Andrey Ptitsyn

List of Publications by Year in descending order

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ΔΝΠΦΕΥ ΡΤΙΤΟΥΝ

#	Article	IF	CITATIONS
1	The ctenophore genome and the evolutionary origins of neural systems. Nature, 2014, 510, 109-114.	27.8	606
2	Characterization of Peripheral Circadian Clocks in Adipose Tissues. Diabetes, 2006, 55, 962-970.	0.6	443
3	A Comprehensive Approach to Clustering of Expressed Human Gene Sequence: The Sequence Tag Alignment and Consensus Knowledge Base. Genome Research, 1999, 9, 1143-1155.	5.5	175
4	Small RNA profiling of Dengue virus-mosquito interactions implicates the PIWI RNA pathway in anti-viral defense. BMC Microbiology, 2011, 11, 45.	3.3	155
5	Circadian Clocks Are Resounding in Peripheral Tissues. PLoS Computational Biology, 2006, 2, e16.	3.2	117
6	Yield and characterization of subcutaneous human adipose-derived stem cells by flow cytometric and adipogenic mRNA analyzes. Cytotherapy, 2010, 12, 538-546.	0.7	111
7	Circadian Oscillation of Gene Expression in Murine Calvarial Bone. Journal of Bone and Mineral Research, 2007, 22, 357-365.	2.8	91
8	Circadian Expression of Genes Regulating Food Intake. Obesity, 2007, 15, 607-615.	3.0	63
9	Resveratrol suppresses human colon cancer cell proliferation and induces apoptosis via targeting the pentose phosphate and the talin-FAK signaling pathways-A proteomic approach. Proteome Science, 2011, 9, 49.	1.7	57
10	Eukaryotic genes in Mycobacterium tuberculosis could have a role in pathogenesis and immunomodulation. Trends in Genetics, 2002, 18, 5-8.	6.7	53
11	Digital Signal Processing Reveals Circadian Baseline Oscillation in Majority of Mammalian Genes. PLoS Computational Biology, 2007, 3, e120.	3.2	45
12	Prospective influences of circadian clocks in adipose tissue and metabolism. Nature Reviews Endocrinology, 2011, 7, 98-107.	9.6	38
13	True or false: All genes are rhythmic. Annals of Medicine, 2011, 43, 1-12.	3.8	33
14	Circadian Rhythm of Osteocalcin in the Maxillomandibular Complex. Journal of Dental Research, 2009, 88, 45-50.	5.2	31
15	Proteome of Human Subcutaneous Adipose Tissue Stromal Vascular Fraction Cells versus Mature Adipocytes Based on DIGE. Journal of Proteome Research, 2011, 10, 1519-1527.	3.7	28
16	Persistent fetal infection with bovine viral diarrhea virus differentially affects maternal blood cell signal transduction pathways. Physiological Genomics, 2009, 36, 129-139.	2.3	27
17	The American lobster genome reveals insights on longevity, neural, and immune adaptations. Science Advances, 2021, 7, .	10.3	27
18	Unsupervised clustering of gene expression data points at hypoxia as possible trigger for metabolic syndrome. BMC Genomics, 2006, 7, 318.	2.8	26

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19	A systems biology approach to identify molecular pathways altered by HDAC inhibition in osteosarcoma. Journal of Cellular Biochemistry, 2012, 113, 773-783.	2.6	19
20	CLU: A new algorithm for EST clustering. BMC Bioinformatics, 2005, 6, S3.	2.6	16
21	Comprehensive analysis of circadian periodic pattern in plant transcriptome. BMC Bioinformatics, 2008, 9, S18.	2.6	14
22	Stochastic Resonance Reveals "Pilot Light―Expression in Mammalian Genes. PLoS ONE, 2008, 3, e1842.	2.5	10
23	Computational workflow for analysis of gain and loss of genes in distantly related genomes. BMC Bioinformatics, 2012, 13, S5.	2.6	7
24	Class Discovery Analysis of the Lung Cancer Gene Expression Data. DNA and Cell Biology, 2004, 23, 715-721.	1.9	6
25	The role of alternative Polyadenylation in regulation of rhythmic gene expression. BMC Genomics, 2017, 18, 576.	2.8	6
26	Regional Societies: Fostering Competitive Research Through Virtual Infrastructures. PLoS Biology, 2004, 2, e372.	5.6	4
27	Transformation of metabolism with age and lifestyle in Antarctic seals: a case study of systems biology approach to cross-species microarray experiment. BMC Systems Biology, 2010, 4, 133.	3.0	4
28	Application of an RNA amplification method for reliable single-cell transcriptome analysis. BioTechniques, 2015, 59, 137-148.	1.8	4
29	Semantic Assembly and Annotation of Draft RNAseq Transcripts without a Reference Genome. PLoS ONE, 2015, 10, e0138006.	2.5	4
30	Computational analysis of gene expression space associated with metastatic cancer. BMC Bioinformatics, 2009, 10, S6.	2.6	3
31	Confidence in Phase Definition for Periodicity in Genes Expression Time Series. PLoS ONE, 2015, 10, e0131111.	2.5	2
32	Optimization of data-intensive next generation sequencing in high performance computing. , 2015, , .		2
33	Circadian succession of molecular processes in living tissues. BMC Medical Genomics, 2018, 11, 14.	1.5	1
34	Modulation of mRNA circadian transcription cycle by microRNAs. , 2014, , .		0