Christopher Benner

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4606466/publications.pdf

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56 25,386 33 57 papers citations h-index g-index

69 69 69 44168

times ranked

citing authors

docs citations

all docs

#	Article	IF	CITATIONS
1	Simple Combinations of Lineage-Determining Transcription Factors Prime cis-Regulatory Elements Required for Macrophage and B Cell Identities. Molecular Cell, 2010, 38, 576-589.	4.5	10,215
2	Metascape provides a biologist-oriented resource for the analysis of systems-level datasets. Nature Communications, 2019, 10, 1523.	5.8	7,886
3	The selection and function of cell type-specific enhancers. Nature Reviews Molecular Cell Biology, 2015, 16, 144-154.	16.1	859
4	Reprogramming transcription by distinct classes of enhancers functionally defined by eRNA. Nature, 2011, 474, 390-394.	13.7	777
5	Discovery of SARS-CoV-2 antiviral drugs through large-scale compound repurposing. Nature, 2020, 586, 113-119.	13.7	672
6	Remodeling of the Enhancer Landscape during Macrophage Activation Is Coupled to Enhancer Transcription. Molecular Cell, 2013, 51, 310-325.	4.5	616
7	Rev-Erbs repress macrophage gene expression by inhibiting enhancer-directed transcription. Nature, 2013, 498, 511-515.	13.7	480
8	Inflammation-induced IgA+ cells dismantle anti-liver cancer immunity. Nature, 2017, 551, 340-345.	13.7	396
9	Transcription Elongation Can Affect Genome 3D Structure. Cell, 2018, 174, 1522-1536.e22.	13.5	369
10	Global changes in the nuclear positioning of genes and intra- and interdomain genomic interactions that orchestrate B cell fate. Nature Immunology, 2012, 13, 1196-1204.	7.0	249
11	Kdo2-Lipid A of Escherichia coli, a defined endotoxin that activates macrophages via TLR-4. Journal of Lipid Research, 2006, 47, 1097-1111.	2.0	202
12	Identification of Novel Long Noncoding RNAs Underlying Vertebrate Cardiovascular Development. Circulation, 2015, 131, 1278-1290.	1.6	185
13	Regulation of DNA repair pathway choice in S and G2 phases by the NHEJ inhibitor CYREN. Nature, 2017, 549, 548-552.	13.7	184
14	53BP1 and USP28 mediate p53 activation and G1 arrest after centrosome loss or extended mitotic duration. Journal of Cell Biology, 2016, 214, 155-166.	2.3	178
15	Chromatin accessibility underlies synthetic lethality of SWI/SNF subunits in ARID1A-mutant cancers. ELife, 2017, 6, .	2.8	138
16	Functional landscape of SARS-CoV-2 cellular restriction. Molecular Cell, 2021, 81, 2656-2668.e8.	4.5	137
17	Nascent RNA sequencing reveals distinct features in plant transcription. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 12316-12321.	3.3	136
18	Topoisomerase 1 inhibition suppresses inflammatory genes and protects from death by inflammation. Science, 2016, 352, aad7993.	6.0	132

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19	Human Promoters Are Intrinsically Directional. Molecular Cell, 2015, 57, 674-684.	4.5	115
20	IL-17 signaling in steatotic hepatocytes and macrophages promotes hepatocellular carcinoma in alcohol-related liver disease. Journal of Hepatology, 2020, 72, 946-959.	1.8	113
21	Identification and dynamic quantification of regulatory elements using total RNA. Genome Research, 2019, 29, 1836-1846.	2.4	102
22	Heterochromatin-Encoded Satellite RNAs Induce Breast Cancer. Molecular Cell, 2018, 70, 842-853.e7.	4.5	96
23	MHC class II proteins mediate cross-species entry of bat influenza viruses. Nature, 2019, 567, 109-112.	13.7	91
24	Paradoxical association of TET loss of function with genome-wide DNA hypomethylation. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 16933-16942.	3.3	81
25	TOP1 inhibition therapy protects against SARS-CoV-2-induced lethal inflammation. Cell, 2021, 184, 2618-2632.e17.	13.5	80
26	The Dfm1 Derlin Is Required for ERAD Retrotranslocation of Integral Membrane Proteins. Molecular Cell, 2018, 69, 306-320.e4.	4.5	76
27	Genomic Decoding of Neuronal Depolarization by Stimulus-Specific NPAS4 Heterodimers. Cell, 2019, 179, 373-391.e27.	13.5	73
28	Mutant p53 shapes the enhancer landscape of cancer cells in response to chronic immune signaling. Nature Communications, 2017, 8, 754.	5.8	71
29	Senataxin suppresses the antiviral transcriptional response and controls viral biogenesis. Nature Immunology, 2015, 16, 485-494.	7.0	50
30	Diverse motif ensembles specify non-redundant DNA binding activities of AP-1 family members in macrophages. Nature Communications, 2019, 10, 414.	5.8	49
31	ApoC-III ASO promotes tissue LPL activity in the absence of apoE-mediated TRL clearance. Journal of Lipid Research, 2019, 60, 1379-1395.	2.0	48
32	Control of VEGF-A transcriptional programs by pausing and genomic compartmentalization. Nucleic Acids Research, 2014, 42, 12570-12584.	6.5	47
33	Influenza virus repurposes the antiviral protein IFIT2 to promote translation of viral mRNAs. Nature Microbiology, 2020, 5, 1490-1503.	5.9	45
34	Deconvolution of pro- and antiviral genomic responses in Zika virus-infected and bystander macrophages. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E9172-E9181.	3.3	44
35	Massively Parallel Sequencing of Peritoneal and Splenic B Cell Repertoires Highlights Unique Properties of B-1 Cell Antibodies. Journal of Immunology, 2018, 200, 1702-1717.	0.4	36
36	Circular synthesized CRISPR/Cas gRNAs for functional interrogations in the coding and noncoding genome. ELife, 2019, 8, .	2.8	34

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37	Hybrid Gene Origination Creates Human-Virus Chimeric Proteins during Infection. Cell, 2020, 181, 1502-1517.e23.	13.5	33
38	ZNF263 is a transcriptional regulator of heparin and heparan sulfate biosynthesis. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 9311-9317.	3.3	30
39	LSD1-mediated enhancer silencing attenuates retinoic acid signalling during pancreatic endocrine cell development. Nature Communications, 2020, 11, 2082.	5.8	28
40	Restriction factor compendium for influenza A virus reveals a mechanism for evasion of autophagy. Nature Microbiology, 2021, 6, 1319-1333.	5.9	23
41	ARTDeco: automatic readthrough transcription detection. BMC Bioinformatics, 2020, 21, 214.	1.2	21
42	Shear stress associated with cardiopulmonary bypass induces expression of inflammatory cytokines and necroptosis in monocytes. JCI Insight, 2021, 6, .	2.3	20
43	A Genome-Wide CRISPR-Cas9 Screen Reveals the Requirement of Host Cell Sulfation for Schmallenberg Virus Infection. Journal of Virology, 2020, 94, .	1.5	18
44	Ncor2/PPARα-Dependent Upregulation of MCUb in the Type 2 Diabetic Heart Impacts Cardiac Metabolic Flexibility and Function. Diabetes, 2021, 70, 665-679.	0.3	17
45	DNMT3A haploinsufficiency causes dichotomous DNA methylation defects at enhancers in mature human immune cells. Journal of Experimental Medicine, 2021, 218, .	4.2	16
46	MAGGIE: leveraging genetic variation to identify DNA sequence motifs mediating transcription factor binding and function. Bioinformatics, 2020, 36, i84-i92.	1.8	12
47	NCoR1 Protects Mice From Dextran Sodium Sulfate–Induced Colitis by Guarding Colonic Crypt Cells From Luminal Insult. Cellular and Molecular Gastroenterology and Hepatology, 2020, 10, 133-147.	2.3	11
48	Decoding Transcription Regulatory Mechanisms Associated with <i>Coccidioides immitis</i> Phase Transition Using Total RNA. MSystems, 2022, 7, e0140421.	1.7	8
49	A Chinese hamster transcription start site atlas that enables targeted editing of CHO cells. NAR Genomics and Bioinformatics, 2021, 3, Iqab061.	1.5	7
50	Glucocorticoid Receptor-Regulated Enhancers Play a Central Role in the Gene Regulatory Networks Underlying Drug Addiction. Frontiers in Neuroscience, 2022, 16, .	1.4	7
51	A longitudinal systems immunologic investigation of acute Zika virus infection in an individual infected while traveling to Caracas, Venezuela. PLoS Neglected Tropical Diseases, 2018, 12, e0007053.	1.3	6
52	Inner-nuclear-membrane–associated degradation employs Dfm1-independent retrotranslocation and alleviates misfolded transmembrane-protein toxicity. Molecular Biology of the Cell, 2021, 32, 521-537.	0.9	6
53	Wiskott-Aldrich syndrome protein forms nuclear condensates and regulates alternative splicing. Nature Communications, 2022, 13, .	5.8	6
54	MEIRLOP: improving score-based motif enrichment by incorporating sequence bias covariates. BMC Bioinformatics, 2020, 21, 410.	1.2	5

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55	Viral Determinants in H5N1 Influenza A Virus Enable Productive Infection of HeLa Cells. Journal of Virology, 2020, 94, .	1.5	5
56	Negative Regulation of Enhancerâ€Associated RNA in Macrophages. FASEB Journal, 2012, 26, 912.1.	0.2	0