

# James A Coffman

## List of Publications by Year in descending order

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Version: 2024-02-01

51  
papers

2,760  
citations

304743

22  
h-index

206112

48  
g-index

53  
all docs

53  
docs citations

53  
times ranked

2789  
citing authors

#	ARTICLE	IF	CITATIONS
1	The Genome of the Sea Urchin <i>Strongylocentrotus purpuratus</i> . <i>Science</i> , 2006, 314, 941-952.	12.6	1,018
2	Runx transcription factors and the developmental balance between cell proliferation and differentiation. <i>Cell Biology International</i> , 2003, 27, 315-324.	3.0	173
3	The genomic underpinnings of apoptosis in <i>Strongylocentrotus purpuratus</i> . <i>Developmental Biology</i> , 2006, 300, 321-334.	2.0	111
4	Oral-aboral axis specification in the sea urchin embryo. <i>Developmental Biology</i> , 2004, 273, 160-171.	2.0	101
5	Oral-Aboral Axis Specification in the Sea Urchin Embryo. <i>Developmental Biology</i> , 2001, 230, 18-28.	2.0	94
6	Structural analysis of proteins by capillary HPLC electrospray tandem mass spectrometry. <i>International Journal of Mass Spectrometry and Ion Processes</i> , 1991, 111, 131-149.	1.8	86
7	The sea urchin kinome: A first look. <i>Developmental Biology</i> , 2006, 300, 180-193.	2.0	84
8	The evolution of Runx genes I. A comparative study of sequences from phylogenetically diverse model organisms. <i>BMC Evolutionary Biology</i> , 2003, 3, 4.	3.2	81
9	Cis-regulatory control of the nodal gene, initiator of the sea urchin oral ectoderm gene network. <i>Developmental Biology</i> , 2007, 306, 860-869.	2.0	78
10	Cell Cycle Development. <i>Developmental Cell</i> , 2004, 6, 321-327.	7.0	77
11	A hyaline layer protein that becomes localized to the oral ectoderm and foregut of sea urchin embryos. <i>Developmental Biology</i> , 1990, 140, 93-104.	2.0	72
12	Oral-aboral axis specification in the sea urchin embryo. <i>Developmental Biology</i> , 2009, 330, 123-130.	2.0	69
13	Cortisol-treated zebrafish embryos develop into pro-inflammatory adults with aberrant immune gene regulation. <i>Biology Open</i> , 2016, 5, 1134-1141.	1.2	61
14	SpRunt-1, a New Member of the Runt Domain Family of Transcription Factors, Is a Positive Regulator of the Aboral Ectoderm-Specific <i>CyIII</i> Gene in Sea Urchin Embryos. <i>Developmental Biology</i> , 1996, 174, 43-54.	2.0	59
15	The genomic repertoire for cell cycle control and DNA metabolism in <i>S. purpuratus</i> . <i>Developmental Biology</i> , 2006, 300, 238-251.	2.0	48
16	Mitochondria, redox signaling and axis specification in metazoan embryos. <i>Developmental Biology</i> , 2007, 308, 266-280.	2.0	43
17	Gene Expression Changes Associated With the Developmental Plasticity of Sea Urchin Larvae in Response to Food Availability. <i>Biological Bulletin</i> , 2015, 228, 171-180.	1.8	38
18	The expression of SpRunt during sea urchin embryogenesis. <i>Mechanisms of Development</i> , 2002, 117, 327-330.	1.7	37

#	ARTICLE	IF	CITATIONS
19	SpGCF1, a Sea Urchin Embryo DNA-Binding Protein, Exists as Five Nested Variants Encoded by a Single mRNA. <i>Developmental Biology</i> , 1995, 169, 713-727.	2.0	31
20	Expression of spatially regulated genes in the sea urchin embryo. <i>Current Opinion in Genetics and Development</i> , 1992, 2, 260-268.	3.3	30
21	Evaluation of developmental phenotypes produced by morpholino antisense targeting of a sea urchin Runx gene. <i>BMC Biology</i> , 2004, 2, 6.	3.8	27
22	Is Runx a linchpin for developmental signaling in metazoans?. <i>Journal of Cellular Biochemistry</i> , 2009, 107, 194-202.	2.6	24
23	Klf9 is a key feedforward regulator of the transcriptomic response to glucocorticoid receptor activity. <i>Scientific Reports</i> , 2020, 10, 11415.	3.3	24
24	Runx Expression Is Mitogenic and Mutually Linked to Wnt Activity in Blastula-Stage Sea Urchin Embryos. <i>PLoS ONE</i> , 2008, 3, e3770.	2.5	22
25	Oral-aboral axis specification in the sea urchin embryo, IV: Hypoxia radializes embryos by preventing the initial spatialization of nodal activity. <i>Developmental Biology</i> , 2014, 386, 302-307.	2.0	22
26	Redox regulation of development and regeneration. <i>Current Opinion in Genetics and Development</i> , 2019, 57, 9-15.	3.3	22
27	Developmental Ascendency: From Bottom-up to Top-down Control. <i>Biological Theory</i> , 2006, 1, 165-178.	1.5	21
28	Runx-dependent expression of PKC is critical for cell survival in the sea urchin embryo. <i>BMC Biology</i> , 2005, 3, 18.	3.8	20
29	Mitochondria and metazoan epigenesis. <i>Seminars in Cell and Developmental Biology</i> , 2009, 20, 321-329.	5.0	18
30	Simple and fast quantification of DNA damage by real-time PCR, and its application to nuclear and mitochondrial DNA from multiple tissues of aging zebrafish. <i>BMC Research Notes</i> , 2017, 10, 269.	1.4	17
31	The evolution of Runx genes II. The C-terminal Groucho recruitment motif is present in both eumetazoans and homoscleromorphs but absent in a haplosclerid demosponge. <i>BMC Research Notes</i> , 2009, 2, 59.	1.4	13
32	Chronic stress, physiological adaptation and developmental programming of the neuroendocrine stress system. <i>Future Neurology</i> , 2020, 15, FNL39.	0.5	13
33	CBFbeta is a facultative Runx partner in the sea urchin embryo. <i>BMC Biology</i> , 2006, 4, 4.	3.8	12
34	Comparative biology of tissue repair, regeneration and aging. <i>Npj Regenerative Medicine</i> , 2016, 1, .	5.2	12
35	On Causality in Nonlinear Complex Systems. , 2011, , 287-309.		11
36	Oxygen, pH, and oral-aboral axis specification in the sea urchin embryo. <i>Molecular Reproduction and Development</i> , 2011, 78, 68-68.	2.0	10

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37	<i>Nodal</i> -mediated epigenesis requires dynamin-mediated endocytosis. <i>Developmental Dynamics</i> , 2011, 240, 704-711.	1.8	10
38	Glucocorticoid-Mediated Developmental Programming of Vertebrate Stress Responsivity. <i>Frontiers in Physiology</i> , 2021, 12, 812195.	2.8	10
39	On the Meaning of Chance in Biology. <i>Biosemiotics</i> , 2014, 7, 377-388.	1.4	8
40	An Elk transcription factor is required for Runx-dependent survival signaling in the sea urchin embryo. <i>Developmental Biology</i> , 2016, 416, 173-186.	2.0	8
41	Chronic cortisol exposure in early development leads to neuroendocrine dysregulation in adulthood. <i>BMC Research Notes</i> , 2020, 13, 366.	1.4	8
42	Glucocorticoid-Responsive Transcription Factor Krüppel-Like Factor 9 Regulates <i>flbp5</i> and Metabolism. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 727037.	3.7	7
43	Developmental cis-regulatory analysis of the cyclin D gene in the sea urchin <i>Strongylocentrotus purpuratus</i> . <i>Biochemical and Biophysical Research Communications</i> , 2013, 440, 413-418.	2.1	6
44	Sea urchin <i>akt</i> activity is Runx-dependent and required for post-cleavage stage cell division. <i>Biology Open</i> , 2013, 2, 472-478.	1.2	6
45	Information as a Manifestation of Development. <i>Information (Switzerland)</i> , 2011, 2, 102-116.	2.9	5
46	Identification of Sequence-Specific DNA Binding Proteins. <i>Methods in Cell Biology</i> , 2004, 74, 653-675.	1.1	4
47	On reductionism, organicism, somatic mutations and cancer. <i>BioEssays</i> , 2005, 27, 459-459.	2.5	4
48	Ping Ao's "Darwinian Dynamics Implies Developmental Ascendency ( <i>Biological Theory</i> 2: 113-115, 2007). <i>Biological Theory</i> , 2007, 2, 179-180.	1.5	2
49	Mitochondrial patterns and function in animal development. <i>Seminars in Cell and Developmental Biology</i> , 2009, 20, 320.	5.0	1
50	Interview with James Coffman: early-life stress in adult illness. <i>Future Neurology</i> , 2017, 12, 9-11.	0.5	0
51	Why Functional Genomics Is the Central Concern of Biology and the Hard Problem of Abiogenesis. <i>Springer Proceedings in Complexity</i> , 2019, , 327-337.	0.3	0