Yong Kong

List of Publications by Year in descending order

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Version: 2024-02-01

102 papers	8,708 citations	C	94433 37 h-index	4	89 g-index
111 all docs	111 docs citations		111 times ranked		17192 citing authors

#	Article	IF	CITATIONS
1	A Protein Interaction Map of Drosophila melanogaster. Science, 2003, 302, 1727-1736.	12.6	2,151
2	Exome sequencing identifies recurrent somatic RAC1 mutations in melanoma. Nature Genetics, 2012, 44, 1006-1014.	21.4	1,052
3	Btrim: A fast, lightweight adapter and quality trimming program for next-generation sequencing technologies. Genomics, 2011, 98, 152-153.	2.9	531
4	Macrophage function in tissue repair and remodeling requires IL-4 or IL-13 with apoptotic cells. Science, 2017, 356, 1072-1076.	12.6	408
5	Exome sequencing identifies recurrent mutations in NF1 and RASopathy genes in sun-exposed melanomas. Nature Genetics, 2015, 47, 996-1002.	21.4	348
6	Integrative Annotation of Variants from 1092 Humans: Application to Cancer Genomics. Science, 2013, 342, 1235587.	12.6	341
7	AlleleSeq: analysis of alleleâ€specific expression and binding in a network framework. Molecular Systems Biology, 2011, 7, 522.	7.2	284
8	Type I interferons instigate fetal demise after Zika virus infection. Science Immunology, 2018, 3, .	11.9	212
9	Microbial Communities of the Upper Respiratory Tract and Otitis Media in Children. MBio, 2011, 2, e00245-10.	4.1	193
10	Negligible impact of rare autoimmune-locus coding-region variants on missing heritability. Nature, 2013, 498, 232-235.	27.8	184
11	Upper Respiratory Tract Microbial Communities, Acute Otitis Media Pathogens, and Antibiotic Use in Healthy and Sick Children. Applied and Environmental Microbiology, 2012, 78, 6262-6270.	3.1	172
12	ERVmap analysis reveals genome-wide transcription of human endogenous retroviruses. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 12565-12572.	7.1	134
13	Substrate Discrimination among Mitogen-activated Protein Kinases through Distinct Docking Sequence Motifs. Journal of Biological Chemistry, 2008, 283, 19511-19520.	3.4	130
14	Glycerol phosphate shuttle enzyme GPD2 regulates macrophage inflammatory responses. Nature Immunology, 2019, 20, 1186-1195.	14.5	126
15	Dynamic Changes in the Streptococcus pneumoniae Transcriptome during Transition from Biofilm Formation to Invasive Disease upon Influenza A Virus Infection. Infection and Immunity, 2014, 82, 4607-4619.	2.2	121
16	Nanoparticles that deliver triplex-forming peptide nucleic acid molecules correct F508del CFTR in airway epithelium. Nature Communications, 2015, 6, 6952.	12.8	114
17	Aging impairs both primary and secondary RIG-I signaling for interferon induction in human monocytes. Science Signaling, 2017, 10, .	3.6	113
18	Crystal structures of the Klenow fragment of <i>Thermus aquaticus</i> DNA polymerase I complexed with deoxyribonucleoside triphosphates. Protein Science, 1998, 7, 1116-1123.	7.6	102

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19	An international effort towards developing standards for best practices in analysis, interpretation and reporting of clinical genome sequencing results in the CLARITY Challenge. Genome Biology, 2014, 15, R53.	9.6	101
20	Ketogenic diet activates protective $\hat{I}^3\hat{I}^{'}T$ cell responses against influenza virus infection. Science Immunology, 2019, 4, .	11.9	98
21	Analysis of genomic variation in non-coding elements using population-scale sequencing data from the 1000 Genomes Project. Nucleic Acids Research, 2011, 39, 7058-7076.	14.5	81
22	A uniform survey of allele-specific binding and expression over 1000-Genomes-Project individuals. Nature Communications, 2016, 7, 11101.	12.8	78
23	Analysis of gene–environment interactions in postnatal development of the mammalian intestine. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 1929-1936.	7.1	77
24	Dolosigranulum pigrum Cooperation and Competition in Human Nasal Microbiota. MSphere, 2020, 5, .	2.9	65
25	Rare SF3B1 R625 mutations in cutaneous melanoma. Melanoma Research, 2014, 24, 332-334.	1.2	64
26	The Lupus Susceptibility Locus Sgp3 Encodes the Suppressor of Endogenous Retrovirus Expression SNERV. Immunity, 2019, 50, 334-347.e9.	14.3	61
27	Alleles of a Polymorphic ETV6 Binding Site in DCDC2 Confer Risk of Reading and Language Impairment. American Journal of Human Genetics, 2013, 93, 19-28.	6.2	60
28	Viral Spread to Enteric Neurons Links Genital HSV-1 Infection to Toxic Megacolon and Lethality. Cell Host and Microbe, 2016, 19, 788-799.	11.0	58
29	<i>Haemophilus influenzae</i> genome evolution during persistence in the human airways in chronic obstructive pulmonary disease. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E3256-E3265.	7.1	57
30	Calculation of the reaction field due to off-center point multipoles. Journal of Chemical Physics, 1997, 107, 481-492.	3.0	55
31	Intratumoral delivery of RIG-I agonist SLR14 induces robust antitumor responses. Journal of Experimental Medicine, 2019, 216, 2854-2868.	8.5	49
32	Regional Differences in Airway Epithelial Cells Reveal Tradeoff between Defense against Oxidative Stress and Defense against Rhinovirus. Cell Reports, 2018, 24, 3000-3007.e3.	6.4	46
33	Type 3 Inositol 1,4,5â€Trisphosphate Receptor Is Increased and Enhances Malignant Properties in Cholangiocarcinoma. Hepatology, 2020, 71, 583-599.	7. 3	45
34	Forward genetics defines Xylt1 as a key, conserved regulator of early chondrocyte maturation and skeletal length. Developmental Biology, 2014, 385, 67-82.	2.0	44
35	Association of sputum microbiota profiles with severity of community-acquired pneumonia in children. BMC Infectious Diseases, 2016, 16, 317.	2.9	44
36	Gastrointestinal Microbiota Disruption and Risk of Colonization With Carbapenem-resistant Pseudomonas aeruginosa in Intensive Care Unit Patients. Clinical Infectious Diseases, 2019, 69, 604-613.	5.8	43

3

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37	Heterologous quaternary structure of CXCL12 and its relationship to the CC chemokine family. Proteins: Structure, Function and Bioinformatics, 2010, 78, 1331-1337.	2.6	40
38	Forward genetics uncovers Transmembrane protein 107 as a novel factor required for ciliogenesis and Sonic hedgehog signaling. Developmental Biology, 2012, 368, 382-392.	2.0	40
39	MicroRNA: Biological and Computational Perspective. Genomics, Proteomics and Bioinformatics, 2005, 3, 62-72.	6.9	39
40	Theory of multivalent binding in one and two-dimensional lattices. Biophysical Chemistry, 1996, 61, 107-124.	2.8	36
41	Fluorescence Anisotropy Studies on the Ku-DNA Interaction. Journal of Biological Chemistry, 2004, 279, 42826-42835.	3.4	35
42	Antimicrobial activity of antisense peptide–peptide nucleic acid conjugates against non-typeable <i>Haemophilus influenzae</i> in planktonic and biofilm forms. Journal of Antimicrobial Chemotherapy, 2017, 72, 137-144.	3.0	33
43	Forward genetics identifies Kdf1/1810019J16Rik as an essential regulator of the proliferation–differentiation decision in epidermal progenitor cells. Developmental Biology, 2013, 383, 201-213.	2.0	31
44	Desynchronization of the molecular clock contributes to the heterogeneity of the inflammatory response. Science Signaling, 2019, 12, .	3.6	30
45	Expression of IgA Proteases by <i>Haemophilus influenzae </i> in the Respiratory Tract of Adults With Chronic Obstructive Pulmonary Disease. Journal of Infectious Diseases, 2015, 212, 1798-1805.	4.0	29
46	The regulatory element READ1 epistatically influences reading and language, with both deleterious and protective alleles. Journal of Medical Genetics, 2016, 53, 163-171.	3.2	29
47	A phase II evaluation of pembrolizumab in recurrent microsatellite instability-high (MSI-H) endometrial cancer patients with Lynch-like versus MLH-1 methylated characteristics (NCT02899793). Annals of Oncology, 2021, 32, 1045-1046.	1.2	29
48	A phase 2 evaluation of pembrolizumab for recurrent Lynchâ€like versus sporadic endometrial cancers with microsatellite instability. Cancer, 2022, 128, 1206-1218.	4.1	28
49	Packing dimers on(2p+1)×(2q+1)lattices. Physical Review E, 2006, 73, 016106.	2.1	26
50	Effect of Fluoroquinolones and Macrolides on Eradication and Resistance of Haemophilus influenzae in Chronic Obstructive Pulmonary Disease. Antimicrobial Agents and Chemotherapy, 2016, 60, 4151-4158.	3.2	26
51	Antibodies against human endogenous retrovirus K102 envelope activate neutrophils in systemic lupus erythematosus. Journal of Experimental Medicine, 2021, 218, .	8.5	26
52	IFT56 regulates vertebrate developmental patterning by maintaining IFTB integrity and ciliary microtubule architecture. Development (Cambridge), 2017, 144, 1544-1553.	2.5	25
53	Distribution of Runs and Longest Runs. Journal of the American Statistical Association, 2006, 101, 1253-1263.	3.1	23
54	Logarithmic corrections in the free energy of monomer-dimer model on plane lattices with free boundaries. Physical Review E, 2006, 74, 011102.	2.1	22

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55	Monomer-dimer model in two-dimensional rectangular lattices with fixed dimer density. Physical Review E, 2006, 74, 061102.	2.1	20
56	A Simple Method for Evaluating Partition Functions of Linear Polymers. Journal of Physical Chemistry B, 2001, 105, 10111-10114.	2.6	16
57	Exon capture and bulk segregant analysis: rapid discovery of causative mutations using high-throughput sequencing. BMC Genomics, 2012, 13, 649.	2.8	16
58	Streptococcus pneumoniae Clonal Complex 199: Genetic Diversity and Tissue-Specific Virulence. PLoS ONE, 2011, 6, e18649.	2.5	16
59	Changes in IgA Protease Expression Are Conferred by Changes in Genomes during Persistent Infection by Nontypeable Haemophilus influenzae in Chronic Obstructive Pulmonary Disease. Infection and Immunity, 2018, 86, .	2.2	15
60	Human APOBEC3G Prevents Emergence of Infectious Endogenous Retrovirus in Mice. Journal of Virology, 2019, 93, .	3.4	15
61	Exosomal MicroRNAs Associate With Neuropsychological Performance in Individuals With HIV Infection on Antiretroviral Therapy. Journal of Acquired Immune Deficiency Syndromes (1999), 2019, 82, 514-522.	2.1	15
62	Integrating Sequencing Technologies in Personal Genomics: Optimal Low Cost Reconstruction of Structural Variants. PLoS Computational Biology, 2009, 5, e1000432.	3.2	14
63	Generalized Correlation Functions and Their Applications in Selection of Optimal Multiple Spaced Seeds for Homology Search. Journal of Computational Biology, 2007, 14, 238-254.	1.6	10
64	Adjusting for Background Mutation Frequency Biases Improves the Identification of Cancer Driver Genes. IEEE Transactions on Nanobioscience, 2013, 12, 150-157.	3.3	10
65	General recurrence theory of ligand binding on a three-dimensional lattice. Journal of Chemical Physics, 1999, 111, 4790-4799.	3.0	9
66	Differential immunomodulatory effect of PARP inhibition in BRCA1 deficient and competent tumor cells. Biochemical Pharmacology, 2021, 184, 114359.	4.4	8
67	Extension of Goulden–Jackson cluster method on pattern occurrences in random sequences and comparison with Régnier–Szpankowski method. Journal of Difference Equations and Applications, 2005, 11, 1265-1271.	1.1	7
68	MU2Aâ€"reconciling the genome and transcriptome to determine the effects of base substitutions. Bioinformatics, 2011, 27, 416-418.	4.1	7
69	Distributions of Runs Revisited. Communications in Statistics - Theory and Methods, 2015, 44, 4663-4678.	1.0	7
70	Number of appearances of events in random sequences: a new generating function approach to Type II and Type III runs. Annals of the Institute of Statistical Mathematics, 2017, 69, 489-495.	0.8	7
71	APOBEC3A regulates transcription from interferon-stimulated response elements. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e2011665119.	7.1	7
72	Calculating complexity of large randomized libraries. Journal of Theoretical Biology, 2009, 259, 641-645.	1.7	6

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73	Construction and maintenance of randomized retroviral expression libraries for transmembrane protein engineering. Protein Engineering, Design and Selection, 2011, 24, 311-320.	2.1	6
74	Number of appearances of events in random sequences: A new approach to non-overlapping runs. Communications in Statistics - Theory and Methods, 2016, 45, 6765-6772.	1.0	6
75	Joint distribution of rises, falls, and number of runs in random sequences. Communications in Statistics - Theory and Methods, 2019, 48, 493-499.	1.0	6
76	RUNX Binding Sites Are Enriched in Herpesvirus Genomes, and RUNX1 Overexpression Leads to Herpes Simplex Virus 1 Suppression. Journal of Virology, 2020, 94, .	3.4	6
77	Gastrointestinal Microbiome Disruption and Antibiotic-Associated Diarrhea in Children Receiving Antibiotic Therapy for Community-Acquired Pneumonia. Journal of Infectious Diseases, 2022, 226, 1109-1119.	4.0	6
78	Asymptotics of the monomer-dimer model on two-dimensional semi-infinite lattices. Physical Review E, 2007, 75, 051123.	2.1	5
79	Note: Solution of one-dimensional self-assembly problem in multiple-component systems with correction of overcounting of non-symmetrical complexes. Journal of Chemical Physics, 2011, 135, 156101.	3.0	5
80	The mth longest runs of multivariate random sequences. Annals of the Institute of Statistical Mathematics, 2017, 69, 497-512.	0.8	5
81	Decoupling Combinatorial Complexity: a Two-Step Approach to Distributions of Runs. Methodology and Computing in Applied Probability, 2019, 21, 789-803.	1.2	5
82	A phase II evaluation of pembrolizumab in recurrent microsatellite instability-high (MSI-H) endometrial cancer patients with Lynch-like versus <i>MLH</i> -1 methylated characteristics (NCT02899793) Journal of Clinical Oncology, 2021, 39, 5523-5523.	1.6	5
83	Bacterial Signatures of Paediatric Respiratory Disease: An Individual Participant Data Meta-Analysis. Frontiers in Microbiology, 2021, 12, 711134.	3.5	5
84	Ligand binding on ladder lattices. Biophysical Chemistry, 1999, 81, 7-21.	2.8	4
85	Statistical Distributions of Pyrosequencing. Journal of Computational Biology, 2009, 16, 31-42.	1.6	4
86	Statistical Distributions of Sequencing by Synthesis with Probabilistic Nucleotide Incorporation. Journal of Computational Biology, 2009, 16, 817-827.	1.6	4
87	Streptococcus pneumoniae Serotype 6C: an Intra- and Interclonal Complex Comparison. Journal of Bacteriology, 2011, 193, 3409-3410.	2.2	4
88	A note on the quantitative properties of McGhee–von Hippel model. Biophysical Chemistry, 2002, 95, 1-6.	2.8	3
89	Keck Foundation Biotechnology Resource Laboratory, Yale University. Yale Journal of Biology and Medicine, 2007, 80, 195-211.	0.2	3
90	The Expectation of Sample Central Moments. American Statistician, 2011, 65, 198-199.	1.6	2

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91	Generating Function Methods for Run and Scan Statistics. , 2019, , 1-19.		2
92	Reply to Iñiguez et al.: ERVmap is a validated approach to mapping proviral endogenous retroviruses in the human genome. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 21352-21353.	7.1	2
93	Distributions of successions of arbitrary multisets. Communications in Statistics - Theory and Methods, 2022, 51, 1693-1705.	1.0	2
94	Multiple consecutive runs of multi-state trials: Distributions of <mml:math <="" display="inline" id="d1e22" td="" xmlns:mml="http://www.w3.org/1998/Math/MathML"><td></td><td></td></mml:math>		