

# Andrade Rp

## List of Publications by Year in descending order

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Version: 2024-02-01

32  
papers

816  
citations

516215

16  
h-index

500791

28  
g-index

36  
all docs

36  
docs citations

36  
times ranked

1062  
citing authors

#	ARTICLE	IF	CITATIONS
1	gga-miRNOME, a microRNA-sequencing dataset from chick embryonic tissues. <i>Scientific Data</i> , 2022, 9, 29.	2.4	4
2	Cellâ€™Fibronectin Interactions and Actomyosin Contractility Regulate the Segmentation Clock and Spatio-Temporal Somite Cleft Formation during Chick Embryo Somitogenesis. <i>Cells</i> , 2022, 11, 2003.	1.8	1
3	Altered Cogs of the Clock: Insights into the Embryonic Etiology of Spondylocostal Dysostosis. <i>Journal of Developmental Biology</i> , 2021, 9, 5.	0.9	5
4	Brachyury Is Associated with Glioma Differentiation and Response to Temozolomide. <i>Neurotherapeutics</i> , 2020, 17, 2015-2027.	2.1	7
5	NineTeen Complex-subunit Salsa is required for efficient splicing of a subset of introns and dorsalâ€™ventral patterning. <i>Rna</i> , 2020, 26, 1935-1956.	1.6	2
6	The Tâ€™box transcription factor brachyury behaves as a tumor suppressor in gliomas. <i>Journal of Pathology</i> , 2020, 251, 87-99.	2.1	10
7	StemMapper: a curated gene expression database for stem cell lineage analysis. <i>Nucleic Acids Research</i> , 2018, 46, D788-D793.	6.5	28
8	rdml: A Mathematica package for parsing and importing Real-Time qPCR data. <i>BMC Research Notes</i> , 2017, 10, 208.	0.6	1
9	Getting a handle on embryo limb development: Molecular interactions driving limb outgrowth and patterning. <i>Seminars in Cell and Developmental Biology</i> , 2016, 49, 92-101.	2.3	20
10	The embryonic Brachyury transcription factor is a novel biomarker of GIST aggressiveness and poor survival. <i>Gastric Cancer</i> , 2016, 19, 651-659.	2.7	18
11	Mechanisms of vertebrate embryo segmentation: Common themes in trunk and limb development. <i>Seminars in Cell and Developmental Biology</i> , 2016, 49, 125-134.	2.3	20
12	Brachyury as a potential modulator of androgen receptor activity and a key player in therapy resistance in prostate cancer. <i>Oncotarget</i> , 2016, 7, 28891-28902.	0.8	19
13	Patterning in time and space: HoxB cluster gene expression in the developing chick embryo. <i>Cell Cycle</i> , 2015, 14, 135-145.	1.3	10
14	Timing Embryo Segmentation: Dynamics and Regulatory Mechanisms of the Vertebrate Segmentation Clock. <i>BioMed Research International</i> , 2014, 2014, 1-12.	0.9	26
15	T-box Transcription Factor Brachyury Is Associated with Prostate Cancer Progression and Aggressiveness. <i>Clinical Cancer Research</i> , 2014, 20, 4949-4961.	3.2	67
16	MicroRNA processing machinery in the developing chick embryo. <i>Gene Expression Patterns</i> , 2014, 16, 114-121.	0.3	4
17	Limb Patterning: From Signaling Gradients to Molecular Oscillations. <i>Journal of Molecular Biology</i> , 2014, 426, 780-784.	2.0	16
18	Joint interpretation of AER/FGF and ZPA/SHH over time and space underlies <i>hairy2</i> expression in the chick limb. <i>Biology Open</i> , 2012, 1, 1102-1110.	0.6	13

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19	Retinoic acid signaling regulates embryonic clock hairy2 gene expression in the developing chick limb. <i>Biochemical and Biophysical Research Communications</i> , 2012, 423, 889-894.	1.0	5
20	Sonic hedgehog in temporal control of somite formation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 12907-12912.	3.3	62
21	Comprehensive analysis of fibroblast growth factor receptor expression patterns during chick forelimb development. <i>International Journal of Developmental Biology</i> , 2010, 54, 1515-1524.	0.3	21
22	Redefining the role of ectoderm in somitogenesis: a player in the formation of the fibronectin matrix of presomitic mesoderm. <i>Development (Cambridge)</i> , 2007, 134, 3155-3165.	1.2	59
23	Progressive mRNA decay establishes an mkp3 expression gradient in the chick limb bud. <i>Biochemical and Biophysical Research Communications</i> , 2007, 352, 153-157.	1.0	11
24	Molecular clocks underlying vertebrate embryo segmentation: A 10-year-old hairy-go-round. <i>Birth Defects Research Part C: Embryo Today Reviews</i> , 2007, 81, 65-83.	3.6	19
25	Chick Hairy1 protein interacts with Sap18, a component of the Sin3/HDAC transcriptional repressor complex. <i>BMC Developmental Biology</i> , 2007, 7, 83.	2.1	8
26	Multiple transcripts regulate glucose-triggered mRNA decay of the lactate transporter JEN1 from <i>Saccharomyces cerevisiae</i> . <i>Biochemical and Biophysical Research Communications</i> , 2005, 332, 254-262.	1.0	30
27	Thinking clockwise. <i>Brain Research Reviews</i> , 2005, 49, 114-119.	9.1	13
28	Functional expression of the lactate permease Jen1p of <i>Saccharomyces cerevisiae</i> in <i>Pichia pastoris</i> . <i>Biochemical Journal</i> , 2003, 376, 781-787.	1.7	35
29	Expression of the Lactate Permease Gene JEN1 from the Yeast <i>Saccharomyces cerevisiae</i> . <i>Fungal Genetics and Biology</i> , 2001, 32, 105-111.	0.9	38
30	RNase II removes the oligo(A) tails that destabilize the rpsO mRNA of <i>Escherichia coli</i> . <i>Rna</i> , 2000, 6, 1185-1193.	1.6	73
31	The Lactate-Proton Symport of <i>Saccharomyces cerevisiae</i> Is Encoded by <i>JEN1</i> . <i>Journal of Bacteriology</i> , 1999, 181, 2620-2623.	1.0	166
32	The posterior limit of the <i>area pellucida</i> (pPL) as a reliable proxy for the end of the primitive streak in chick elongation studies. <i>Matters</i> , 0, , .	1.0	0