

Martin J Howard

List of Publications by Year in descending order

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83
papers

5,343
citations

87888

38
h-index

95266

68
g-index

94
all docs

94
docs citations

94
times ranked

4325
citing authors

#	ARTICLE	IF	CITATIONS
1	Investigating Histone Modification Dynamics by Mechanistic Computational Modeling. <i>Methods in Molecular Biology</i> , 2022, , 441-473.	0.9	2
2	Using computational modelling to reveal mechanisms of epigenetic Polycomb control. <i>Biochemical Society Transactions</i> , 2021, 49, 71-77.	3.4	14
3	Cell size controlled in plants using DNA content as an internal scale. <i>Science</i> , 2021, 372, 1176-1181.	12.6	70
4	Digital paradigm for Polycomb epigenetic switching and memory. <i>Current Opinion in Plant Biology</i> , 2021, 61, 102012.	7.1	15
5	Diffusion-mediated HEI10 coarsening can explain meiotic crossover positioning in <i>Arabidopsis</i> . <i>Nature Communications</i> , 2021, 12, 4674.	12.8	62
6	Hybrid protein assembly-histone modification mechanism for PRC2-based epigenetic switching and memory. <i>ELife</i> , 2021, 10, .	6.0	23
7	Feeling Every Bit of Winter “ Distributed Temperature Sensitivity in Vernalization. <i>Frontiers in Plant Science</i> , 2021, 12, 628726.	3.6	14
8	A cis-acting mechanism mediates transcriptional memory at Polycomb target genes in mammals. <i>Nature Genetics</i> , 2021, 53, 1686-1697.	21.4	53
9	Temperature-dependent growth contributes to long-term cold sensing. <i>Nature</i> , 2020, 583, 825-829.	27.8	77
10	A theoretical model of Polycomb/Trithorax action unites stable epigenetic memory and dynamic regulation. <i>Nature Communications</i> , 2020, 11, 4782.	12.8	24
11	The 3' processing of antisense RNAs physically links to chromatin-based transcriptional control. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 15316-15321.	7.1	40
12	Noncoding SNPs influence a distinct phase of Polycomb silencing to destabilize long-term epigenetic memory at <i>Arabidopsis FLC</i> . <i>Genes and Development</i> , 2020, 34, 446-461.	5.9	30
13	Natural variation in autumn expression is the major adaptive determinant distinguishing <i>Arabidopsis FLC</i> haplotypes. <i>ELife</i> , 2020, 9, .	6.0	28
14	Reassessment of the Basis of Cell Size Control Based on Analysis of Cell-to-Cell Variability. <i>Biophysical Journal</i> , 2019, 117, 1728-1738.	0.5	21
15	Center Finding in <i>E. coli</i> and the Role of Mathematical Modeling: Past, Present and Future. <i>Journal of Molecular Biology</i> , 2019, 431, 928-938.	4.2	7
16	Reprogramming Cdr2-Dependent Geometry-Based Cell Size Control in Fission Yeast. <i>Current Biology</i> , 2019, 29, 350-358.e4.	3.9	62
17	Absence of warmth permits epigenetic memory of winter in <i>Arabidopsis</i> . <i>Nature Communications</i> , 2018, 9, 639.	12.8	90
18	Temperature Sensing Is Distributed throughout the Regulatory Network that Controls FLC Epigenetic Silencing in Vernalization. <i>Cell Systems</i> , 2018, 7, 643-655.e9.	6.2	46

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19	A gated relaxation oscillator mediated by FrzX controls morphogenetic movements in <i>Myxococcus xanthus</i> . <i>Nature Microbiology</i> , 2018, 3, 948-959.	13.3	44
20	Dissecting chromatin-mediated gene regulation and epigenetic memory through mathematical modelling. <i>Current Opinion in Systems Biology</i> , 2017, 3, 7-14.	2.6	19
21	Cell-Size-Dependent Transcription of FLC and Its Antisense Long Non-coding RNA COOLAIR Explain Cell-to-Cell Expression Variation. <i>Cell Systems</i> , 2017, 4, 622-635.e9.	6.2	70
22	Slow Chromatin Dynamics Allow Polycomb Target Genes to Filter Fluctuations in Transcription Factor Activity. <i>Cell Systems</i> , 2017, 4, 445-457.e8.	6.2	99
23	Controlling cell size through sizer mechanisms. <i>Current Opinion in Systems Biology</i> , 2017, 5, 86-92.	2.6	74
24	Distinct phases of Polycomb silencing to hold epigenetic memory of cold in <i>Arabidopsis</i> . <i>Science</i> , 2017, 357, 1142-1145.	12.6	167
25	Disruption of an RNA-binding hinge region abolishes LHP1-mediated epigenetic repression. <i>Genes and Development</i> , 2017, 31, 2115-2120.	5.9	33
26	Quantitative Environmentally Triggered Switching Between Stable Epigenetic States. , 2017, , 169-187.		0
27	Physical coupling of activation and derepression activities to maintain an active transcriptional state at <i>FLC</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 9369-9374.	7.1	55
28	Quantitative regulation of <i>FLC</i> via coordinated transcriptional initiation and elongation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 218-223.	7.1	76
29	Local chromatin environment of a Polycomb target gene instructs its own epigenetic inheritance. <i>ELife</i> , 2015, 4, .	6.0	92
30	How plants manage food reserves at night: quantitative models and open questions. <i>Frontiers in Plant Science</i> , 2015, 6, 204.	3.6	35
31	Vernalizing cold is registered digitally at <i>FLC</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 4146-4151.	7.1	78
32	Competing ParA Structures Space Bacterial Plasmids Equally over the Nucleoid. <i>PLoS Computational Biology</i> , 2014, 10, e1004009.	3.2	60
33	Antagonistic Roles for H3K36me3 and H3K27me3 in the Cold-Induced Epigenetic Switch at <i>Arabidopsis FLC</i> . <i>Current Biology</i> , 2014, 24, 1793-1797.	3.9	201
34	Cortical regulation of cell size by a sizer <i>cdr2p</i> . <i>ELife</i> , 2014, 3, e02040.	6.0	111
35	Computational and Genetic Reduction of a Cell Cycle to Its Simplest, Primordial Components. <i>PLoS Biology</i> , 2013, 11, e1001749.	5.6	60
36	<i>Arabidopsis</i> plants perform arithmetic division to prevent starvation at night. <i>ELife</i> , 2013, 2, e00669.	6.0	134

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37	Mechanistic Basis of Branch-Site Selection in Filamentous Bacteria. <i>PLoS Computational Biology</i> , 2012, 8, e1002423.	3.2	41
38	Quantitative Dynamics of Telomere Bouquet Formation. <i>PLoS Computational Biology</i> , 2012, 8, e1002812.	3.2	37
39	Regulation of apical growth and hyphal branching in <i>Streptomyces</i> . <i>Current Opinion in Microbiology</i> , 2012, 15, 737-743.	5.1	92
40	Vernalization – a cold-induced epigenetic switch. <i>Journal of Cell Science</i> , 2012, 125, 3723-31.	2.0	193
41	Noise Reduction in the Intracellular Pom1p Gradient by a Dynamic Clustering Mechanism. <i>Developmental Cell</i> , 2012, 22, 558-572.	7.0	83
42	How to build a robust intracellular concentration gradient. <i>Trends in Cell Biology</i> , 2012, 22, 311-317.	7.9	48
43	A Polycomb-based switch underlying quantitative epigenetic memory. <i>Nature</i> , 2011, 476, 105-108.	27.8	414
44	What is the mechanism of ParA-mediated DNA movement?. <i>Molecular Microbiology</i> , 2010, 78, 9-12.	2.5	18
45	Shaping a Morphogen Gradient for Positional Precision. <i>Biophysical Journal</i> , 2010, 99, 697-707.	0.5	46
46	Pushing and Pulling in Prokaryotic DNA Segregation. <i>Cell</i> , 2010, 141, 927-942.	28.9	281
47	Morphogen profiles can be optimized to buffer against noise. <i>Physical Review E</i> , 2009, 80, 041902.	2.1	39
48	Role of Spatial Averaging in the Precision of Gene Expression Patterns. <i>Physical Review Letters</i> , 2009, 103, 258101.	7.8	70
49	A mechanical bottleneck explains the variation in cup growth during <i>FcγR</i> phagocytosis. <i>Molecular Systems Biology</i> , 2009, 5, 298.	7.2	44
50	When it pays to rush: interpreting morphogen gradients prior to steady-state. <i>Physical Biology</i> , 2009, 6, 046020.	1.8	31
51	Movement and equi-positioning of plasmids by ParA filament disassembly. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 19369-19374.	7.1	171
52	Cell Division: Experiments and Modelling Unite to Resolve the Middle. <i>Current Biology</i> , 2009, 19, R67-R69.	3.9	0
53	Modeling the Establishment of PAR Protein Polarity in the One-Cell <i>C. elegans</i> Embryo. <i>Biophysical Journal</i> , 2008, 95, 4512-4522.	0.5	39
54	Fundamental Limits to Position Determination by Concentration Gradients. <i>PLoS Computational Biology</i> , 2007, 3, e78.	3.2	111

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55	An experimentalist's guide to computational modelling of the Min system. <i>Molecular Microbiology</i> , 2007, 63, 1279-1284.	2.5	77
56	Cell Signalling: Changing Shape Changes the Signal. <i>Current Biology</i> , 2006, 16, R673-R675.	3.9	6
57	The Cell-End Factor Pom1p Inhibits Mid1p in Specification of the Cell Division Plane in Fission Yeast. <i>Current Biology</i> , 2006, 16, 2480-2487.	3.9	126
58	A stochastic model of Min oscillations in <i>Escherichia coli</i> and Min protein segregation during cell division. <i>Physical Biology</i> , 2006, 3, 1-12.	1.8	43
59	Modeling dual pathways for the metazoan spindle assembly checkpoint. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 16758-16763.	7.1	42
60	Applications of field-theoretic renormalization group methods to reaction-diffusion problems. <i>Journal of Physics A</i> , 2005, 38, R79-R131.	1.6	241
61	Finding the Center Reliably: Robust Patterns of Developmental Gene Expression. <i>Physical Review Letters</i> , 2005, 95, 208103.	7.8	68
62	Stochastic model for Soj relocation dynamics in <i>Bacillus subtilis</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 9808-9813.	7.1	29
63	Cellular organization by self-organization. <i>Journal of Cell Biology</i> , 2005, 168, 533-536.	5.2	83
64	Fundamental Limits to Position Determination by Concentration Gradients. <i>PLoS Computational Biology</i> , 2005, preprint, e78.	3.2	1
65	Dynamics and stability of vortex-antivortex fronts in type-II superconductors. <i>Physical Review E</i> , 2004, 70, 026209.	2.1	5
66	A Mechanism for Polar Protein Localization in Bacteria. <i>Journal of Molecular Biology</i> , 2004, 335, 655-663.	4.2	36
67	Pattern Formation inside Bacteria: Fluctuations due to the Low Copy Number of Proteins. <i>Physical Review Letters</i> , 2003, 90, 128102.	7.8	102
68	Hole-defect chaos in the one-dimensional complex Ginzburg-Landau equation. <i>Physical Review E</i> , 2003, 68, 026213.	2.1	11
69	DIRECTED PERCOLATION AND OTHER SYSTEMS WITH ABSORBING STATES: IMPACT OF BOUNDARIES. <i>International Journal of Modern Physics B</i> , 2001, 15, 1761-1797.	2.0	32
70	FLUCTUATIONS AND CORRELATIONS IN POPULATION MODELS WITH AGE STRUCTURE. <i>International Journal of Modern Physics B</i> , 2001, 15, 391-402.	2.0	5
71	Branching and annihilating Lévy flights. <i>Physical Review E</i> , 2001, 63, 041116.	2.1	31
72	Ordered and Self-Disordered Dynamics of Holes and Defects in the One-Dimensional Complex Ginzburg-Landau Equation. <i>Physical Review Letters</i> , 2001, 86, 2018-2021.	7.8	38

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73	Dynamic Compartmentalization of Bacteria: Accurate Division in <i>E. Coli</i> . <i>Physical Review Letters</i> , 2001, 87, 278102.	7.8	164
74	Surface critical behavior in systems with nonequilibrium phase transitions. <i>Physical Review E</i> , 2000, 61, 167-183.	2.1	13
75	Nonequilibrium critical behavior in unidirectionally coupled stochastic processes. <i>Physical Review E</i> , 1999, 59, 6381-6408.	2.1	32
76	Surface Critical Behavior in Systems with Absorbing States. <i>Physical Review Letters</i> , 1998, 81, 2104-2107.	7.8	20
77	Directed percolation with a wall or edge. <i>Journal of Physics A</i> , 1998, 31, 2311-2320.	1.6	21
78	Persistence in the Voter model: continuum reaction-diffusion approach. <i>Journal of Physics A</i> , 1998, 31, L209-L215.	1.6	27
79	Multicritical Behavior in Coupled Directed Percolation Processes. <i>Physical Review Letters</i> , 1998, 80, 2165-2168.	7.8	43
80	TÄuber, Howard, and Hinrichsen Reply:. <i>Physical Review Letters</i> , 1998, 81, 2179-2179.	7.8	2
81	'Real' versus 'imaginary' noise in diffusion-limited reactions. <i>Journal of Physics A</i> , 1997, 30, 7721-7731.	1.6	101
82	Fluctuation kinetics in a multispecies reaction - diffusion system. <i>Journal of Physics A</i> , 1996, 29, 3437-3460.	1.6	36
83	Fluctuation effects and multiscaling of the reaction-diffusion front for $A+B \rightarrow OE$. <i>Journal of Physics A</i> , 1995, 28, 3599-3621.	1.6	51