Vincent L Butty

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7742493/publications.pdf

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34 papers 3,048 citations

331670
21
h-index

377865 34 g-index

42 all docs 42 docs citations

42 times ranked 5566 citing authors

#	Article	IF	Citations
1	Braveheart, a Long Noncoding RNA Required for Cardiovascular Lineage Commitment. Cell, 2013, 152, 570-583.	28.9	839
2	Single-cell transcriptomic profiling of the aging mouse brain. Nature Neuroscience, 2019, 22, 1696-1708.	14.8	432
3	Ketone Body Signaling Mediates Intestinal Stem Cell Homeostasis and Adaptation to Diet. Cell, 2019, 178, 1115-1131.e15.	28.9	231
4	Digital genotyping and haplotyping with polymerase colonies. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 5926-5931.	7.1	141
5	Altered Natural Killer Cells in Type 1 Diabetic Patients. Diabetes, 2007, 56, 177-185.	0.6	135
6	Transcriptional Reversion of Cardiac Myocyte Fate During Mammalian Cardiac Regeneration. Circulation Research, 2015, 116, 804-815.	4.5	131
7	Gut-Liver Physiomimetics Reveal Paradoxical Modulation of IBD-Related Inflammation by Short-Chain Fatty Acids. Cell Systems, 2020, 10, 223-239.e9.	6.2	115
8	Multiplexed DNA repair assays for multiple lesions and multiple doses via transcription inhibition and transcriptional mutagenesis. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E1823-32.	7.1	114
9	Transcriptional Analysis of Murine Macrophages Infected with Different Toxoplasma Strains Identifies Novel Regulation of Host Signaling Pathways. PLoS Pathogens, 2013, 9, e1003779.	4.7	111
10	Genomic mapping of phosphorothioates reveals partial modification of short consensus sequences. Nature Communications, 2014, 5, 3951.	12.8	90
11	Methods for time series analysis of RNA-seq data with application to human Th17 cell differentiation. Bioinformatics, 2014, 30, i113-i120.	4.1	62
12	Three <i>Toxoplasma gondii</i> Dense Granule Proteins Are Required for Induction of Lewis Rat Macrophage Pyroptosis. MBio, 2019, 10, .	4.1	59
13	Global transcriptional regulation of innate immunity by ATF-7 in C. elegans. PLoS Genetics, 2019, 15, e1007830.	3.5	56
14	H2A.Z Acidic Patch Couples Chromatin Dynamics to Regulation of Gene Expression Programs during ESC Differentiation. PLoS Genetics, 2013, 9, e1003725.	3.5	53
15	An Adaptable Platform for Directed Evolution in Human Cells. Journal of the American Chemical Society, 2018, 140, 18093-18103.	13.7	52
16	Signatures of strong population differentiation shape extended haplotypes across the human CD28, CTLA4, and ICOS costimulatory genes. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 570-575.	7.1	40
17	Host proteostasis modulates influenza evolution. ELife, 2017, 6, .	6.0	34
18	H3K27me3-mediated silencing of structural genes is required for zebrafish heart regeneration. Development (Cambridge), 2019, 146, .	2.5	33

#	Article	IF	CITATIONS
19	A broad analysis of <i>IL1</i> polymorphism and rheumatoid arthritis. Arthritis and Rheumatism, 2008, 58, 1947-1957.	6.7	31
20	Impact of Diabetes Susceptibility Loci on Progression From Pre-Diabetes to Diabetes in At-Risk Individuals of the Diabetes Prevention Trial–Type 1 (DPT-1). Diabetes, 2008, 57, 2348-2359.	0.6	31
21	Destabilized adaptive influenza variants critical for innate immune system escape are potentiated by host chaperones. PLoS Biology, 2018, 16, e3000008.	5.6	28
22	Galectin-3 Regulates Î ³ -Herpesvirus Specific CD8ÂT Cell Immunity. IScience, 2018, 9, 101-119.	4.1	25
23	Alternative RNA splicing in the endothelium mediated in part by Rbfox2 regulates the arterial response to low flow. ELife, 2018, 7, .	6.0	25
24	Enhanced ER proteostasis and temperature differentially impact the mutational tolerance of influenza hemagglutinin. ELife, 2018, 7, .	6.0	25
25	Geometry-dependent functional changes in iPSC-derived cardiomyocytes probed by functional imaging and RNA sequencing. PLoS ONE, 2017, 12, e0172671.	2.5	23
26	Polycomb Repressive Complex 2 Regulates Lineage Fidelity during Embryonic Stem Cell Differentiation. PLoS ONE, 2014, 9, e110498.	2.5	22
27	Transcriptional and Linkage Analyses Identify Loci that Mediate the Differential Macrophage Response to Inflammatory Stimuli and Infection. PLoS Genetics, 2015, 11, e1005619.	3.5	21
28	A Method for Selective Depletion of Zn(II) Ions from Complex Biological Media and Evaluation of Cellular Consequences of Zn(II) Deficiency. Journal of the American Chemical Society, 2018, 140, 2413-2416.	13.7	19
29	Failed Progenitor Specification Underlies the Cardiopharyngeal Phenotypes in a Zebrafish Model of 22q11.2 Deletion Syndrome. Cell Reports, 2018, 24, 1342-1354.e5.	6.4	18
30	The genetic basis for individual differences in mRNA splicing and APOBEC1 editing activity in murine macrophages. Genome Research, 2014, 24, 377-389.	5.5	13
31	Latent TGF \hat{l}^2 -binding proteins 1 and 3 protect the larval zebrafish outflow tract from aneurysmal dilatation. DMM Disease Models and Mechanisms, 2022, 15, .	2.4	10
32	The endoplasmic reticulum proteostasis network profoundly shapes the protein sequence space accessible to HIV envelope. PLoS Biology, 2022, 20, e3001569.	5.6	7
33	HSF1 Activation Can Restrict HIV Replication. ACS Infectious Diseases, 2020, 6, 1659-1666.	3.8	6
34	High-throughput Minitaturized RNA-Seq Library Preparation. Journal of Biomolecular Techniques, 2020, 31, jbt.20-3104-004.	1.5	5