

Hans-Peter Herzel

List of Publications by Year in descending order

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95
papers

6,366
citations

76326

40
h-index

74163

75
g-index

107
all docs

107
docs citations

107
times ranked

5596
citing authors

#	ARTICLE	IF	CITATIONS
1	Spontaneous Synchronization of Coupled Circadian Oscillators. <i>Biophysical Journal</i> , 2005, 89, 120-129.	0.5	401
2	Differential effects of PER2 phosphorylation: molecular basis for the human familial advanced sleep phase syndrome (FASPS). <i>Genes and Development</i> , 2006, 20, 2660-2672.	5.9	339
3	Lymphocyte Circadian Clocks Control Lymph Node Trafficking and Adaptive Immune Responses. <i>Immunity</i> , 2017, 46, 120-132.	14.3	324
4	Coupling governs entrainment range of circadian clocks. <i>Molecular Systems Biology</i> , 2010, 6, 438.	7.2	297
5	Bifurcations in an asymmetric vocal fold model. <i>Journal of the Acoustical Society of America</i> , 1995, 97, 1874-1884.	1.1	286
6	Regulation of Clock-Controlled Genes in Mammals. <i>PLoS ONE</i> , 2009, 4, e4882.	2.5	251
7	Guidelines for Genome-Scale Analysis of Biological Rhythms. <i>Journal of Biological Rhythms</i> , 2017, 32, 380-393.	2.6	237
8	Mathematical Modeling Identifies Inhibitors of Apoptosis as Mediators of Positive Feedback and Bistability. <i>PLoS Computational Biology</i> , 2006, 2, e120.	3.2	217
9	Synchronization-Induced Rhythmicity of Circadian Oscillators in the Suprachiasmatic Nucleus. <i>PLoS Computational Biology</i> , 2007, 3, e68.	3.2	184
10	Tuning the Mammalian Circadian Clock: Robust Synergy of Two Loops. <i>PLoS Computational Biology</i> , 2011, 7, e1002309.	3.2	179
11	High-accuracy determination of internal circadian time from a single blood sample. <i>Journal of Clinical Investigation</i> , 2018, 128, 3826-3839.	8.2	174
12	Modeling Feedback Loops of the Mammalian Circadian Oscillator. <i>Biophysical Journal</i> , 2004, 87, 3023-3034.	0.5	151
13	Effects of sequestration on signal transduction cascades. <i>FEBS Journal</i> , 2006, 273, 895-906.	4.7	148
14	Ras-Mediated Deregulation of the Circadian Clock in Cancer. <i>PLoS Genetics</i> , 2014, 10, e1004338.	3.5	140
15	Bifurcation analysis of the regulatory modules of the mammalian G1/S transition. <i>Bioinformatics</i> , 2004, 20, 1506-1511.	4.1	125
16	Species independence of mutual information in coding and noncoding DNA. <i>Physical Review E</i> , 2000, 61, 5624-5629.	2.1	120
17	The choroid plexus is an important circadian clock component. <i>Nature Communications</i> , 2018, 9, 1062.	12.8	118
18	Spatio-temporal analysis of irregular vocal fold oscillations: Biphonation due to desynchronization of spatial modes. <i>Journal of the Acoustical Society of America</i> , 2001, 110, 3179-3192.	1.1	114

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19	Correlations in DNA sequences: The role of protein coding segments. <i>Physical Review E</i> , 1997, 55, 800-810.	2.1	98
20	Timing of circadian genes in mammalian tissues. <i>Scientific Reports</i> , 2014, 4, 5782.	3.3	97
21	Human Chronotypes from a Theoretical Perspective. <i>PLoS ONE</i> , 2013, 8, e59464.	2.5	92
22	Quantification of Circadian Rhythms in Single Cells. <i>PLoS Computational Biology</i> , 2009, 5, e1000580.	3.2	88
23	Tuning the phase of circadian entrainment. <i>Journal of the Royal Society Interface</i> , 2015, 12, 20150282.	3.4	85
24	Modeling the role of nonhuman vocal membranes in phonation. <i>Journal of the Acoustical Society of America</i> , 1999, 105, 2020-2028.	1.1	83
25	Modelling transcriptional feedback loops: the role of Gro/TLE1 in Hes1 oscillations. <i>Philosophical Transactions Series A, Mathematical, Physical, and Engineering Sciences</i> , 2006, 364, 1155-1170.	3.4	83
26	Functioning and robustness of a bacterial circadian clock. <i>Molecular Systems Biology</i> , 2007, 3, 90.	7.2	83
27	Global parameter search reveals design principles of the mammalian circadian clock. <i>BMC Systems Biology</i> , 2008, 2, 22.	3.0	82
28	Competing Docking Interactions can Bring About Bistability in the MAPK Cascade. <i>Biophysical Journal</i> , 2007, 93, 2279-2288.	0.5	78
29	Mechanism for 12 Hr Rhythm Generation by the Circadian Clock. <i>Cell Reports</i> , 2013, 3, 1228-1238.	6.4	78
30	Feedback Loops of the Mammalian Circadian Clock Constitute Repressilator. <i>PLoS Computational Biology</i> , 2016, 12, e1005266.	3.2	75
31	Positive Feedback Promotes Oscillations in Negative Feedback Loops. <i>PLoS ONE</i> , 2014, 9, e104761.	2.5	74
32	The Interplay of cis-Regulatory Elements Rules Circadian Rhythms in Mouse Liver. <i>PLoS ONE</i> , 2012, 7, e46835.	2.5	68
33	Correlations in Protein Sequences and Property Codes. <i>Journal of Theoretical Biology</i> , 1998, 190, 341-353.	1.7	60
34	High reproducibility of large-gel two-dimensional electrophoresis. <i>Electrophoresis</i> , 2004, 25, 3040-3047.	2.4	56
35	Co-existing feedback loops generate tissue-specific circadian rhythms. <i>Life Science Alliance</i> , 2018, 1, e201800078.	2.8	55
36	Statistical analysis of the DNA sequence of human chromosome 22. <i>Physical Review E</i> , 2001, 64, 041917.	2.1	53

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37	Identification of Novel Nuclear Factor of Activated T Cell (NFAT)-associated Proteins in T Cells. <i>Journal of Biological Chemistry</i> , 2016, 291, 24172-24187.	3.4	51
38	A Theoretical Study on Seasonality. <i>Frontiers in Neurology</i> , 2015, 6, 94.	2.4	50
39	Quantitative analysis of ultrasensitive responses. <i>FEBS Journal</i> , 2005, 272, 4071-4079.	4.7	49
40	Periodicities of 10â€“11bp as Indicators of the Supercoiled State of Genomic DNA. <i>Journal of Molecular Biology</i> , 2004, 343, 891-901.	4.2	47
41	Assembly of a Comprehensive Regulatory Network for the Mammalian Circadian Clock: A Bioinformatics Approach. <i>PLoS ONE</i> , 2015, 10, e0126283.	2.5	43
42	Measuring Relative Coupling Strength in Circadian Systems. <i>Journal of Biological Rhythms</i> , 2018, 33, 84-98.	2.6	43
43	Phonation onset: Vocal fold modeling and high-speed glottography. <i>Journal of the Acoustical Society of America</i> , 1998, 104, 464-470.	1.1	42
44	Live-cell imaging of circadian clock protein dynamics in CRISPR-generated knock-in cells. <i>Nature Communications</i> , 2021, 12, 3796.	12.8	42
45	Venn diagram analysis overestimates the extent of circadian rhythm reprogramming. <i>FEBS Journal</i> , 2022, 289, 6605-6621.	4.7	40
46	Timing of Neuropeptide Coupling Determines Synchrony and Entrainment in the Mammalian Circadian Clock. <i>PLoS Computational Biology</i> , 2014, 10, e1003565.	3.2	38
47	Sequence Periodicity in Complete Genomes of Archaea Suggests Positive Supercoiling. <i>Journal of Biomolecular Structure and Dynamics</i> , 1998, 16, 341-345.	3.5	37
48	Excitability in the p53 network mediates robust signaling with tunable activation thresholds in single cells. <i>Scientific Reports</i> , 2017, 7, 46571.	3.3	37
49	Intercellular coupling between peripheral circadian oscillators by TGF- β^2 signaling. <i>Science Advances</i> , 2021, 7, .	10.3	37
50	Clocks in the Wild: Entrainment to Natural Light. <i>Frontiers in Physiology</i> , 2020, 11, 272.	2.8	33
51	Amplitude Effects Allow Short Jet Lags and Large Seasonal Phase Shifts in Minimal Clock Models. <i>Journal of Molecular Biology</i> , 2020, 432, 3722-3737.	4.2	31
52	Moranâ€™s I quantifies spatio-temporal pattern formation in neural imaging data. <i>Bioinformatics</i> , 2017, 33, 3072-3079.	4.1	30
53	Ultrasensitization: Switch-Like Regulation of Cellular Signaling by Transcriptional Induction. <i>PLoS Computational Biology</i> , 2005, 1, e54.	3.2	28
54	Regulation of mammalian cell cycle progression in the regenerating liver. <i>Journal of Theoretical Biology</i> , 2011, 283, 103-112.	1.7	28

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55	Robustness: A Key to Evolutionary Design. <i>Biological Theory</i> , 2006, 1, 90-93.	1.5	26
56	Extracting information from cDNA arrays. <i>Chaos</i> , 2001, 11, 98.	2.5	23
57	Coupling Controls the Synchrony of Clock Cells in Development and Knockouts. <i>Biophysical Journal</i> , 2015, 109, 2159-2170.	0.5	22
58	Circadian rhythms in septic shock patients. <i>Annals of Intensive Care</i> , 2021, 11, 64.	4.6	22
59	Gating Characteristics Control Glutamate Receptor Distribution and Trafficking In Vivo. <i>Current Biology</i> , 2014, 24, 2059-2065.	3.9	20
60	Are Noncoding Sequences of <i>Rickettsia prowazekii</i> Remnants of "Neutralized" Genes?. <i>Journal of Molecular Evolution</i> , 2000, 51, 353-362.	1.8	19
61	Weak coupling between intracellular feedback loops explains dissociation of clock gene dynamics. <i>PLoS Computational Biology</i> , 2019, 15, e1007330.	3.2	19
62	A Robust Model for Circadian Redox Oscillations. <i>International Journal of Molecular Sciences</i> , 2019, 20, 2368.	4.1	18
63	Ultradian Rhythms in the Transcriptome of <i>Neurospora crassa</i> . <i>IScience</i> , 2018, 9, 475-486.	4.1	15
64	An Inactivation Switch Enables Rhythms in a <i>Neurospora</i> Clock Model. <i>International Journal of Molecular Sciences</i> , 2019, 20, 2985.	4.1	15
65	Conceptual Models of Entrainment, Jet Lag, and Seasonality. <i>Frontiers in Physiology</i> , 2020, 11, 334.	2.8	15
66	How to Quantify 'Small-World Networks'?. <i>Fractals</i> , 1998, 06, 301-303.	3.7	14
67	Transcription factor co-occupied regions in the murine genome constitute T helper cell subtype-specific enhancers. <i>European Journal of Immunology</i> , 2015, 45, 3150-3157.	2.9	13
68	Coherency of circadian rhythms in the SCN is governed by the interplay of two coupling factors. <i>PLoS Computational Biology</i> , 2018, 14, e1006607.	3.2	13
69	Mining for novel candidate clock genes in the circadian regulatory network. <i>BMC Systems Biology</i> , 2015, 9, 78.	3.0	12
70	Phasegram Analysis of Vocal Fold Vibration Documented With Laryngeal High-speed Video Endoscopy. <i>Journal of Voice</i> , 2016, 30, 771.e1-771.e15.	1.5	12
71	Nonlinear phenomena in models of the circadian clock. <i>Journal of the Royal Society Interface</i> , 2020, 17, 20200556.	3.4	12
72	AVERAGE MUTUAL INFORMATION OF CODING AND NONCODING DNA. , 1999, , 614-23.		12

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73	Principles underlying the complex dynamics of temperature entrainment by a circadian clock. <i>IScience</i> , 2021, 24, 103370.	4.1	12
74	The structural code of cyanobacterial genomes. <i>Nucleic Acids Research</i> , 2014, 42, 8873-8883.	14.5	11
75	Searching Novel Clock Genes Using RNAi-Based Screening. <i>Methods in Molecular Biology</i> , 2021, 2130, 103-114.	0.9	11
76	Quantitative analysis of circadian single cell oscillations in response to temperature. <i>PLoS ONE</i> , 2018, 13, e0190004.	2.5	11
77	Adequate immune response ensured by binary IL-2 and graded CD25 expression in a murine transfer model. <i>ELife</i> , 2016, 5, .	6.0	11
78	A mesoscale model of G1/S phase transition in liver regeneration. <i>Journal of Theoretical Biology</i> , 2008, 252, 465-473.	1.7	10
79	Stable IL-2 Decision Making by Endogenous c-Fos Amounts in Peripheral Memory T-helper Cells. <i>Journal of Biological Chemistry</i> , 2012, 287, 18386-18397.	3.4	10
80	Elucidating the adaptation and temporal coordination of metabolic pathways using in-silico evolution. <i>BioSystems</i> , 2014, 117, 68-76.	2.0	10
81	Multiple random phosphorylations in clock proteins provide long delays and switches. <i>Scientific Reports</i> , 2020, 10, 22224.	3.3	9
82	SPOTTED HYAENA WHOOPS: FREQUENT INCIDENCE OF VOCAL INSTABILITIES IN A MAMMALIAN LOUD CALL. <i>Bioacoustics</i> , 2004, 14, 99-109.	1.7	8
83	Synergies of Multiple Zeitgebers Tune Entrainment. <i>Frontiers in Network Physiology</i> , 2022, 1, .	1.8	7
84	Beyond spikes: Multiscale computational analysis of <i>in vivo</i> long-term recordings in the cockroach circadian clock. <i>Network Neuroscience</i> , 2019, 3, 944-968.	2.6	6
85	Mathematical modelling identifies conditions for maintaining and escaping feedback control in the intestinal epithelium. <i>Scientific Reports</i> , 2022, 12, 5569.	3.3	6
86	GRAPH-THEORETICAL COMPARISON REVEALS STRUCTURAL DIVERGENCE OF HUMAN PROTEIN INTERACTION NETWORKS. , 2007, , .		5
87	Death of neuronal clusters contributes to variance of age at onset in Huntington's disease. <i>Neurogenetics</i> , 2006, 7, 21-25.	1.4	4
88	Mathematical Modeling in Circadian Rhythmicity. <i>Methods in Molecular Biology</i> , 2022, , 55-80.	0.9	4
89	Flexible web-based integration of distributed large-scale human protein interaction maps. <i>Journal of Integrative Bioinformatics</i> , 2007, 4, 40-50.	1.5	3
90	Functional and Transcriptional Coherency of Modules in the Human Protein Interaction Network. <i>Journal of Integrative Bioinformatics</i> , 2007, 4, 198-207.	1.5	3

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91	Neither <i>per</i> , nor <i>tim1</i> , nor <i>cry2</i> alone are essential components of the molecular circadian clockwork in the Madeira cockroach. PLoS ONE, 2020, 15, e0235930.	2.5	3
92	PREDICTION OF REGULATORY TRANSCRIPTION FACTORS IN T HELPER CELL DIFFERENTIATION AND MAINTENANCE. , 2010, , .		2
93	MODELING IL-2 GENE EXPRESSION IN HUMAN REGULATORY T CELLS. , 2008, , .		1
94	Simple Kinetic Models in Molecular Chronobiology. Methods in Molecular Biology, 2021, 2130, 87-100.	0.9	1
95	Modeling IL-2 gene expression in human regulatory T cells. Genome Informatics, 2008, 20, 222-30.	0.4	1