Anis Karimpour-Fard

List of Publications by Year in descending order

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39 papers 3,119 citations

394421 19 h-index 302126 39 g-index

40 all docs

40 docs citations

40 times ranked

5166 citing authors

#	Article	IF	CITATIONS
1	Evolutionary and Biomedical Insights from the Rhesus Macaque Genome. Science, 2007, 316, 222-234.	12.6	1,283
2	Gibbon genome and the fast karyotype evolution of small apes. Nature, 2014, 513, 195-201.	27.8	320
3	Rapid profiling of a microbial genome using mixtures of barcoded oligonucleotides. Nature Biotechnology, 2010, 28, 856-862.	17.5	265
4	Lineage-Specific Gene Duplication and Loss in Human and Great Ape Evolution. PLoS Biology, 2004, 2, e207.	5.6	263
5	Gene copy number variation spanning 60 million years of human and primate evolution. Genome Research, 2007, 17, 1266-1277.	5.5	153
6	Coordinate Loss of <i>MAP3K7</i> and <i>CHD1</i> Promotes Aggressive Prostate Cancer. Cancer Research, 2015, 75, 1021-1034.	0.9	74
7	Circulating microRNA as a biomarker for recovery in pediatric dilated cardiomyopathy. Journal of Heart and Lung Transplantation, 2015, 34, 724-733.	0.6	65
8	Metabolic cycles in a circannual hibernator. Physiological Genomics, 2011, 43, 799-807.	2.3	64
9	Genome-scale identification and characterization of ethanol tolerance genes in Escherichia coli. Metabolic Engineering, 2013, 15, 124-133.	7.0	52
10	Metabolic changes associated with the long winter fast dominate the liver proteome in 13-lined ground squirrels. Physiological Genomics, 2014, 46, 348-361.	2.3	49
11	Skeletal muscle proteomics: carbohydrate metabolism oscillates with seasonal and torpor-arousal physiology of hibernation. American Journal of Physiology - Regulatory Integrative and Comparative Physiology, 2011, 301, R1440-R1452.	1.8	47
12	Pediatric dilated cardiomyopathy hearts display a unique gene expression profile. JCI Insight, 2017, 2, .	5.0	46
13	Multistate proteomics analysis reveals novel strategies used by a hibernator to precondition the heart and conserve ATP for winter heterothermy. Physiological Genomics, 2011, 43, 1263-1275.	2.3	42
14	Myocardial microRNAs associated with reverse remodeling in human heart failure. JCI Insight, 2017, 2, e89169.	5.0	42
15	Micro-RNA Expression in Hypoplastic Left Heart Syndrome. Journal of Cardiac Failure, 2015, 21, 83-88.	1.7	40
16	Amino acids in a targeted versus a non-targeted metabolomics LC-MS/MS assay. Are the results consistent?. Clinical Biochemistry, 2016, 49, 955-961.	1.9	32
17	Phosphodiesterase-5 Is Elevated in Failing Single Ventricle Myocardium and Affects Cardiomyocyte Remodeling In Vitro. Circulation: Heart Failure, 2018, 11, e004571.	3.9	32
18	Analysis of Keloid Response to 5-Fluorouracil Treatment and Long-Term Prevention of Keloid Recurrence. Plastic and Reconstructive Surgery, 2019, 143, 490-494.	1.4	30

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19	Kidney proteome changes provide evidence for a dynamic metabolism and regional redistribution of plasma proteins during torpor-arousal cycles of hibernation. Physiological Genomics, 2012, 44, 717-727.	2.3	21
20	Temporal analysis of mRNA and miRNA expression in transgenic mice overexpressing Arg- and Gly389 polymorphic variants of the \hat{l}^2 (sub) -adrenergic receptor. Physiological Genomics, 2011, 43, 1294-1306.	2.3	17
21	Investigation of factors affecting prediction of protein-protein interaction networks by phylogenetic profiling. BMC Genomics, 2007, 8, 393.	2.8	16
22	Predicting protein linkages in bacteria: Which method is best depends on task. BMC Bioinformatics, 2008, 9, 397.	2.6	16
23	The topology of the bacterial co-conserved protein network and its implications for predicting protein function. BMC Genomics, 2008, 9, 313.	2.8	16
24	A survey of computational tools for downstream analysis of proteomic and other omic datasets. Human Genomics, 2015, 9, 28.	2.9	16
25	Circulating microRNAs differentiate Kawasaki Disease from infectious febrile illnesses in childhood. Journal of Molecular and Cellular Cardiology, 2020, 146, 12-18.	1.9	16
26	Improved Detection of Circulating miRNAs in Serum and Plasma Following Rapid Heat/Freeze Cycling. MicroRNA (Shariqah, United Arab Emirates), 2018, 7, 138-147.	1.2	15
27	Cross-species cluster co-conservation: a new method for generating protein interaction networks. Genome Biology, 2007, 8, R185.	9.6	12
28	Integrated analysis of miRNA–mRNA interaction in pediatric dilated cardiomyopathy. Pediatric Research, 2022, 92, 98-108.	2.3	12
29	Elevated serum vascular endothelial growth factor and development of cardiac allograft vasculopathy in children. Journal of Heart and Lung Transplantation, 2018, 37, 1075-1082.	0.6	9
30	R213G polymorphism in SOD3 protects against bleomycin-induced inflammation and attenuates induction of proinflammatory pathways. Physiological Genomics, 2018, 50, 807-816.	2.3	9
31	<i>MAP3K7</i> Loss Drives Enhanced Androgen Signaling and Independently Confers Risk of Recurrence in Prostate Cancer with Joint Loss of <i>CHD1</i> Molecular Cancer Research, 2021, 19, 1123-1136.	3.4	8
32	Serum circulating proteins from pediatric patients with dilated cardiomyopathy cause pathologic remodeling and cardiomyocyte stiffness. JCI Insight, 2021, 6, .	5.0	7
33	A survey of analysis software for array-comparative genomic hybridisation studies to detect copy number variation. Human Genomics, 2010, 4, 421.	2.9	6
34	Amniotic fluid microRNA profiles in twin-twin transfusion syndrome with and without severe recipient cardiomyopathy. American Journal of Obstetrics and Gynecology, 2021, 225, 439.e1-439.e10.	1.3	5
35	Amniotic Fluid microRNA in Severe Twin-Twin Transfusion Syndrome Cardiomyopathy—Identification of Differences and Predicting Demise. Journal of Cardiovascular Development and Disease, 2022, 9, 37.	1.6	2
36	MicroRNA regulation postbleomycin due to the R213G extracellular superoxide dismutase variant is predicted to suppress inflammatory and immune pathways. Physiological Genomics, 2020, 52, 245-254.	2.3	1

#	Article	IF	CITATIONS
37	Circulating cyclic adenosine monophosphate concentrations in milrinone treated paediatric patients after congenital heart surgery. Cardiology in the Young, 2021, 31, 1393-1400.	0.8	1
38	Serum response factor deletion 5 regulates phospholamban phosphorylation and calcium uptake. Journal of Molecular and Cellular Cardiology, 2021, 159, 28-37.	1.9	1
39	Origins of Intraindividual Genetic Variation in Human Fetuses. Reproductive Sciences, 2019, 26, 1139-1145.	2.5	0