

# John B Hogenesch

## List of Publications by Year in descending order

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143  
papers

28,378  
citations

13099

68  
h-index

10734

138  
g-index

158  
all docs

158  
docs citations

158  
times ranked

27508  
citing authors

#	ARTICLE	IF	CITATIONS
1	Coordinated Transcription of Key Pathways in the Mouse by the Circadian Clock. <i>Cell</i> , 2002, 109, 307-320.	28.9	2,099
2	A circadian gene expression atlas in mammals: Implications for biology and medicine. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 16219-16224.	7.1	1,802
3	Mop3 Is an Essential Component of the Master Circadian Pacemaker in Mammals. <i>Cell</i> , 2000, 103, 1009-1017.	28.9	1,380
4	Large-scale analysis of the human and mouse transcriptomes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 4465-4470.	7.1	1,366
5	The PAS Superfamily: Sensors of Environmental and Developmental Signals. <i>Annual Review of Pharmacology and Toxicology</i> , 2000, 40, 519-561.	9.4	959
6	JTK_CYCLE: An Efficient Nonparametric Algorithm for Detecting Rhythmic Components in Genome-Scale Data Sets. <i>Journal of Biological Rhythms</i> , 2010, 25, 372-380.	2.6	919
7	A Functional Genomics Strategy Reveals Rora as a Component of the Mammalian Circadian Clock. <i>Neuron</i> , 2004, 43, 527-537.	8.1	909
8	Circadian rhythms from flies to human. <i>Nature</i> , 2002, 417, 329-335.	27.8	860
9	BMAL1 and CLOCK, Two Essential Components of the Circadian Clock, Are Involved in Glucose Homeostasis. <i>PLoS Biology</i> , 2004, 2, e377.	5.6	860
10	A Heat-Sensitive TRP Channel Expressed in Keratinocytes. <i>Science</i> , 2002, 296, 2046-2049.	12.6	828
11	Melanopsin ( <i>Opn4</i> ) Requirement for Normal Light-Induced Circadian Phase Shifting. <i>Science</i> , 2002, 298, 2213-2216.	12.6	768
12	Melanopsin Is Required for Non-Image-Forming Photoc Responses in Blind Mice. <i>Science</i> , 2003, 301, 525-527.	12.6	635
13	Harmonics of Circadian Gene Transcription in Mammals. <i>PLoS Genetics</i> , 2009, 5, e1000442.	3.5	616
14	TORCs. <i>Molecular Cell</i> , 2003, 12, 413-423.	9.7	564
15	ESRP1 and ESRP2 Are Epithelial Cell-Type-Specific Regulators of FGFR2 Splicing. <i>Molecular Cell</i> , 2009, 33, 591-601.	9.7	509
16	LXR-Dependent Gene Expression Is Important for Macrophage Survival and the Innate Immune Response. <i>Cell</i> , 2004, 119, 299-309.	28.9	498
17	Circadian and CLOCK-controlled regulation of the mouse transcriptome and cell proliferation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 3342-3347.	7.1	439
18	A Genome-wide RNAi Screen for Modifiers of the Circadian Clock in Human Cells. <i>Cell</i> , 2009, 139, 199-210.	28.9	437

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19	Characterization of a Subset of the Basic-Helix-Loop-Helix-PAS Superfamily That Interacts with Components of the Dioxin Signaling Pathway. <i>Journal of Biological Chemistry</i> , 1997, 272, 8581-8593.	3.4	425
20	Illumination of the Melanopsin Signaling Pathway. <i>Science</i> , 2005, 307, 600-604.	12.6	421
21	MetaCycle: an integrated R package to evaluate periodicity in large scale data. <i>Bioinformatics</i> , 2016, 32, 3351-3353.	4.1	413
22	A new view of transcriptome complexity and regulation through the lens of local splicing variations. <i>ELife</i> , 2016, 5, e11752.	6.0	385
23	A Chemical, Genetic, and Structural Analysis of the Nuclear Bile Acid Receptor FXR. <i>Molecular Cell</i> , 2003, 11, 1079-1092.	9.7	359
24	Feedback repression is required for mammalian circadian clock function. <i>Nature Genetics</i> , 2006, 38, 312-319.	21.4	344
25	Extensive Variation in Chromatin States Across Humans. <i>Science</i> , 2013, 342, 750-752.	12.6	338
26	Genome-Wide Expression Analysis in <i>Drosophila</i> Reveals Genes Controlling Circadian Behavior. <i>Journal of Neuroscience</i> , 2002, 22, 9305-9319.	3.6	329
27	Mammalian Per-Arnt-Sim Proteins in Environmental Adaptation. <i>Annual Review of Physiology</i> , 2010, 72, 625-645.	13.1	321
28	Identification of the circadian transcriptome in adult mouse skeletal muscle. <i>Physiological Genomics</i> , 2007, 31, 86-95.	2.3	300
29	CLOCK and BMAL1 regulate <i>MyoD</i> and are necessary for maintenance of skeletal muscle phenotype and function. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 19090-19095.	7.1	299
30	CircaDB: a database of mammalian circadian gene expression profiles. <i>Nucleic Acids Research</i> , 2012, 41, D1009-D1013.	14.5	285
31	c-Myb and p300 Regulate Hematopoietic Stem Cell Proliferation and Differentiation. <i>Developmental Cell</i> , 2005, 8, 153-166.	7.0	251
32	Medicine in the Fourth Dimension. <i>Cell Metabolism</i> , 2019, 30, 238-250.	16.2	245
33	Guidelines for Genome-Scale Analysis of Biological Rhythms. <i>Journal of Biological Rhythms</i> , 2017, 32, 380-393.	2.6	237
34	Genome-Wide Analysis of CREB Target Genes Reveals A Core Promoter Requirement for cAMP Responsiveness. <i>Molecular Cell</i> , 2003, 11, 1101-1108.	9.7	232
35	Network Features of the Mammalian Circadian Clock. <i>PLoS Biology</i> , 2009, 7, e1000052.	5.6	228
36	MYC Disrupts the Circadian Clock and Metabolism in Cancer Cells. <i>Cell Metabolism</i> , 2015, 22, 1009-1019.	16.2	217

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37	A database of tissue-specific rhythmically expressed human genes has potential applications in circadian medicine. <i>Science Translational Medicine</i> , 2018, 10, .	12.4	217
38	The imprinted gene <i>Magel2</i> regulates normal circadian output. <i>Nature Genetics</i> , 2007, 39, 1266-1272.	21.4	196
39	The <i>Nephila clavipes</i> genome highlights the diversity of spider silk genes and their complex expression. <i>Nature Genetics</i> , 2017, 49, 895-903.	21.4	190
40	A Comparison of the Celera and Ensembl Predicted Gene Sets Reveals Little Overlap in Novel Genes. <i>Cell</i> , 2001, 106, 413-415.	28.9	185
41	CYCLOPS reveals human transcriptional rhythms in health and disease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 5312-5317.	7.1	184
42	Tissue specific expression of the rat Ah-receptor and ARNT mRNAs. <i>Nucleic Acids Research</i> , 1994, 22, 3038-3044.	14.5	162
43	Clock Regulation of Metabolites Reveals Coupling between Transcription and Metabolism. <i>Cell Metabolism</i> , 2017, 25, 961-974.e4.	16.2	162
44	Dosing time matters. <i>Science</i> , 2019, 365, 547-549.	12.6	161
45	Modeling of RNA-seq fragment sequence bias reduces systematic errors in transcript abundance estimation. <i>Nature Biotechnology</i> , 2016, 34, 1287-1291.	17.5	159
46	mTOR signaling regulates central and peripheral circadian clock function. <i>PLoS Genetics</i> , 2018, 14, e1007369.	3.5	154
47	The Circadian Clock Interacts with Metabolic Physiology to Influence Reproductive Fitness. <i>Cell Metabolism</i> , 2011, 13, 639-654.	16.2	149
48	Bioinformatic Analysis of Circadian Gene Oscillation in Mouse Aorta. <i>Circulation</i> , 2005, 112, 2716-2724.	1.6	141
49	Categorically Distinct Acute Stressors Elicit Dissimilar Transcriptional Profiles in the Paraventricular Nucleus of the Hypothalamus. <i>Journal of Neuroscience</i> , 2003, 23, 5607-5616.	3.6	136
50	IVT-seq reveals extreme bias in RNA sequencing. <i>Genome Biology</i> , 2014, 15, R86.	9.6	134
51	Night/Day Changes in Pineal Expression of >600 Genes. <i>Journal of Biological Chemistry</i> , 2009, 284, 7606-7622.	3.4	130
52	miR-210 Inhibits Trophoblast Invasion and Is a Serum Biomarker for Preeclampsia. <i>American Journal of Pathology</i> , 2013, 183, 1437-1445.	3.8	126
53	Understanding systems-level properties: timely stories from the study of clocks. <i>Nature Reviews Genetics</i> , 2011, 12, 407-416.	16.3	124
54	Cell Type-Specific Functions of Period Genes Revealed by Novel Adipocyte and Hepatocyte Circadian Clock Models. <i>PLoS Genetics</i> , 2014, 10, e1004244.	3.5	119

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55	Genome-scale functional profiling of the mammalian AP-1 signaling pathway. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 12153-12158.	7.1	115
56	Machine Learning Helps Identify CHRONO as a Circadian Clock Component. PLoS Biology, 2014, 12, e1001840.	5.6	109
57	The Basic Helix-Loop-Helix-PAS Protein MOP9 Is a Brain-Specific Heterodimeric Partner of Circadian and Hypoxia Factors. Journal of Neuroscience, 2000, 20, RC83-RC83.	3.6	104
58	A Novel <i>BHLHE41</i> Variant is Associated with Short Sleep and Resistance to Sleep Deprivation in Humans. Sleep, 2014, 37, 1327-1336.	1.1	104
59	A coactivator trap identifies NONO (p54 <sup>nrb</sup> ) as a component of the cAMP-signaling pathway. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 20314-20319.	7.1	103
60	The Transcription Factor Encyclopedia. Genome Biology, 2012, 13, R24.	9.6	103
61	It's All in the Timing: Many Clocks, Many Outputs. Journal of Biological Rhythms, 2004, 19, 374-387.	2.6	102
62	Ribosome profiling reveals an important role for translational control in circadian gene expression. Genome Research, 2015, 25, 1836-1847.	5.5	99
63	Brain-Specific Rescue of Clock Reveals System-Driven Transcriptional Rhythms in Peripheral Tissue. PLoS Genetics, 2012, 8, e1002835.	3.5	97
64	Population-level rhythms in human skin with implications for circadian medicine. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 12313-12318.	7.1	97
65	Benchmark analysis of algorithms for determining and quantifying full-length mRNA splice forms from RNA-seq data. Bioinformatics, 2015, 31, 3938-3945.	4.1	90
66	The Role of Clock Genes in Pharmacology. Annual Review of Pharmacology and Toxicology, 2010, 50, 187-214.	9.4	88
67	A Genome-wide Screen Identifies PAPP-AA-Mediated IGFR Signaling as a Novel Regulator of Habituation Learning. Neuron, 2015, 85, 1200-1211.	8.1	85
68	Assessing the prevalence of mycoplasma contamination in cell culture via a survey of NCBI's RNA-seq archive. Nucleic Acids Research, 2015, 43, 2535-2542.	14.5	80
69	Role for <i>LSM</i> genes in the regulation of circadian rhythms. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 15166-15171.	7.1	76
70	The Transcriptional Repressor STRA13 Regulates a Subset of Peripheral Circadian Outputs. Journal of Biological Chemistry, 2004, 279, 1141-1150.	3.4	75
71	Applications of a Rat Multiple Tissue Gene Expression Data Set. Genome Research, 2004, 14, 742-749.	5.5	73
72	Analysis and synthesis of high-amplitude Cis-elements in the mammalian circadian clock. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 14946-14951.	7.1	69

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73	Considerations for RNA-seq Analysis of Circadian Rhythms. <i>Methods in Enzymology</i> , 2015, 551, 349-367.	1.0	68
74	Placental Expression of miR-517a/b and miR-517c Contributes to Trophoblast Dysfunction and Preeclampsia. <i>PLoS ONE</i> , 2015, 10, e0122707.	2.5	67
75	Cisplatin-DNA adduct repair of transcribed genes is controlled by two circadian programs in mouse tissues. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E4777-E4785.	7.1	65
76	Discovering Biology in Periodic Data through Phase Set Enrichment Analysis (PSEA). <i>Journal of Biological Rhythms</i> , 2016, 31, 244-257.	2.6	63
77	The $\hat{3}$ -Secretase Cleavage Product of Polycystin-1 Regulates TCF and CHOP-Mediated Transcriptional Activation through a p300-Dependent Mechanism. <i>Developmental Cell</i> , 2012, 22, 197-210.	7.0	61
78	Neural clocks and Neuropeptide F/Y regulate circadian gene expression in a peripheral metabolic tissue. <i>ELife</i> , 2016, 5, .	6.0	61
79	An array of insights: application of DNA chip technology in the study of cell biology—Trends in Cell Biology, 2003, 13, 151-156.	7.9	56
80	Intracellular and intercellular processes determine robustness of the circadian clock. <i>FEBS Letters</i> , 2011, 585, 1427-1434.	2.8	54
81	Wnt ligands signal in a cooperative manner to promote foregut organogenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 15348-15353.	7.1	54
82	Adaptive Thermogenesis in Mice Is Enhanced by Opsin 3-Dependent Adipocyte Light Sensing. <i>Cell Reports</i> , 2020, 30, 672-686.e8.	6.4	53
83	Pax3 and Hippo Signaling Coordinate Melanocyte Gene Expression in Neural Crest. <i>Cell Reports</i> , 2014, 9, 1885-1895.	6.4	49
84	Computational and experimental insights into the circadian effects of SIRT1. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 11643-11648.	7.1	49
85	Identification of novel mammalian growth regulatory factors by genome-scale quantitative image analysis. <i>Genome Research</i> , 2005, 15, 1136-1144.	5.5	45
86	WAVECLOCK: wavelet analysis of circadian oscillation. <i>Bioinformatics</i> , 2008, 24, 2794-2795.	4.1	43
87	The Circadian Clock Gene, <i>Bmal1</i> , Regulates Intestinal Stem Cell Signaling and Represses Tumor Initiation. <i>Cellular and Molecular Gastroenterology and Hepatology</i> , 2021, 12, 1847-1872.e0.	4.5	43
88	Interrogation of nonconserved human adipose lincRNAs identifies a regulatory role of <i>linc-ADAL</i> in adipocyte metabolism. <i>Science Translational Medicine</i> , 2018, 10, .	12.4	42
89	Genome-wide effect of pulmonary airway epithelial cell-specific <i>Bmal1</i> deletion. <i>FASEB Journal</i> , 2019, 33, 6226-6238.	0.5	40
90	NF- $\hat{B}$ modifies the mammalian circadian clock through interaction with the core clock protein <i>BMAL1</i> . <i>PLoS Genetics</i> , 2021, 17, e1009933.	3.5	39

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91	Genomics and systems approaches in the mammalian circadian clock. <i>Current Opinion in Genetics and Development</i> , 2010, 20, 581-587.	3.3	38
92	Shift Work Disrupts Circadian Regulation of the Transcriptome in Hospital Nurses. <i>Journal of Biological Rhythms</i> , 2019, 34, 167-177.	2.6	38
93	KPNB1 mediates PER/CRY nuclear translocation and circadian clock function. <i>ELife</i> , 2015, 4, .	6.0	37
94	Identification and Characterization of Genes Susceptible to Transcriptional Cross-Talk between the Hypoxia and Dioxin Signaling Cascades. <i>Chemical Research in Toxicology</i> , 2006, 19, 1284-1293.	3.3	35
95	Ubiquitin ligase Siah2 regulates RevErb1± degradation and the mammalian circadian clock. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 12420-12425.	7.1	34
96	A population-based gene expression signature of molecular clock phase from a single epidermal sample. <i>Genome Medicine</i> , 2020, 12, 73.	8.2	34
97	Adhesion Regulates MAP Kinase/Ternary Complex Factor Exchange to Control a Proliferative Transcriptional Switch. <i>Current Biology</i> , 2012, 22, 2017-2026.	3.9	32
98	DNA Arrays: Applications and Implications for Circadian Biology. <i>Journal of Biological Rhythms</i> , 2003, 18, 96-105.	2.6	31
99	The Liver Clock Controls Cholesterol Homeostasis through Trib1 Protein-mediated Regulation of PCSK9/Low Density Lipoprotein Receptor (LDLR) Axis. <i>Journal of Biological Chemistry</i> , 2015, 290, 31003-31012.	3.4	31
100	CRY1±CBS binding regulates circadian clock function and metabolism. <i>FEBS Journal</i> , 2021, 288, 614-639.	4.7	29
101	Lowering Nighttime Blood Pressure With Bedtime Dosing of Antihypertensive Medications: Controversies in Hypertension - Con Side of the Argument. <i>Hypertension</i> , 2021, 78, 871-878.	2.7	26
102	Generation of a Novel Allelic Series of Cryptochrome Mutants via Mutagenesis Reveals Residues Involved in Protein-Protein Interaction and CRY2-Specific Repression. <i>Molecular and Cellular Biology</i> , 2009, 29, 5465-5476.	2.3	25
103	Polycystin-1 regulates bone development through an interaction with the transcriptional coactivator TAZ. <i>Human Molecular Genetics</i> , 2019, 28, 16-30.	2.9	25
104	The Local Edge Machine: inference of dynamic models of gene regulation. <i>Genome Biology</i> , 2016, 17, 214.	8.8	24
105	Ontogeny and function of the circadian clock in intestinal organoids. <i>EMBO Journal</i> , 2022, 41, e106973.	7.8	24
106	Circadian Dysregulation: The Next Frontier in Obstructive Sleep Apnea Research. <i>Otolaryngology - Head and Neck Surgery</i> , 2018, 159, 948-955.	1.9	23
107	The NRON complex controls circadian clock function through regulated PER and CRY nuclear translocation. <i>Scientific Reports</i> , 2019, 9, 11883.	3.3	23
108	RNA Profiling in Circadian Biology. <i>Methods in Enzymology</i> , 2005, 393, 366-376.	1.0	22

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109	The Growth and Impact of Alzheimer Disease Centers as Measured by Social Network Analysis. <i>JAMA Neurology</i> , 2014, 71, 412.	9.0	22
110	Short-term exposure to intermittent hypoxia leads to changes in gene expression seen in chronic pulmonary disease. <i>ELife</i> , 2021, 10, .	6.0	22
111	A functional map of NF $\kappa$ B signaling identifies novel modulators and multiple system controls. <i>Genome Biology</i> , 2007, 8, R104.	9.6	20
112	Network Dynamics to Evaluate Performance of an Academic Institution. <i>Science Translational Medicine</i> , 2010, 2, 53ps49.	12.4	20
113	A large-scale study reveals 24-h operational rhythms in hospital treatment. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 20953-20958.	7.1	20
114	Systematic Analysis of Mouse Genome Reveals Distinct Evolutionary and Functional Properties Among Circadian and Ultradian Genes. <i>Frontiers in Physiology</i> , 2018, 9, 1178.	2.8	19
115	Zebrafish <i>foxc1a</i> drives appendage-specific neural circuit development. <i>Development (Cambridge)</i> , 2015, 142, 753-762.	2.5	16
116	Experimental and statistical reevaluation provides no evidence for <i>Drosophila</i> courtship song rhythms. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 9978-9983.	7.1	14
117	Soluble syntaxin 3 functions as a transcriptional regulator. <i>Journal of Biological Chemistry</i> , 2018, 293, 5478-5491.	3.4	14
118	Normalized coefficient of variation (nCV): a method to evaluate circadian clock robustness in population scale data. <i>Bioinformatics</i> , 2021, 37, 4581-4583.	4.1	13
119	Comparative Analysis of Human Genome Assemblies Reveals Genome-Level Differences. <i>Genomics</i> , 2002, 80, 138-139.	2.9	12
120	Exploring Trafficking GTPase Function by mRNA Expression Profiling: Use of the SymAtlas Web Application and the Membrane Datasets. <i>Methods in Enzymology</i> , 2005, 403, 1-10.	1.0	11
121	The network as the target. <i>Wiley Interdisciplinary Reviews: Systems Biology and Medicine</i> , 2010, 2, 127-133.	6.6	11
122	Sleep and Circadian Medicine. <i>Neurologic Clinics</i> , 2019, 37, 615-629.	1.8	11
123	Genome-wide studies of time of day in the brain: Design and analysis. <i>Brain Science Advances</i> , 2020, 6, 92-105.	0.9	10
124	Clean Thoughts about Dirty Genes. <i>Journal of Biological Rhythms</i> , 2004, 19, 3-9.	2.6	9
125	When Should You Take Your Medicines?. <i>Journal of Biological Rhythms</i> , 2019, 34, 582-583.	2.6	9
126	Detection Theory in Identification of RNA-DNA Sequence Differences Using RNA-Sequencing. <i>PLoS ONE</i> , 2014, 9, e112040.	2.5	7



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127	The central melanocortin system mediates the benefits of time-restricted feeding on energy balance. <i>Physiology and Behavior</i> , 2020, 227, 113132.	2.1	7
128	Circadian Rhythms: Move Over Neurons – Astrocytes Mediate SCN Clock Function. <i>Current Biology</i> , 2017, 27, R350-R352.	3.9	6
129	<i>duper</i> is a null mutation of Cryptochrome 1 in Syrian hamsters. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2123560119.	7.1	6
130	A CRY in the Night. <i>Developmental Cell</i> , 2011, 20, 144-145.	7.0	5
131	NetAtlas: a Cytoscape plugin to examine signaling networks based on tissue gene expression. <i>In Silico Biology</i> , 2008, 8, 47-52.	0.9	5
132	It's all in a day's work: Regulation of DNA excision repair by the circadian clock. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 2481-2482.	7.1	4
133	Analysis of Diurnal Variations in Heart Rate: Potential Applications for Chronobiology and Cardiovascular Medicine. <i>Frontiers in Physiology</i> , 2022, 13, 835198.	2.8	3
134	Comparative genomics as a tool in the understanding of eukaryotic transcriptional regulation. <i>Current Opinion in Genetics and Development</i> , 2005, 15, 634-639.	3.3	2
135	High Throughput Genomic Screen Identifies Multiple Factors That Promote Cooperative Wnt Signaling. <i>PLoS ONE</i> , 2013, 8, e55782.	2.5	2
136	CAMPAREE: a robust and configurable RNA expression simulator. <i>BMC Genomics</i> , 2021, 22, 692.	2.8	2
137	Clock Gene Wikis Available: Join the “Long Tail”. <i>Journal of Biological Rhythms</i> , 2008, 23, 456-457.	2.6	1
138	Polycystin-1 stimulates skeletogenesis via TAZ-mediated activation of RunX2. <i>FASEB Journal</i> , 2012, 26, 1b811.	0.5	1
139	It's not all in the brain. <i>ELife</i> , 2017, 6, .	6.0	0
140	Reply to Furlan et al.: The role of SIRT1 in cell autonomous clock function. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 13173-13173.	7.1	0
141	Circadian biology in translation. <i>FASEB Journal</i> , 2019, 33, 344.3.	0.5	0
142	Response to Lowering Nighttime Blood Pressure with Bedtime Dosing of Antihypertensive Medications: Controversies in Hypertension - Pro Side of the Argument. <i>Hypertension</i> , 2021, 78, 893.	2.7	0
143	Poor Sleep Quality in Pediatric HSCT Recipients. <i>Transplantation and Cellular Therapy</i> , 2022, 28, S398.	1.2	0