

Raimond L Winslow

List of Publications by Year in descending order

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121
papers

7,608
citations

61984

43
h-index

53230

85
g-index

128
all docs

128
docs citations

128
times ranked

6165
citing authors

#	ARTICLE	IF	CITATIONS
1	Computational signatures for post-cardiac arrest trajectory prediction: Importance of early physiological time series. <i>Anaesthesia, Critical Care & Pain Medicine</i> , 2022, 41, 101015.	1.4	8
2	771: MACHINE LEARNING PREDICTION OF RESPONSIVENESS PHENOTYPES IN NON-NEUROLOGIC ICU PATIENTS. <i>Critical Care Medicine</i> , 2022, 50, 379-379.	0.9	0
3	Critical Requirements for the Initiation of a Cardiac Arrhythmia in Rat Ventricle: How Many Myocytes?. <i>Cells</i> , 2022, 11, 1878.	4.1	3
4	Prediction of Impending Septic Shock in Children With Sepsis. , 2021, 3, e0442.		7
5	Early Prediction of Multiple Organ Dysfunction in the Pediatric Intensive Care Unit. <i>Frontiers in Pediatrics</i> , 2021, 9, 711104.	1.9	10
6	PREDICTION OF PHYSIOLOGICAL DETERIORATION AND MORTALITY IN MECHANICALLY VENTILATED PATIENTS ADMITTED TO THE ICU. <i>Chest</i> , 2021, 160, A1061-A1062.	0.8	0
7	Estimating ectopic beat probability with simplified statistical models that account for experimental uncertainty. <i>PLoS Computational Biology</i> , 2021, 17, e1009536.	3.2	1
8	356: Predicting Hypoxemia in ICU Patients. <i>Critical Care Medicine</i> , 2021, 49, 167-167.	0.9	1
9	Predicting Flow Rate Escalation for Pediatric Patients on High Flow Nasal Cannula Using Machine Learning. <i>Frontiers in Pediatrics</i> , 2021, 9, 734753.	1.9	2
10	SWIFT: A deep learning approach to prediction of hypoxemic events in critically-ill patients using SpO2 waveform prediction. <i>PLoS Computational Biology</i> , 2021, 17, e1009712.	3.2	5
11	Simplified Models Predict Cellular Arrhythmia Probabilities and Reveal the Impact of Experimental Parameter Uncertainty on the Predicted Distribution of Arrhythmic Events. <i>Biophysical Journal</i> , 2020, 118, 409a.	0.5	0
12	Spectral clustering of risk score trajectories stratifies sepsis patients by clinical outcome and interventions received. <i>ELife</i> , 2020, 9, .	6.0	15
13	Natural Language Processing of Clinical Notes for Improved Early Prediction of Septic Shock in the ICU. , 2019, 2019, 6103-6108.		19
14	Data-driven discovery of a novel sepsis pre-shock state predicts impending septic shock in the ICU. <i>Scientific Reports</i> , 2019, 9, 6145.	3.3	56
15	Na ⁺ microdomains and sparks: Role in cardiac excitation-contraction coupling and arrhythmias in ankyrin-B deficiency. <i>Journal of Molecular and Cellular Cardiology</i> , 2019, 128, 145-157.	1.9	10
16	Please Welcome the New Team Member. <i>Pediatric Critical Care Medicine</i> , 2019, 20, 1200-1201.	0.5	8
17	A bilobal model of Ca ²⁺ -dependent inactivation to probe the physiology of L-type Ca ²⁺ channels. <i>Journal of General Physiology</i> , 2018, 150, 1688-1701.	1.9	10
18	Mechanisms of the cyclic nucleotide cross-talk signaling network in cardiac L-type calcium channel regulation. <i>Journal of Molecular and Cellular Cardiology</i> , 2017, 106, 29-44.	1.9	7

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19	Estimating the probabilities of rare arrhythmic events in multiscale computational models of cardiac cells and tissue. <i>PLoS Computational Biology</i> , 2017, 13, e1005783.	3.2	16
20	Modeling Na ⁺ -Ca ²⁺ exchange in the heart: Allosteric activation, spatial localization, sparks and excitation-contraction coupling. <i>Journal of Molecular and Cellular Cardiology</i> , 2016, 99, 174-187.	1.9	20
21	Modeling calcium regulation of contraction, energetics, signaling, and transcription in the cardiac myocyte. <i>Wiley Interdisciplinary Reviews: Systems Biology and Medicine</i> , 2016, 8, 37-67.	6.6	31
22	WaveformECG: A Platform for Visualizing, Annotating, and Analyzing ECG Data. <i>Computing in Science and Engineering</i> , 2016, 18, 36-46.	1.2	9
23	Roles of phosphodiesterases in the regulation of the cardiac cyclic nucleotide cross-talk signaling network. <i>Journal of Molecular and Cellular Cardiology</i> , 2016, 91, 215-227.	1.9	33
24	Models and Simulations as a Service: Exploring the Use of Galaxy for Delivering Computational Models. <i>Biophysical Journal</i> , 2016, 110, 1038-1043.	0.5	9
25	Shape analysis of hypertrophic and hypertensive heart disease using MRI-based 3D surface models of left ventricular geometry. <i>Medical Image Analysis</i> , 2016, 29, 12-23.	11.6	12
26	Metadata-driven Clinical Data Loading into i2b2 for Clinical and Translational Science Institutes. <i>AMIA Summits on Translational Science Proceedings</i> , 2016, 2016, 184-93.	0.4	6
27	On the Adjacency Matrix of RyR2 Cluster Structures. <i>PLoS Computational Biology</i> , 2015, 11, e1004521.	3.2	33
28	Roles of Phosphodiesterases in Cyclic Nucleotide Cross-Talk in Cardiac Myocytes. <i>Biophysical Journal</i> , 2015, 108, 260a.	0.5	0
29	Dynamics of Ca ²⁺ -Dependent Regulation of the Cardiac Na ⁺ /Ca ²⁺ Exchanger. <i>Biophysical Journal</i> , 2015, 108, 265a.	0.5	0
30	Lessons on Quality Control in Large Scale Imaging Trials: the Multi-Ethnic Study of Atherosclerosis (MESA). <i>Current Cardiovascular Imaging Reports</i> , 2015, 8, 1.	0.6	5
31	Interaction between phosphodiesterases in the regulation of the cardiac β^2 -adrenergic pathway. <i>Journal of Molecular and Cellular Cardiology</i> , 2015, 88, 29-38.	1.9	22
32	Mechanistic Investigation of the Arrhythmogenic Role of Oxidized CaMKII in the Heart. <i>Biophysical Journal</i> , 2015, 109, 838-849.	0.5	27
33	Modeling CaMKII-mediated regulation of L-type Ca ²⁺ channels and ryanodine receptors in the heart. <i>Frontiers in Pharmacology</i> , 2014, 5, 60.	3.5	11
34	Systems biology approaches to understanding the cause and treatment of heart, lung, blood, and sleep disorders. <i>Frontiers in Physiology</i> , 2014, 5, 107.	2.8	1
35	Superresolution Modeling of Calcium Release in the Heart. <i>Biophysical Journal</i> , 2014, 107, 3018-3029.	0.5	96
36	Integrating Mitochondrial Energetics, Redox and ROS Metabolic Networks: A Two-Compartment Model. <i>Biophysical Journal</i> , 2013, 104, 332-343.	0.5	94

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37	Extinguishing the Sparks. <i>Biophysical Journal</i> , 2013, 104, 2115-2117.	0.5	6
38	A Computational Model of Reactive Oxygen Species and Redox Balance in Cardiac Mitochondria. <i>Biophysical Journal</i> , 2013, 105, 1045-1056.	0.5	55
39	An Integrated Mitochondrial ROS Production and Scavenging Model: Implications for Heart Failure. <i>Biophysical Journal</i> , 2013, 105, 2832-2842.	0.5	36
40	Mechanistic Electron Transport Chain Model Explains ROS Production in Different Respiratory Modes. <i>Biophysical Journal</i> , 2013, 104, 304a-305a.	0.5	0
41	Cardiac Resynchronization Therapy Improves Altered Na Channel Gating in Canine Model of Dyssynchronous Heart Failure. <i>Circulation: Arrhythmia and Electrophysiology</i> , 2013, 6, 546-554.	4.8	27
42	Screening Entire Health System ECG Databases to Identify Patients at Increased Risk of Death. <i>Circulation: Arrhythmia and Electrophysiology</i> , 2013, 6, 1156-1162.	4.8	29
43	Toward an Integrative Computational Model of the Guinea Pig Cardiac Myocyte. <i>Frontiers in Physiology</i> , 2012, 3, 244.	2.8	25
44	Computational Medicine: Translating Models to Clinical Care. <i>Science Translational Medicine</i> , 2012, 4, 158rv11.	12.4	171
45	Dynamics of matrix-free Ca ²⁺ in cardiac mitochondria: two components of Ca ²⁺ uptake and role of phosphate buffering. <i>Journal of General Physiology</i> , 2012, 139, 465-478.	1.9	69
46	Local control model illustrates how action potential morphology affects Ca ²⁺ release. <i>FASEB Journal</i> , 2012, 26, 1053.1.	0.5	0
47	Ontological Labels for Automated Location of Left Ventricular Remodeling. , 2011, , .		0
48	Grand Challenges in Computational Physiology and Medicine. <i>Frontiers in Physiology</i> , 2011, 2, 79.	2.8	12
49	Cardiac myocytes and local signaling in nano-domains. <i>Progress in Biophysics and Molecular Biology</i> , 2011, 107, 48-59.	2.9	17
50	Integrative modeling of the cardiac ventricular myocyte. <i>Wiley Interdisciplinary Reviews: Systems Biology and Medicine</i> , 2011, 3, 392-413.	6.6	30
51	CaMKII-dependent activation of late I _{Na} contributes to cellular arrhythmia in a model of the cardiac myocyte. , 2011, 2011, 4665-8.		10
52	Integrative Systems Models of Cardiac Excitation-Contractile Coupling. <i>Circulation Research</i> , 2011, 108, 70-84.	4.5	56
53	A Reaction-Diffusion Model of ROS-Induced ROS Release in a Mitochondrial Network. <i>PLoS Computational Biology</i> , 2010, 6, e1000657.	3.2	131
54	Role of CaMKII in RyR leak, EC coupling and action potential duration: A computational model. <i>Journal of Molecular and Cellular Cardiology</i> , 2010, 49, 617-624.	1.9	57

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55	Control and Regulation of Integrated Mitochondrial Function in Metabolic and Transport Networks. International Journal of Molecular Sciences, 2009, 10, 1500-1513.	4.1	25
56	Identification of Novel Serological Biomarkers for Inflammatory Bowel Disease Using Escherichia coli Proteome Chip. Molecular and Cellular Proteomics, 2009, 8, 1765-1776.	3.8	63
57	Computational Method for Identifying and Quantifying Shape Features of Human Left Ventricular Remodeling. Annals of Biomedical Engineering, 2009, 37, 1043-1054.	2.5	24
58	Improved reproducibility of reverse-phase protein microarrays using array microenvironment normalization. Proteomics, 2009, 9, 5562-5566.	2.2	31
59	CaMKII-Induced Shift in Modal Gating Explains L-Type Ca ²⁺ Current Facilitation: A Modeling Study. Biophysical Journal, 2009, 96, 1770-1785.	0.5	48
60	Control and Regulation of Mitochondrial Energetics in an Integrated Model of Cardiomyocyte Function. Biophysical Journal, 2009, 96, 2466-2478.	0.5	70
61	Modeling Cardiac Action Potential Shortening Driven by Oxidative Stress-Induced Mitochondrial Oscillations in Guinea Pig Cardiomyocytes. Biophysical Journal, 2009, 97, 1843-1852.	0.5	77
62	CaMKII-Induced Shift in Modal Gating Explains L-type Ca ²⁺ Current Facilitation: A Modeling Study. Biophysical Journal, 2009, 96, 540a.	0.5	1
63	Transport of Relational Structures in Groups of Diffeomorphisms. Journal of Mathematical Imaging and Vision, 2008, 32, 41-56.	1.3	44
64	Allele-specific expression in the germline of patients with familial pancreatic cancer: An unbiased approach to cancer gene discovery. Cancer Biology and Therapy, 2008, 7, 135-144.	3.4	42
65	Key pathways associated with heart failure development revealed by gene networks correlated with cardiac remodeling. Physiological Genomics, 2008, 35, 222-230.	2.3	44
66	Protein Geometry and Placement in the Cardiac Dyad Influence Macroscopic Properties of Calcium-Induced Calcium Release. Biophysical Journal, 2007, 92, 3379-3396.	0.5	57
67	Discovering robust protein biomarkers for disease from relative expression reversals in 2-D DIGE data.. Proteomics, 2007, 7, 1197-1207.	2.2	21
68	Integrative Structurally Detailed Model of Calcium Dynamics in the Cardiac Diad. Multiscale Modeling and Simulation, 2006, 5, 1280-1296.	1.6	3
69	AB34-6. Heart Rhythm, 2006, 3, S72.	0.7	3
70	Mechanisms of Excitation-Contraction Coupling in an Integrative Model of the Cardiac Ventricular Myocyte. Biophysical Journal, 2006, 90, 77-91.	0.5	133
71	A Computational Model Integrating Electrophysiology, Contraction, and Mitochondrial Bioenergetics in the Ventricular Myocyte. Biophysical Journal, 2006, 91, 1564-1589.	0.5	198
72	Transcriptomic profiling of the canine tachycardia-induced heart failure model: global comparison to human and murine heart failure. Journal of Molecular and Cellular Cardiology, 2006, 40, 76-86.	1.9	51

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73	MASCOT HTML and XML parser: An implementation of a novel object model for protein identification data. <i>Proteomics</i> , 2006, 6, 5688-5693.	2.2	7
74	Multiscale Modeling of Calcium Signaling in the Cardiac Dyad. <i>Annals of the New York Academy of Sciences</i> , 2006, 1080, 362-375.	3.8	19
75	Evidence of Structural Remodeling in the Dyssynchronous Failing Heart. <i>Circulation Research</i> , 2006, 98, 125-132.	4.5	573
76	Measuring and Mapping Cardiac Fiber and Laminar Architecture Using Diffusion Tensor MR Imaging. <i>Annals of the New York Academy of Sciences</i> , 2005, 1047, 296-307.	3.8	216
77	Using models of the myocyte for functional interpretation of cardiac proteomic data. <i>Journal of Physiology</i> , 2005, 563, 73-81.	2.9	19
78	Ex vivo 3D diffusion tensor imaging and quantification of cardiac laminar structure. <i>Magnetic Resonance in Medicine</i> , 2005, 54, 850-859.	3.0	208
79	Candidate Gene Discovery in Cardiovascular Disease. <i>Circulation Research</i> , 2005, 96, 605-606.	4.5	5
80	The Role of Stochastic and Modal Gating of Cardiac L-Type Ca ²⁺ Channels on Early After-Depolarizations. <i>Biophysical Journal</i> , 2005, 88, 85-95.	0.5	138
81	The Ongoing Journey to Understand Heart Function Through Integrative Modeling. <i>Circulation Research</i> , 2004, 95, 1135-1136.	4.5	4
82	Modeling the Actions of β^2 -Adrenergic Signaling on Excitation-Contraction Coupling Processes. <i>Annals of the New York Academy of Sciences</i> , 2004, 1015, 16-27.	3.8	27
83	A Mitochondrial Oscillator Dependent on Reactive Oxygen Species. <i>Biophysical Journal</i> , 2004, 87, 2060-2073.	0.5	206
84	A Computational Model of the Human Left-Ventricular Epicardial Myocyte. <i>Biophysical Journal</i> , 2004, 87, 1507-1525.	0.5	244
85	Classifying Gene Expression Profiles from Pairwise mRNA Comparisons. <i>Statistical Applications in Genetics and Molecular Biology</i> , 2004, 3, 1-19.	0.6	297
86	Gene expression profiles in end-stage human idiopathic dilated cardiomyopathy: altered expression of apoptotic and cytoskeletal genes. <i>Genomics</i> , 2004, 83, 281-297.	2.9	89
87	An Integrated Model of Cardiac Mitochondrial Energy Metabolism and Calcium Dynamics. <i>Biophysical Journal</i> , 2003, 84, 2734-2755.	0.5	345
88	Role of Sodium-Calcium Exchanger in Modulating the Action Potential of Ventricular Myocytes From Normal and Failing Hearts. <i>Circulation Research</i> , 2003, 93, 46-53.	4.5	165
89	Genome Informatics. <i>Circulation Research</i> , 2003, 92, 953-961.	4.5	43
90	An Integrative Model of the Cardiac Ventricular Myocyte Incorporating Local Control of Ca ²⁺ Release. <i>Biophysical Journal</i> , 2002, 83, 2918-2945.	0.5	173

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91	Ectopic expression of KCNE3 accelerates cardiac repolarization and abbreviates the QT interval. <i>Journal of Clinical Investigation</i> , 2002, 109, 1083-1090.	8.2	28
92	Imaging-based integrative models of the heart: closing the loop between experiment and simulation. <i>Novartis Foundation Symposium</i> , 2002, 247, 129-41; discussion 141-3, 144-50, 244-52.	1.1	1
93	Final Report of the Meeting "Modeling & Simulation in Medicine: Towards an Integrated Framework" July 20-21, 2000, National Library of Medicine, National Institutes of Health, Bethesda, Maryland, USA. <i>Computer Aided Surgery</i> , 2001, 6, 32-39.	1.8	5
94	Molecular Interactions Between Two Long-QT Syndrome Gene Products, HERG and KCNE2 , Rationalized by In Vitro and In Silico Analysis. <i>Circulation Research</i> , 2001, 89, 33-38.	4.5	104
95	Direct histological validation of diffusion tensor MRI in formaldehyde-fixed myocardium. <i>Magnetic Resonance in Medicine</i> , 2000, 44, 157-161.	3.0	208
96	Modeling short-term interval-force relations in cardiac muscle. <i>American Journal of Physiology - Heart and Circulatory Physiology</i> , 2000, 278, H913-H931.	3.2	60
97	Role of the Calcium-Independent Transient Outward Current I_{to1} in Shaping Action Potential Morphology and Duration. <i>Circulation Research</i> , 2000, 87, 1026-1033.	4.5	197
98	Computational modelling of biological systems: tools and visions. <i>Philosophical Transactions Series A, Mathematical, Physical, and Engineering Sciences</i> , 2000, 358, 579-610.	3.4	84
99	Direct histological validation of diffusion tensor MRI in formaldehyde-fixed myocardium. <i>Magnetic Resonance in Medicine</i> , 2000, 44, 157-161.	3.0	2
100	Comparison of putative cooperative mechanisms in cardiac muscle: length dependence and dynamic responses. <i>American Journal of Physiology - Heart and Circulatory Physiology</i> , 1999, 276, H1734-H1754.	3.2	102
101	Optimal Detection of Flash Intensity Differences Using Rod Photocurrent Observations. <i>Neural Computation</i> , 1999, 11, 1097-1111.	2.2	4
102	Mechanisms of Altered Excitation-Contraction Coupling in Canine Tachycardia-Induced Heart Failure, II. <i>Circulation Research</i> , 1999, 84, 571-586.	4.5	557
103	Modeling Gain and Gradedness of Ca ²⁺ Release in the Functional Unit of the Cardiac Diadic Space. <i>Biophysical Journal</i> , 1999, 77, 1871-1884.	0.5	105
104	Cardiac Sodium Channel Markov Model with Temperature Dependence and Recovery from Inactivation. <i>Biophysical Journal</i> , 1999, 76, 1868-1885.	0.5	70
105	Modeling Short-Term Interval-Force Relations in Cardiac Muscle. <i>Annals of the New York Academy of Sciences</i> , 1998, 853, 345-349.	3.8	7
106	Modeling the cellular basis of altered excitation-contraction coupling in heart failure. <i>Progress in Biophysics and Molecular Biology</i> , 1998, 69, 497-514.	2.9	35
107	Model Studies of the Role of Mechano-sensitive Currents in the Generation of Cardiac Arrhythmias. <i>Journal of Theoretical Biology</i> , 1998, 190, 295-312.	1.7	32
108	Cardiac Ca ²⁺ Dynamics: The Roles of Ryanodine Receptor Adaptation and Sarcoplasmic Reticulum Load. <i>Biophysical Journal</i> , 1998, 74, 1149-1168.	0.5	300

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109	Functional Roles of Sodium-Calcium Exchange in Normal and Abnormal Cardiac Rhythm. <i>Annals of the New York Academy of Sciences</i> , 1996, 779, 480-488.	3.8	12
110	Generation and propagation of normal and abnormal pacemaker activity in network models of cardiac sinus node and atrium. <i>Chaos, Solitons and Fractals</i> , 1995, 5, 491-512.	5.1	22
111	Geometric Properties of the Chaotic Saddle Responsible for Supertransients in Spatiotemporal Chaotic Systems. <i>Physical Review Letters</i> , 1995, 74, 5208-5211.	7.8	78
112	The Functional Role of Excitatory and Inhibitory Interactions in Chopper Cells of the Anteroventral Cochlear Nucleus. <i>Neural Computation</i> , 1994, 6, 1127-1140.	2.2	7
113	Riddled parameter space in spatiotemporal chaotic dynamical systems. <i>Physical Review Letters</i> , 1994, 72, 1640-1643.	7.8	48
114	A model of selective processing of auditory-nerve inputs by stellate cells of the antero-ventral cochlear nucleus. <i>Journal of Computational Neuroscience</i> , 1994, 1, 167-194.	1.0	25
115	Extreme sensitive dependence on parameters and initial conditions in spatio-temporal chaotic dynamical systems. <i>Physica D: Nonlinear Phenomena</i> , 1994, 74, 353-371.	2.8	36
116	Simulating cardiac sinus and atrial network dynamics on the connection machine. <i>Physica D: Nonlinear Phenomena</i> , 1993, 64, 281-298.	2.8	92
117	Complex dynamics in coupled cardiac pacemaker cells. <i>Physical Review Letters</i> , 1993, 71, 2501-2504.	7.8	25
118	Dynamic models of the retinal horizontal cell network. <i>Progress in Biophysics and Molecular Biology</i> , 1991, 56, 107-133.	2.9	6
119	A computational model for rate-level functions from cat auditory-nerve fibers. <i>Hearing Research</i> , 1989, 41, 61-69.	2.0	89
120	Single-tone intensity discrimination based on auditory-nerve rate responses in backgrounds of quiet, noise, and with stimulation of the crossed olivocochlear bundle. <i>Hearing Research</i> , 1988, 35, 165-189.	2.0	228
121	Imaging-Based Integrative Models of the Heart: Closing the Loop between Experiment and Simulation. <i>Novartis Foundation Symposium</i> , 0, , 129-143.	1.1	4