

Byron Goldstein

List of Publications by Year in descending order

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78
papers

3,575
citations

185998

28
h-index

138251

58
g-index

80
all docs

80
docs citations

80
times ranked

2805
citing authors

#	ARTICLE	IF	CITATIONS
1	Combinatorial diversity of Syk recruitment driven by its multivalent engagement with FċRİ ³ . <i>Molecular Biology of the Cell</i> , 2019, 30, 2331-2347.	0.9	11
2	Membrane-Mediated Regulation of the Intrinsically Disordered CD3̇ [̇] Cytoplasmic Tail of the TCR. <i>Biophysical Journal</i> , 2015, 108, 2481-2491.	0.2	21
3	Estimating the Probability of Polyreactive Antibodies 4E10 and 2F5 Disabling a gp41 Trimer after T Cell-HIV Adhesion. <i>PLoS Computational Biology</i> , 2014, 10, e1003431.	1.5	3
4	Modeling and Simulation of Aggregation of Membrane Protein LAT with Molecular Variability in the Number of Binding Sites for Cytosolic Grb2-SOS1-Grb2. <i>PLoS ONE</i> , 2012, 7, e28758.	1.1	18
5	A Mechanistic Model of Early FċRİ Signaling: Lipid Rafts and the Question of Protection from Dephosphorylation. <i>PLoS ONE</i> , 2012, 7, e51669.	1.1	25
6	Guidelines for visualizing and annotating rule-based models. <i>Molecular BioSystems</i> , 2011, 7, 2779.	2.9	36
7	A Biophysical Model of Cell Adhesion Mediated by Immunoadhesin Drugs and Antibodies. <i>PLoS ONE</i> , 2011, 6, e19701.	1.1	3
8	Quantifying Intramolecular Binding in Multivalent Interactions: A Structure-Based Synergistic Study on Grb2-Sos1 Complex. <i>PLoS Computational Biology</i> , 2011, 7, e1002192.	1.5	17
9	Shaping the response: the role of FċRİ and Syk expression levels in mast cell signalling. <i>IET Systems Biology</i> , 2010, 4, 334-347.	0.8	12
10	A Detailed Mathematical Model Predicts That Serial Engagement of IgĖ-FċRİ Complexes Can Enhance Syk Activation in Mast Cells. <i>Journal of Immunology</i> , 2010, 185, 3268-3276.	0.4	14
11	Soluble MD2 increases TLR4 levels on the epithelial cell surface. <i>Cellular Immunology</i> , 2009, 255, 8-16.	1.4	12
12	Aggregation of Membrane Proteins by Cytosolic Cross-Linkers: Theory and Simulation of the LAT-Grb2-SOS1 System. <i>Biophysical Journal</i> , 2009, 96, 2604-2623.	0.2	63
13	Binding Mechanisms of PEGylated Ligands Reveal Multiple Effects of the PEG Scaffold. <i>Biochemistry</i> , 2008, 47, 1017-1030.	1.2	12
14	Kinetic Proofreading Model. <i>Advances in Experimental Medicine and Biology</i> , 2008, 640, 82-94.	0.8	16
15	Quantification and Modeling of Tripartite CD2-, CD58FC Chimera (Afacept)-, and CD16-mediated Cell Adhesion. <i>Journal of Biological Chemistry</i> , 2007, 282, 34748-34757.	1.6	23
16	Kinetic Proofreading of Ligand-FċRİ Interactions May Persist beyond LAT Phosphorylation. <i>Journal of Immunology</i> , 2007, 178, 3530-3535.	0.4	13
17	Diffusion Limited Reactions. <i>SIAM Journal on Applied Mathematics</i> , 2007, 67, 1147-1165.	0.8	5
18	A network model of early events in epidermal growth factor receptor signaling that accounts for combinatorial complexity. <i>BioSystems</i> , 2006, 83, 136-151.	0.9	141

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19	A Theoretical and Experimental Study of Competition Between Solution and Surface Receptors for Ligand in a Biacore Flow Cell. <i>Bulletin of Mathematical Biology</i> , 2006, 68, 1125-1150.	0.9	11
20	T cell activation: Kinetic proofreading, serial engagement and cell adhesion. <i>Journal of Computational and Applied Mathematics</i> , 2005, 184, 121-139.	1.1	25
21	'On-the-fly' or 'generate-first' modeling?. <i>Nature Biotechnology</i> , 2005, 23, 1344-1345.	9.4	16
22	Rule-based modeling of biochemical networks. <i>Complexity</i> , 2005, 10, 22-41.	0.9	110
23	T cell receptor binding kinetics required for T cell activation depend on the density of cognate ligand on the antigen-presenting cell. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 4824-4829.	3.3	151
24	BioNetGen: software for rule-based modeling of signal transduction based on the interactions of molecular domains. <i>Bioinformatics</i> , 2004, 20, 3289-3291.	1.8	381
25	Mathematical and computational models of immune-receptor signalling. <i>Nature Reviews Immunology</i> , 2004, 4, 445-456.	10.6	210
26	Effects of the Geometry of the Immunological Synapse on the Delivery of Effector Molecules. <i>Biophysical Journal</i> , 2004, 87, 2215-2220.	0.2	38
27	Equilibrium Thermodynamics of Cell-Cell Adhesion Mediated by Multiple Ligand-Receptor Pairs. <i>Biophysical Journal</i> , 2004, 86, 1408-1423.	0.2	85
28	The complexity of complexes in signal transduction. <i>Biotechnology and Bioengineering</i> , 2003, 84, 783-794.	1.7	176
29	Investigation of Early Events in Fc γ RI-Mediated Signaling Using a Detailed Mathematical Model. <i>Journal of Immunology</i> , 2003, 170, 3769-3781.	0.4	172
30	Analysis of Cholera Toxin α -Ganglioside Interactions by Flow Cytometry. <i>Biochemistry</i> , 2002, 41, 1742-1751.	1.2	120
31	Effective Rate Models for Receptors Distributed in a Layer above a Surface: Application to Cells and Biacore. <i>Biophysical Journal</i> , 2002, 82, 1743-1755.	0.2	45
32	Modeling the early signaling events mediated by Fc γ RI. <i>Molecular Immunology</i> , 2002, 38, 1213-1219.	1.0	54
33	Analysis of the Role of the Interleukin-2 Receptor β Chain in Ligand Binding. <i>Biochemistry</i> , 2002, 41, 2543-2551.	1.2	35
34	Kinetic Proofreading in Receptor-Mediated Transduction of Cellular Signals: Receptor Aggregation, Partially Activated Receptors, and Cytosolic Messengers. <i>Bulletin of Mathematical Biology</i> , 2002, 64, 887-911.	0.9	30
35	Activated TCRs remain marked for internalization after dissociation from pMHC. <i>Nature Immunology</i> , 2002, 3, 926-931.	7.0	103
36	Calculations Show Substantial Serial Engagement of T Cell Receptors. <i>Biophysical Journal</i> , 2001, 80, 606-612.	0.2	63

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37	Lattice Boltzmann Simulation of Diffusion-Convection Systems with Surface Chemical Reaction. Molecular Simulation, 2000, 25, 145-156.	0.9	65
38	A quantitative approach to signal transduction. Immunology Letters, 1999, 68, 53-57.	1.1	8
39	Transport effects on surface-volume biological reactions. Journal of Mathematical Biology, 1999, 39, 533-561.	0.8	24
40	The influence of transport on the kinetics of binding to surface receptors: application to cells and BIAcore. Journal of Molecular Recognition, 1999, 12, 293-299.	1.1	92
41	Effective rate models for the analysis of transport-dependent biosensor data. Mathematical Biosciences, 1999, 159, 123-144.	0.9	57
42	Solution assembly of the pseudo-high affinity and intermediate affinity interleukin-2 receptor complexes. Protein Science, 1999, 8, 482-489.	3.1	11
43	The influence of transport on the kinetics of binding to surface receptors: application to cells and BIAcore. , 1999, 12, 293.		2
44	Transport effects on surface-volume biological reactions. Journal of Mathematical Biology, 1999, 39, 533.	0.8	28
45	Extending the Range of Rate Constants Available from BIACORE: Interpreting Mass Transport-Influenced Binding Data. Biophysical Journal, 1998, 75, 583-594.	0.2	389
46	Altered Patterns of Tyrosine Phosphorylation and Syk Activation for Sterically Restricted Cyclic Dimers of IgE-Fc μ R1. Biochemistry, 1997, 36, 2237-2242.	1.2	22
47	The Fc Segment of IgE Influences the Kinetics of Dissociation of a Symmetrical Bivalent Ligand from Cyclic Dimeric Complexes. Biochemistry, 1996, 35, 5518-5527.	1.2	26
48	Solution assembly of cytokine receptor ectodomain complexes. Techniques in Protein Chemistry, 1996, , 45-56.	0.3	0
49	Kinetics of Tyrosine Phosphorylation When IgE Dimers Bind to FC γ 3 Receptors on Rat Basophilic Leukemia Cells. Journal of Biological Chemistry, 1995, 270, 20264-20272.	1.6	29
50	Solution Assembly of a Soluble, Heteromeric, High Affinity Interleukin-2 Receptor Complex. Journal of Biological Chemistry, 1995, 270, 16039-16044.	1.6	26
51	The kinetics of bivalent ligand-bivalent receptor aggregation: Ring formation and the breakdown of the equivalent site approximation. Mathematical Biosciences, 1995, 126, 171-190.	0.9	30
52	Evidence for p55-p75 heterodimers in the absence of IL-2 from Scatchard plot analysis. International Immunology, 1992, 4, 23-32.	1.8	26
53	Aggregation of IgE-receptor complexes on rat basophilic leukemia cells does not change the intrinsic affinity but can alter the kinetics of the ligand-IgE interaction. Biochemistry, 1992, 31, 5350-5356.	1.2	63
54	Interpretation of Scatchard plots for aggregating receptor systems. Mathematical Biosciences, 1992, 112, 115-154.	0.9	29

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55	Interaction of Interleukin-2 with its Cell Surface Receptors: Interpretation of Equilibrium Binding Experiments via Scatchard Plots. , 1992, , 117-148.		1
56	Dissociation kinetics of bivalent ligand-immunoglobulin E aggregates in solution. Biochemistry, 1991, 30, 2348-2356.	1.2	26
57	Bivalent ligand dissociation kinetics from receptor-bound immunoglobulin E: evidence for a time-dependent increase in ligand rebinding at the cell surface. Biochemistry, 1991, 30, 2357-2363.	1.2	31
58	Analysis of Ligand Binding and Cross-Linking of Receptors in Solution and on Cell Surfaces. , 1991, , 169-195.		0
59	The effect of co-operativity on the equilibrium binding of symmetric bivalent ligands to antibodies: Theoretical results with application to histamine release from basophils. Molecular Immunology, 1987, 24, 151-161.	1.0	15
60	Rates of diffusion-limited reaction in periodic systems. Journal of Statistical Physics, 1987, 49, 725-750.	0.5	12
61	Cross-Linking of IgE-Receptor complexes at the cell surface: A fluorescence method for studying the binding of monovalent and bivalent haptens to IgE. Molecular Immunology, 1986, 23, 769-781.	1.0	60
62	The time development of direct hemolytic plaques: Implications for the binding of IgM to cell surface haptens. Molecular Immunology, 1985, 22, 1323-1332.	1.0	0
63	Kinetic analysis of histamine release due to covalently linked ige dimers. Molecular Immunology, 1982, 19, 421-434.	1.0	16
64	A model of cell activation and desensitization by surface immunoglobulin: The case of histamine release from human basophils. Cell, 1980, 22, 59-67.	13.5	47
65	The effect of indicator layer thickness on the interpretation of hemolytic plaque results. Molecular Immunology, 1979, 16, 395-400.	1.0	0
66	A thermodynamic model of binding of flexible bivalent haptens to antibody. Immunochemistry, 1978, 15, 307-313.	1.3	18
67	A computer method for determining plaque size and plaque morphology. Journal of Immunological Methods, 1977, 16, 301-312.	0.6	3
68	The hemolytic plaque assay: theory for finite layers. Biophysical Chemistry, 1977, 7, 15-32.	1.5	5
69	Immunodiffusion in gels containing erythrocyte antigen ¹ I: Analysis of experiments involving the diffusion of antiserum from a circular well. Immunochemistry, 1976, 13, 29-33.	1.3	2
70	The electrophoretic hemolytic plaque assay - theory. Biophysical Chemistry, 1976, 4, 349-362.	1.5	7
71	Theory of hapten binding to IgM: The question of repulsive interactions between binding sites. Biophysical Chemistry, 1975, 3, 363-367.	1.5	21
72	Immunodiffusion in gels containing erythrocyte antigen I. Theory for diffusion of antiserum from a circular well. Journal of Theoretical Biology, 1975, 52, 317-334.	0.8	10

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73	Effect of antibody emission rates on plaque morphology. Nature, 1975, 253, 637-639.	13.7	14
74	On the mechanism of hemolytic plaque inhibition. Immunochemistry, 1974, 11, 661-665.	1.3	39
75	Theory of melting of the triple helix poly (A + 2U) for a 1 : 2 mixture of Poly A to Poly U. Biopolymers, 1973, 12, 461-475.	1.2	10
76	Anomalies in sedimentation. I. Stability theory of sedimenting entanglements in the "tight-bending" limit. Biopolymers, 1973, 12, 857-867.	1.2	6
77	Effect of Concentration and Intermolecular Forces on the Sedimentation of Polystyrene Spheres. Journal of Chemical Physics, 1971, 54, 4408-4413.	1.2	28
78	Theory of a Helix-to-Coil Transition in a Triple-Stranded Macromolecule. Journal of Chemical Physics, 1969, 51, 431-434.	1.2	7