting the Partitioning of Unfolded Peptides into Phosphatidylcholine Bilayer Interfaces. <i>Biochemistry</i> , 2005, 44, 12614-12619.	1.2	47
136	Imaging Förster Resonance Energy Transfer Measurements of Transmembrane Helix Interactions in Lipid Bilayers on a Solid Support. <i>Langmuir</i> , 2004, 20, 9053-9060.	1.6	24
137	Energetics, stability, and prediction of transmembrane helices. Edited by G. von Heijne. <i>Journal of Molecular Biology</i> , 2001, 312, 927-934.	2.0	229
138	Structure, Location, and Lipid Perturbations of Melittin at the Membrane Interface. <i>Biophysical Journal</i> , 2001, 80, 801-811.	0.2	264
139	How Membranes Shape Protein Structure. <i>Journal of Biological Chemistry</i> , 2001, 276, 32395-32398.	1.6	273
140	An amphipathic α -helix at a membrane interface: a structural study using a novel X-ray diffraction method. Edited by D. C. Rees. <i>Journal of Molecular Biology</i> , 1999, 290, 99-117.	2.0	196
141	[4] Protein folding in membranes: Determining energetics of peptide-bilayer interactions. <i>Methods in Enzymology</i> , 1998, 295, 62-87.	0.4	233
142	Folding of β -sheet membrane proteins: a hydrophobic hexapeptide model. <i>Journal of Molecular Biology</i> , 1998, 277, 1091-1110.	2.0	195
143	Determination of the Hydrocarbon Core Structure of Fluid Dioleoylphosphocholine (DOPC) Bilayers by X-Ray Diffraction Using Specific Bromination of the Double-Bonds: Effect of Hydration. <i>Biophysical Journal</i> , 1998, 74, 2419-2433.	0.2	159
144	Critical Role of Lipid Composition in Membrane Permeabilization by Rabbit Neutrophil Defensins. <i>Journal of Biological Chemistry</i> , 1997, 272, 24224-24233.	1.6	135

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145	[23] Mechanism of leakage of contents of membrane vesicles determined by fluorescence reuquenching. <i>Methods in Enzymology</i> , 1997, 278, 474-486.	0.4	56
146	Interactions of Monomeric Rabbit Neutrophil Defensins with Bilayers:Â Comparison with Dimeric Human Defensin HNP-2â€. <i>Biochemistry</i> , 1996, 35, 11888-11894.	1.2	88
147	Effect of Bilayer Composition on the Phase Behavior of Liposomal Suspensions Containing Poly(ethylene glycol)-Lipids. <i>Macromolecules</i> , 1995, 28, 7693-7699.	2.2	101