

Takumi Washio

List of Publications by Year in descending order

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Version: 2024-02-01

77
papers

1,220
citations

361413

20
h-index

395702

33
g-index

80
all docs

80
docs citations

80
times ranked

1055
citing authors

#	ARTICLE	IF	CITATIONS
1	UT-Heart: A Finite Element Model Designed for the Multiscale and Multiphysics Integration of our Knowledge on the Human Heart. <i>Methods in Molecular Biology</i> , 2022, , 221-245.	0.9	3
2	Using incomplete Cholesky factorization to increase the time step in molecular dynamics simulations. <i>Journal of Computational and Applied Mathematics</i> , 2022, 415, 114519.	2.0	1
3	Chloroquine and hydroxychloroquine provoke arrhythmias at concentrations higher than those clinically used to treat COVID-19: A simulation study. <i>Clinical and Translational Science</i> , 2021, 14, 1092-1100.	3.1	9
4	An application of a patient-specific cardiac simulator for the prediction of outcomes after mitral valve replacement: a pilot study. <i>Journal of Artificial Organs</i> , 2021, 24, 351-357.	0.9	1
5	A reverse stroke characterizes the force generation of cardiac myofilaments, leading to an understanding of heart function. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	16
6	A Multiple Step Active Stiffness Integration Scheme to Couple a Stochastic Cross-Bridge Model and Continuum Mechanics for Uses in Both Basic Research and Clinical Applications of Heart Simulation. <i>Frontiers in Physiology</i> , 2021, 12, 712816.	2.8	6
7	Semi-Implicit Time Integration with Hessian Eigenvalue Corrections for a Larger Time Step in Molecular Dynamics Simulations. <i>Journal of Chemical Theory and Computation</i> , 2021, 17, 5792-5804.	5.3	2
8	Ionic mechanisms of ST segment elevation in electrocardiogram during acute myocardial infarction. <i>Journal of Physiological Sciences</i> , 2020, 70, 36.	2.1	10
9	Mechanism of contraction rhythm homeostasis for hyperthermal sarcomeric oscillations of neonatal cardiomyocytes. <i>Scientific Reports</i> , 2020, 10, 20468.	3.3	8
10	Using Systolic Local Mechanical Load to Predict Fiber Orientation in Ventricles. <i>Frontiers in Physiology</i> , 2020, 11, 467.	2.8	6
11	Personalized Perioperative Multi-scale, Multi-physics Heart Simulation of Double Outlet Right Ventricle. <i>Annals of Biomedical Engineering</i> , 2020, 48, 1740-1750.	2.5	14
12	Patient-specific heart simulation can identify non-responders to cardiac resynchronization therapy. <i>Heart and Vessels</i> , 2020, 35, 1135-1147.	1.2	10
13	Abstract 13732: Evaluation of the Mechanical Effect of Thickened IPS Derived Cardiomyocyte Patch on the Distressed Left Ventricle Using Cardiac Simulator "Ut-heart". <i>Circulation</i> , 2020, 142, .	1.6	0
14	Effect of myofibril passive elastic properties on the mechanical communication between motor proteins on adjacent sarcomeres. <i>Scientific Reports</i> , 2019, 9, 9355.	3.3	8
15	Longitudinal dissociation and transition in thickness of the His-Purkinje system cause various QRS waveforms of surface ECG under His bundle pacing: A simulation study based on clinical observations. <i>Journal of Cardiovascular Electrophysiology</i> , 2019, 30, 2582-2590.	1.7	4
16	Clinical and pharmacological application of multiscale multiphysics heart simulator, UT-Heart. <i>Korean Journal of Physiology and Pharmacology</i> , 2019, 23, 295.	1.2	5
17	Proposed mechanism for the length dependence of the force developed in maximally activated muscles. <i>Scientific Reports</i> , 2019, 9, 1317.	3.3	12
18	Absence of Rapid Propagation through the Purkinje Network as a Potential Cause of Line Block in the Human Heart with Left Bundle Branch Block. <i>Frontiers in Physiology</i> , 2018, 9, 56.	2.8	1

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19	Coupling Langevin Dynamics With Continuum Mechanics: Exposing the Role of Sarcomere Stretch Activation Mechanisms to Cardiac Function. <i>Frontiers in Physiology</i> , 2018, 9, 333.	2.8	10
20	Arrhythmic hazard map for a 3D whole ventricle model under multiple ion channel block. <i>British Journal of Pharmacology</i> , 2018, 175, 3435-3452.	5.4	21
21	Recent Advances in the Computer Simulation of Heart. <i>The Journal of the Japanese Society of Internal Medicine</i> , 2018, 107, 2532-2538.	0.0	0
22	Analysis of spontaneous oscillations for a three-state power-stroke model. <i>Physical Review E</i> , 2017, 95, 022411.	2.1	11
23	Molecular Mechanism of Synchronous Force Generations among Myosin Molecules. <i>Biophysical Journal</i> , 2017, 112, 263a.	0.5	0
24	Multi-scale, tailor-made heart simulation can predict the effect of cardiac resynchronization therapy. <i>Journal of Molecular and Cellular Cardiology</i> , 2017, 108, 17-23.	1.9	46
25	Titin-mediated thick filament activation, through a mechanosensing mechanism, introduces sarcomere-length dependencies in mathematical models of rat trabecula and whole ventricle. <i>Scientific Reports</i> , 2017, 7, 5546.	3.3	20
26	Coordinated force generation of skeletal myosins in myofilaments through motor coupling. <i>Nature Communications</i> , 2017, 8, 16036.	12.8	55
27	Deformable regions of interest with multiple points for tissue tracking in echocardiography. <i>Medical Image Analysis</i> , 2017, 35, 554-569.	11.6	5
28	Including Thermal Fluctuations in Actomyosin Stable States Increases the Predicted Force per Motor and Macroscopic Efficiency in Muscle Modelling. <i>PLoS Computational Biology</i> , 2016, 12, e1005083.	3.2	20
29	Ventricular fiber optimization utilizing the branching structure. <i>International Journal for Numerical Methods in Biomedical Engineering</i> , 2016, 32, e02753.	2.1	21
30	Oscillatory Behavior in Muscle Myosin. <i>Biophysical Journal</i> , 2016, 110, 463a.	0.5	0
31	Mechanism of Cooperative Force Generations between Skeletal Myosins. <i>Biophysical Journal</i> , 2016, 110, 614a.	0.5	0
32	Tailor-made heart simulation predicts the effect of cardiac resynchronization therapy in a canine model of heart failure. <i>Medical Image Analysis</i> , 2016, 31, 46-62.	11.6	20
33	Cardiac safety assessment of drugs using three-dimensional heart simulator. <i>Journal of Pharmacological and Toxicological Methods</i> , 2016, 81, 351.	0.7	0
34	Intermolecular Cooperativity of Skeletal Myosins Enhances Force Output in Myofilaments. <i>Biophysical Journal</i> , 2015, 108, 338a.	0.5	0
35	An integrated finite element simulation of cardiomyocyte function based on triphasic theory. <i>Frontiers in Physiology</i> , 2015, 6, 287.	2.8	9
36	Distinct Functional Roles of Cardiac Mitochondrial Subpopulations Revealed by a 3D Simulation Model. <i>Biophysical Journal</i> , 2015, 108, 2732-2739.	0.5	17

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37	From Molecule to Organ: A Multiscale Simulator of Heart Contraction. <i>Biophysical Journal</i> , 2015, 108, 443a.	0.5	1
38	Screening system for drug-induced arrhythmogenic risk combining a patch clamp and heart simulator. <i>Science Advances</i> , 2015, 1, e1400142.	10.3	87
39	Integrated 3D Simulation of Cardiomyocyte Revealed the Distinct Functional Characteristics between Subsarcolemmal and Interfibrillar Mitochondria. <i>Biophysical Journal</i> , 2014, 106, 643a.	0.5	0
40	Multiscale Heart Simulation with Cooperative Stochastic Cross-Bridge Dynamics and Cellular Structures. <i>Multiscale Modeling and Simulation</i> , 2013, 11, 965-999.	1.6	47
41	Patient Specific Simulation of Body Surface ECG using the Finite Element Method. <i>PACE - Pacing and Clinical Electrophysiology</i> , 2013, 36, 309-321.	1.2	32
42	Mitochondrial Colocalization with Ca ²⁺ Release Sites is Crucial to Cardiac Metabolism. <i>Biophysical Journal</i> , 2013, 104, 496-504.	0.5	19
43	Tailor-made Medicine Using the Multi-scale Heart Simulator "UT-Heart". <i>Journal of Cardiac Failure</i> , 2013, 19, S107.	1.7	0
44	3SDA-05 A numerical model of cross-bridge cycling and its application to a beating human heart(3SDA) Tj ETQq0 0.0 rgBT /Oerlock 10	0.1	0
45	A Study on Large Scale Analysis of Cardiomyocyte Coupling Electrical, Chemical and Mechanical Phenomena Based on Triphasic Theory. <i>Nihon Kikai Gakkai Ronbunshu, A Hen/Transactions of the Japan Society of Mechanical Engineers, Part A</i> , 2013, 79, 934-949.	0.2	1
46	Multi-scale Multi-physics Heart Simulator UT-Heart. <i>Journal of the Society of Mechanical Engineers</i> , 2013, 116, 74-77.	0.0	0
47	OS1012 Multi-physics Analysis of Intra-cardiomyocyte Phenomena Based on Triphasic Theory. <i>The Proceedings of the Materials and Mechanics Conference</i> , 2013, 2013, _OS1012-1_-_OS1012-3_.	0.0	0
48	Multi-scale simulations of cardiac electrophysiology and mechanics using the University of Tokyo heart simulator. <i>Progress in Biophysics and Molecular Biology</i> , 2012, 110, 380-389.	2.9	83
49	A 3D Integrated Model of Cardiomyocytes Revealed the Important Role of Cardiac T-Tubule Structure for the Maintenance of Contractile Function. <i>Biophysical Journal</i> , 2012, 102, 592a.	0.5	0
50	Approximation for Cooperative Interactions of a Spatially-Detailed Cardiac Sarcomere Model. <i>Cellular and Molecular Bioengineering</i> , 2012, 5, 113-126.	2.1	33
51	A Three-Dimensional Simulation Model of Cardiomyocyte Integrating Excitation-Contraction Coupling and Metabolism. <i>Biophysical Journal</i> , 2011, 101, 2601-2610.	0.5	54
52	Convergence analysis of inexact LU-type preconditioners for indefinite problems arising in incompressible continuum analysis. <i>Japan Journal of Industrial and Applied Mathematics</i> , 2011, 28, 89-117.	0.9	3
53	Transmural and apicobasal gradients in repolarization contribute to T-wave genesis in human surface ECG. <i>American Journal of Physiology - Heart and Circulatory Physiology</i> , 2011, 301, H200-H208.	3.2	64
54	Multiscale Multiphysics Heart Simulator UT-Heart. <i>The Proceedings of the Symposium on Micro-Nano Science and Technology</i> , 2011, 2011.3, 1-2.	0.0	0

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55	A Study on Mechano-Electrochemical Modeling of Cardiomyocyte. Nihon Kikai Gakkai Ronbunshu, A Hen/Transactions of the Japan Society of Mechanical Engineers, Part A, 2010, 76, 1806-1815.	0.2	2
56	Study of efficient homogenization algorithms for nonlinear problems. Computational Mechanics, 2010, 46, 247-258.	4.0	22
57	A Parallel Multilevel Technique for Solving the Bidomain Equation on a Human Heart with Purkinje Fibers and a Torso Model. SIAM Review, 2010, 52, 717-743.	9.5	30
58	A Multi-Scale Heart Simulation on Massively Parallel Computers. , 2010, , .		13
59	Large-scale integrated model is useful for understanding heart mechanisms and developments of medical therapy. , 2009, 2009, 2347-50.		1
60	Nonlinear Homogenization Algorithms with Low Computational Cost. Journal of Computational Science and Technology, 2009, 3, 101-114.	0.4	1
61	A Parallel Multilevel Technique for Solving the Bidomain Equation on a Human Heart with Purkinje Fibers and a Torso Model. SIAM Journal of Scientific Computing, 2008, 30, 2855-2881.	2.8	9
62	2309 Opening and Closing Simulation of Mitral Valves in Human Heart. The Proceedings of the Computational Mechanics Conference, 2007, 2007.20, 139.	0.0	0
63	1P577 Efficient calculation of electrostatic interaction in biomolecular simulation revisited(27.) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Seibutsu Butsuri, 2006, 46, S291.	0.1	0
64	138 Parallelization Strategies for Fluid-Structure Interaction Analysis of a Heart. The Proceedings of the Computational Mechanics Conference, 2006, 2006.19, 333-334.	0.0	0
65	131 The effects of vessel wall properties of cerebral aneurysm on internal blood flow. The Proceedings of the Computational Mechanics Conference, 2006, 2006.19, 319-320.	0.0	0
66	133 Fluid Structure Interaction Analysis of Left and Right Ventricle and Coronary Circulation Based on Porohyperelastic Theory. The Proceedings of the Computational Mechanics Conference, 2006, 2006.19, 323-324.	0.0	0
67	A robust preconditioner for fluid-structure interaction problems. Computer Methods in Applied Mechanics and Engineering, 2005, 194, 4027-4047.	6.6	30
68	Error analysis for a potential problem on locally refined grids. Numerische Mathematik, 2000, 86, 539-563.	1.9	6
69	Fourier Analysis of GMRES(m) Preconditioned by Multigrid. SIAM Journal of Scientific Computing, 2000, 22, 582-603.	2.8	24
70	Ordering strategies and related techniques to overcome the trade-off between parallelism and convergence in incomplete factorizations. Parallel Computing, 1999, 25, 1995-2014.	2.1	39
71	Two-Stage Method for Protein-Ligand Docking. Journal of Medicinal Chemistry, 1999, 42, 4422-4433.	6.4	86
72	Multigrid Line Smoothers for Higher Order Upwind Discretizations of Convection-Dominated Problems. Journal of Computational Physics, 1998, 139, 274-307.	3.8	49

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73	An Evaluation of Parallel Multigrid as a Solver and a Preconditioner for Singularly Perturbed Problems. SIAM Journal of Scientific Computing, 1998, 19, 87-110.	2.8	49
74	Flexible Multiple Semicoarsening for Three-Dimensional Singularly Perturbed Problems. SIAM Journal of Scientific Computing, 1998, 19, 1646-1666.	2.8	31
75	Real applications on the new parallel system NEC Cenju-3. Parallel Computing, 1996, 22, 131-148.	2.1	7
76	Overlapped Multicolor MILU Preconditioning. SIAM Journal of Scientific Computing, 1995, 16, 636-650.	2.8	8
77	Parallel block preconditioning based on SSOR and MILU. Numerical Linear Algebra With Applications, 1994, 1, 533-553.	1.6	18