

Eric Rättinger

List of Publications by Year in descending order

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3,203
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394421

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46
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docs citations

46
times ranked

2714
citing authors

#	ARTICLE	IF	CITATIONS
1	A panmetazoan concept for adult stem cells: the wobbling Penrose landscape. <i>Biological Reviews</i> , 2022, 97, 299-325.	10.4	25
2	Intrinsically High Capacity of Animal Cells From a Symbiotic Cnidarian to Deal With Pro-Oxidative Conditions. <i>Frontiers in Physiology</i> , 2022, 13, 819111.	2.8	3
3	Creating a User-Friendly and Open-Access Gene Expression Database for Comparing Embryonic Development and Regeneration in <i>Nematostella vectensis</i> . <i>Methods in Molecular Biology</i> , 2022, 2450, 649-662.	0.9	0
4	Horizontal acquisition of Symbiodiniaceae in the <i>Anemonia viridis</i> (Cnidaria, Anthozoa) species complex. <i>Molecular Ecology</i> , 2021, 30, 391-405.	3.9	0
5	<i>Nematostella vectensis</i> , an Emerging Model for Deciphering the Molecular and Cellular Mechanisms Underlying Whole-Body Regeneration. <i>Cells</i> , 2021, 10, 2692.	4.1	15
6	Transcriptomic Analysis in the Sea Anemone <i>Nematostella vectensis</i> . <i>Methods in Molecular Biology</i> , 2021, 2219, 231-240.	0.9	4
7	Experimental Tools to Study Regeneration in the Sea Anemone <i>Nematostella vectensis</i> . <i>Methods in Molecular Biology</i> , 2021, 2219, 69-80.	0.9	8
8	Cnidarian Cell Cryopreservation: A Powerful Tool for Cultivation and Functional Assays. <i>Cells</i> , 2020, 9, 2541.	4.1	5
9	The Diversity of Muscles and Their Regenerative Potential across Animals. <i>Cells</i> , 2020, 9, 1925.	4.1	9
10	NvERTx: A gene expression database to compare embryogenesis and regeneration in the sea anemone <i>Nematostella vectensis</i> . <i>Development (Cambridge)</i> , 2018, 145, .	2.5	47
11	A novel technique to combine and analyse spatial and temporal expression datasets: A case study with the sea anemone <i>Nematostella vectensis</i> to identify potential gene interactions. <i>Developmental Biology</i> , 2017, 428, 204-214.	2.0	5
12	A bipolar role of the transcription factor ERG for cnidarian germ layer formation and apical domain patterning. <i>Developmental Biology</i> , 2017, 430, 346-361.	2.0	24
13	The rise of the starlet sea anemone <i>Nematostella vectensis</i> as a model system to investigate development and regeneration. <i>Wiley Interdisciplinary Reviews: Developmental Biology</i> , 2016, 5, 408-428.	5.9	121
14	MAPK signaling is necessary for neurogenesis in <i>Nematostella vectensis</i> . <i>BMC Biology</i> , 2016, 14, 61.	3.8	51
15	Diversity of Cnidarian Muscles: Function, Anatomy, Development and Regeneration. <i>Frontiers in Cell and Developmental Biology</i> , 2016, 4, 157.	3.7	60
16	Domain analysis of the <i>Nematostella vectensis</i> SNAIL ortholog reveals unique nucleolar localization that depends on the zinc-finger domains. <i>Scientific Reports</i> , 2015, 5, 12147.	3.3	6
17	Analysis of a spatial gene expression database for sea anemone <i>Nematostella vectensis</i> during early development. <i>BMC Systems Biology</i> , 2015, 9, 63.	3.0	4
18	Characterization of Morphological and Cellular Events Underlying Oral Regeneration in the Sea Anemone, <i>Nematostella vectensis</i> . <i>International Journal of Molecular Sciences</i> , 2015, 16, 28449-28471.	4.1	62

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19	Nodal signaling is required for mesodermal and ventral but not for dorsal fates in the indirect developing hemichordate, <i>Ptychodera flava</i> . <i>Biology Open</i> , 2015, 4, 830-842.	1.2	33
20	Hemichordata. , 2015, , 59-89.		14
21	Autophagy : Moving Benchside Promises to Patient Bedsides. <i>Current Cancer Drug Targets</i> , 2015, 15, 684-702.	1.6	14
22	A Computational Approach towards a Gene Regulatory Network for the Developing <i>Nematostella vectensis</i> Gut. <i>PLoS ONE</i> , 2014, 9, e103341.	2.5	15
23	In vivo imaging of <i>Nematostella vectensis</i> embryogenesis and late development using fluorescent probes. <i>BMC Cell Biology</i> , 2014, 15, 44.	3.0	20
24	Microinjection of mRNA or morpholinos for reverse genetic analysis in the starlet sea anemone, <i>Nematostella vectensis</i> . <i>Nature Protocols</i> , 2013, 8, 924-934.	12.0	73
25	Reciprocal Signaling between the Ectoderm and a Mesendodermal Left-Right Organizer Directs Left-Right Determination in the Sea Urchin Embryo. <i>PLoS Genetics</i> , 2012, 8, e1003121.	3.5	59
26	A Framework for the Establishment of a Cnidarian Gene Regulatory Network for "Endomesoderm" Specification: The Inputs of β -Catenin/TCF Signaling. <i>PLoS Genetics</i> , 2012, 8, e1003164.	3.5	116
27	Evolutionary crossroads in developmental biology: hemichordates. <i>Development (Cambridge)</i> , 2012, 139, 2463-2475.	2.5	59
28	A comparative gene expression database for invertebrates. <i>EvoDevo</i> , 2011, 2, 17.	3.2	21
29	Ancestral Regulatory Circuits Governing Ectoderm Patterning Downstream of Nodal and BMP2/4 Revealed by Gene Regulatory Network Analysis in an Echinoderm. <i>PLoS Genetics</i> , 2010, 6, e1001259.	3.5	133
30	Centralization of the Deuterostome Nervous System Predates Chordates. <i>Current Biology</i> , 2009, 19, 1264-1269.	3.9	110
31	FGF signals guide migration of mesenchymal cells, control skeletal morphogenesis and regulate gastrulation during sea urchin development. <i>Development (Cambridge)</i> , 2008, 135, 353-365.	2.5	133
32	The Genome of the Sea Urchin <i>Strongylocentrotus purpuratus</i> . <i>Science</i> , 2006, 314, 941-952.	12.6	1,018
33	Expression pattern of three putative RNA-binding proteins during early development of the sea urchin <i>Paracentrotus lividus</i> . <i>Gene Expression Patterns</i> , 2006, 6, 864-872.	0.8	10
34	RTK and TGF- β signaling pathways genes in the sea urchin genome. <i>Developmental Biology</i> , 2006, 300, 132-152.	2.0	140
35	Nemo-like kinase (NLK) acts downstream of Notch/Delta signalling to downregulate TCF during mesoderm induction in the sea urchin embryo. <i>Development (Cambridge)</i> , 2006, 133, 4341-4353.	2.5	52
36	Left-Right Asymmetry in the Sea Urchin Embryo Is Regulated by Nodal Signaling on the Right Side. <i>Developmental Cell</i> , 2005, 9, 147-158.	7.0	242

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37	A Raf/MEK/ERK signaling pathway is required for development of the sea urchin embryo micromere lineage through phosphorylation of the transcription factor Ets. <i>Development (Cambridge)</i> , 2004, 131, 1075-1087.	2.5	110
38	Nodal and BMP2/4 Signaling Organizes the Oral-Aboral Axis of the Sea Urchin Embryo. <i>Developmental Cell</i> , 2004, 6, 397-410.	7.0	331