

Zachary Campbell

List of Publications by Year in descending order

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Version: 2024-02-01

44
papers

1,623
citations

304743

22
h-index

330143

37
g-index

50
all docs

50
docs citations

50
times ranked

2022
citing authors

#	ARTICLE	IF	CITATIONS
1	4D Printing of Engineered Living Materials. <i>Advanced Functional Materials</i> , 2022, 32, 2106843.	14.9	38
2	The space between notes: emerging roles for translationally silent ribosomes. <i>Trends in Biochemical Sciences</i> , 2022, 47, 477-491.	7.5	9
3	Bipartite interaction sites differentially modulate RNA-binding affinity of a protein complex essential for germline stem cell self-renewal. <i>Nucleic Acids Research</i> , 2022, 50, 536-548.	14.5	5
4	Global analyses of mRNA expression in human sensory neurons reveal eIF5A as a conserved target for inflammatory pain. <i>FASEB Journal</i> , 2022, 36, .	0.5	6
5	A compendium of validated pain genes. <i>WIREs Mechanisms of Disease</i> , 2022, 14, .	3.3	5
6	A Highly Selective MNK Inhibitor Rescues Deficits Associated with Fragile X Syndrome in Mice. <i>Neurotherapeutics</i> , 2021, 18, 624-639.	4.4	9
7	Stimuli-responsive engineered living materials. <i>Soft Matter</i> , 2021, 17, 785-809.	2.7	64
8	A peptide encoded within a 5' untranslated region promotes pain sensitization in mice. <i>Pain</i> , 2021, 162, 1864-1875.	4.2	8
9	Intercellular Arc Signaling Regulates Vasodilation. <i>Journal of Neuroscience</i> , 2021, 41, 7712-7726.	3.6	12
10	A role for translational regulation by S6 kinase and a downstream target in inflammatory pain. <i>British Journal of Pharmacology</i> , 2021, 178, 4675-4690.	5.4	5
11	Functionally distinct roles for eEF2K in the control of ribosome availability and p-body abundance. <i>Nature Communications</i> , 2021, 12, 6789.	12.8	18
12	Principles of mRNA control by human PUM proteins elucidated from multimodal experiments and integrative data analysis. <i>Rna</i> , 2020, 26, 1680-1703.	3.5	14
13	Conserved Expression of Nav1.7 and Nav1.8 Contribute to the Spontaneous and Thermally Evoked Excitability in IL-6 and NGF-Sensitized Adult Dorsal Root Ganglion Neurons In Vitro. <i>Bioengineering</i> , 2020, 7, 44.	3.5	9
14	Type I Interferons Act Directly on Nociceptors to Produce Pain Sensitization: Implications for Viral Infection-Induced Pain. <i>Journal of Neuroscience</i> , 2020, 40, 3517-3532.	3.6	62
15	Shape-morphing living composites. <i>Science Advances</i> , 2020, 6, eaax8582.	10.3	53
16	Molecular entrapment by RNA: an emerging tool for disrupting protein-RNA interactions in vivo. <i>RNA Biology</i> , 2020, 17, 417-424.	3.1	4
17	Differences between Dorsal Root and Trigeminal Ganglion Nociceptors in Mice Revealed by Translational Profiling. <i>Journal of Neuroscience</i> , 2019, 39, 6829-6847.	3.6	66
18	RNA control in pain: Blame it on the messenger. <i>Wiley Interdisciplinary Reviews RNA</i> , 2019, 10, e1546.	6.4	12

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19	Activation of the integrated stress response in nociceptors drives methylglyoxal-induced pain. <i>Pain</i> , 2019, 160, 160-171.	4.2	45
20	Emerging neurotechnology for antinociceptive mechanisms and therapeutics discovery. <i>Biosensors and Bioelectronics</i> , 2019, 126, 679-689.	10.1	19
21	Nociceptor Translational Profiling Reveals the Ragulator-Rag GTPase Complex as a Critical Generator of Neuropathic Pain. <i>Journal of Neuroscience</i> , 2019, 39, 393-411.	3.6	95
22	Engineering a conserved RNA regulatory protein repurposes its biological function in vivo. <i>ELife</i> , 2019, 8, .	6.0	13
23	A crystal structure of a collaborative RNA regulatory complex reveals mechanisms to refine target specificity. <i>ELife</i> , 2019, 8, .	6.0	21
24	RNA-binding proteins as targets for pain therapeutics. <i>Neurobiology of Pain (Cambridge, Mass)</i> , 2018, 4, 2-7.	2.5	13
25	Inhibition of Poly(A)-binding protein with a synthetic RNA mimic reduces pain sensitization in mice. <i>Nature Communications</i> , 2018, 9, 10.	12.8	135
26	Global pairwise RNA interaction landscapes reveal core features of protein recognition. <i>Nature Communications</i> , 2018, 9, 2511.	12.8	29
27	Adult mouse sensory neurons on microelectrode arrays exhibit increased spontaneous and stimulus-evoked activity in the presence of interleukin-6. <i>Journal of Neurophysiology</i> , 2018, 120, 1374-1385.	1.8	32
28	Architecture and dynamics of overlapped RNA regulatory networks. <i>Rna</i> , 2017, 23, 1636-1647.	3.5	32
29	Integrated analysis of RNA-binding protein complexes using in vitro selection and high-throughput sequencing and sequence specificity landscapes (SEQRS). <i>Methods</i> , 2017, 118-119, 171-181.	3.8	24
30	Drosophila Nanos acts as a molecular clamp that modulates the RNA-binding and repression activities of Pumilio. <i>ELife</i> , 2016, 5, .	6.0	66
31	RNA regulatory networks diversified through curvature of the PUF protein scaffold. <i>Nature Communications</i> , 2015, 6, 8213.	12.8	56
32	Probing RNA-protein networks: biochemistry meets genomics. <i>Trends in Biochemical Sciences</i> , 2015, 40, 157-164.	7.5	39
33	A protein-RNA specificity code enables targeted activation of an endogenous human transcript. <i>Nature Structural and Molecular Biology</i> , 2014, 21, 732-738.	8.2	74
34	Biochemical Characterization of the Caenorhabditis elegans FBF...CPB-1 Translational Regulation Complex Identifies Conserved Protein Interaction Hotspots. <i>Journal of Molecular Biology</i> , 2013, 425, 725-737.	4.2	18
35	A Protein...Protein Interaction Platform Involved in Recruitment of GLD-3 to the FBF...fem-3 mRNA Complex. <i>Journal of Molecular Biology</i> , 2013, 425, 738-754.	4.2	16
36	Identification of a Conserved Interface between PUF and CPEB Proteins. <i>Journal of Biological Chemistry</i> , 2012, 287, 18854-18862.	3.4	40

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37	Patterns and plasticity in RNA-protein interactions enable recruitment of multiple proteins through a single site. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 6054-6059.	7.1	43
38	Cooperativity in RNA-Protein Interactions: Global Analysis of RNA Binding Specificity. <i>Cell Reports</i> , 2012, 1, 570-581.	6.4	106
39	A conserved PUF-â€œAgo-â€œeEF1A complex attenuates translation elongation. <i>Nature Structural and Molecular Biology</i> , 2012, 19, 176-183.	8.2	128
40	Analysis of the Bacterial Luciferase Mobile Loop by Replica-Exchange Molecular Dynamics. <i>Biophysical Journal</i> , 2010, 99, 4012-4019.	0.5	20
41	Two Lysine Residues in the Bacterial Luciferase Mobile Loop Stabilize Reaction Intermediates. <i>Journal of Biological Chemistry</i> , 2009, 284, 32827-32834.	3.4	21
42	Fre Is the Major Flavin Reductase Supporting Bioluminescence from <i>Vibrio harveyi</i> Luciferase in <i>Escherichia coli</i> . <i>Journal of Biological Chemistry</i> , 2009, 284, 8322-8328.	3.4	44
43	Crystal Structure of the Bacterial Luciferase/Flavin Complex Provides Insight into the Function of the Î² Subunit. <i>Biochemistry</i> , 2009, 48, 6085-6094.	2.5	92
44	Intercellular Arc Signaling Regulates Vasodilation. <i>SSRN Electronic Journal</i> , 0, , .	0.4	2