Alper Kucukural

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2386507/publications.pdf

Version: 2024-02-01

40 papers 9,243 citations

304743 22 h-index 35 g-index

50 all docs

50 docs citations

50 times ranked

19896 citing authors

#	Article	IF	CITATIONS
1	I-TASSER: a unified platform for automated protein structure and function prediction. Nature Protocols, 2010, 5, 725-738.	12.0	5,594
2	Biogenesis and function of tRNA fragments during sperm maturation and fertilization in mammals. Science, 2016, 351, 391-396.	12.6	992
3	Identification of Neuronal RNA Targets of TDP-43-containing Ribonucleoprotein Complexes. Journal of Biological Chemistry, 2011, 286, 1204-1215.	3.4	366
4	The Cellular EJC Interactome Reveals Higher-Order mRNP Structure and an EJC-SR Protein Nexus. Cell, 2012, 151, 750-764.	28.9	287
5	Novel Observations From Next-Generation RNA Sequencing of Highly Purified Human Adult and Fetal Islet Cell Subsets. Diabetes, 2015, 64, 3172-3181.	0.6	268
6	Simultaneous generation of many RNA-seq libraries in a single reaction. Nature Methods, 2015, 12, 323-325.	19.0	256
7	DEBrowser: interactive differential expression analysis and visualization tool for count data. BMC Genomics, 2019, 20, 6.	2.8	186
8	Ebola Virus Glycoprotein with Increased Infectivity Dominated the 2013–2016 Epidemic. Cell, 2016, 167, 1088-1098.e6.	28.9	173
9	Genetic and Epigenetic Variation, but Not Diet, Shape the Sperm Methylome. Developmental Cell, 2015, 35, 750-758.	7.0	130
10	Staufen1 senses overall transcript secondary structure to regulate translation. Nature Structural and Molecular Biology, 2014, 21, 26-35.	8.2	117
11	Diverse repertoire of human adipocyte subtypes develops from transcriptionally distinct mesenchymal progenitor cells. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 17970-17979.	7.1	106
12	HLA Class II Antigen Processing and Presentation Pathway Components Demonstrated by Transcriptome and Protein Analyses of Islet \hat{l}^2 -Cells From Donors With Type 1 Diabetes. Diabetes, 2019, 68, 988-1001.	0.6	90
13	An improved zebrafish transcriptome annotation for sensitive and comprehensive detection of cell type-specific genes. ELife, 2020, 9, .	6.0	72
14	Dietary suppression of MHC class II expression in intestinal epithelial cells enhances intestinal tumorigenesis. Cell Stem Cell, 2021, 28, 1922-1935.e5.	11.1	67
15	DolphinNext: a distributed data processing platform for high throughput genomics. BMC Genomics, 2020, 21, 310.	2.8	66
16	HIV-1-induced cytokines deplete homeostatic innate lymphoid cells and expand TCF7-dependent memory NK cells. Nature Immunology, 2020, 21, 274-286.	14.5	60
17	Evolutionarily conserved pachytene piRNA loci are highly divergent among modern humans. Nature Ecology and Evolution, 2020, 4, 156-168.	7.8	58
18	An atlas of cell types in the mouse epididymis and vas deferens. ELife, 2020, 9, .	6.0	51

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19	AAV-delivered suppressor tRNA overcomes a nonsense mutation in mice. Nature, 2022, 604, 343-348.	27.8	44
20	ASPeak: an abundance sensitive peak detection algorithm for RIP-Seq. Bioinformatics, 2013, 29, 2485-2486.	4.1	41
21	GUIDEseq: a bioconductor package to analyze GUIDE-Seq datasets for CRISPR-Cas nucleases. BMC Genomics, 2017, 18, 379.	2.8	32
22	Genomic Characterization of Endothelial Enhancers Reveals a Multifunctional Role for NR2F2 in Regulation of Arteriovenous Gene Expression. Circulation Research, 2020, 126, 875-888.	4.5	32
23	Comparative Analysis of Immune Cells Reveals a Conserved Regulatory Lexicon. Cell Systems, 2018, 6, 381-394.e7.	6.2	19
24	Comparison of RNA isolation and associated methods for extracellular RNA detection by high-throughput quantitative polymerase chain reaction. Analytical Biochemistry, 2016, 501, 66-74.	2.4	17
25	Generation and Analysis of Expressed Sequence Tags from <i>Olea europaea</i> L Comparative and Functional Genomics, 2010, 2010, 1-9.	2.0	16
26	Reduced Neurog3 Gene Dosage Shifts Enteroendocrine Progenitor Towards Goblet Cell Lineage in the Mouse Intestine. Cellular and Molecular Gastroenterology and Hepatology, 2021, 11, 433-448.	4.5	12
27	CRISPR-SONIC: targeted somatic oncogene knock-in enables rapid in vivo cancer modeling. Genome Medicine, 2019, 11, 21.	8.2	11
28	Evolutionary selection of minimum number of features for classification of gene expression data using genetic algorithms. , 2007, , .		9
29	Platelet functional and transcriptional changes induced by intralipid infusion. Thrombosis and Haemostasis, 2016, 115, 1147-1156.	3.4	6
30	Recovery of viable endocrineâ€specific cells and transcriptomes from human pancreatic isletâ€engrafted mice. FASEB Journal, 2020, 34, 1901-1911.	0.5	6
31	A Translational Model for Venous Thromboembolism: MicroRNA Expression in Hibernating Black Bears. Journal of Surgical Research, 2021, 257, 203-212.	1.6	6
32	Type I IFN–Driven Immune Cell Dysregulation in Rat Autoimmune Diabetes. ImmunoHorizons, 2021, 5, 855-869.	1.8	4
33	Modified Association Rule Mining Approach for the MHC-Peptide Binding Problem. Lecture Notes in Computer Science, 2006, , 165-173.	1.3	2
34	DISCRIMINATION OF NATIVE FOLDS USING NETWORK PROPERTIES OF PROTEIN STRUCTURES. , 2007, , .		2
35	Bioinformatics Core Survey Highlights the Challenges Facing Data Analysis Facilities. Journal of Biomolecular Techniques, 2020, 31, jbt.20-3102-005.	1.5	2
36	Protein Homology Analysis for Function Prediction with Parallel Sub-Graph Isomorphism., 0,, 386-399.		2

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37	Discrimination of proteins using graph theoretic properties. BMC Systems Biology, 2007, 1, .	3.0	1
38	Discovery of Biomarkers for Hexachlorobenzene Toxicity Using Population Based Methods on Gene Expression Data. Lecture Notes in Computer Science, 2008, , 412-423.	1.3	1
39	mRNAs with a complex: The longâ€ŧerm consequences of a nuclear upbringing. FASEB Journal, 2010, 24, 68.1.	0.5	O
40	Protein Homology Analysis for Function Prediction with Parallel Sub-Graph Isomorphism. , 2011 , , $129-144$.		O