

Chew Yee Ngan

List of Publications by Year in descending order

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66
papers

5,813
citations

117453

34
h-index

114278

63
g-index

71
all docs

71
docs citations

71
times ranked

10412
citing authors

#	ARTICLE	IF	CITATIONS
1	Oncogenic extrachromosomal DNA functions as mobile enhancers to globally amplify chromosomal transcription. <i>Cancer Cell</i> , 2021, 39, 694-707.e7.	7.7	115
2	Abstract 2084: Single-cell multimodal glioma analyses reveal epigenetic regulators of cellular plasticity and environmental stress response. , 2021, , .		0
3	Reduced subgenomic RNA expression is a molecular indicator of asymptomatic SARS-CoV-2 infection. <i>Communications Medicine</i> , 2021, 1, .	1.9	13
4	Single-cell multimodal glioma analyses identify epigenetic regulators of cellular plasticity and environmental stress response. <i>Nature Genetics</i> , 2021, 53, 1456-1468.	9.4	111
5	Bioinformatics Tools for PacBio Sequenced Amplicon Data Pre-processing and Target Sequence Extraction. <i>Lecture Notes in Networks and Systems</i> , 2020, , 326-340.	0.5	0
6	Chromatin interaction analyses elucidate the roles of PRC2-bound silencers in mouse development. <i>Nature Genetics</i> , 2020, 52, 264-272.	9.4	104
7	Comparative Molecular Life History of Spontaneous Canine and Human Gliomas. <i>Cancer Cell</i> , 2020, 37, 243-257.e7.	7.7	59
8	Succession of physiological stages hallmarks the transcriptomic response of the fungus <i>Aspergillus niger</i> to lignocellulose. <i>Biotechnology for Biofuels</i> , 2020, 13, 69.	6.2	4
9	Mapping the Global Chromatin Connectivity Network for Sox2 Function in Neural Stem Cell Maintenance. <i>Cell Stem Cell</i> , 2019, 24, 462-476.e6.	5.2	72
10	Ultra-long Read Sequencing for Whole Genomic DNA Analysis. <i>Journal of Visualized Experiments</i> , 2019, , .	0.2	20
11	Multiplex chromatin interactions with single-molecule precision. <i>Nature</i> , 2019, 566, 558-562.	13.7	180
12	GENE-57. COMPARATIVE MOLECULAR LIFE HISTORY OF SPONTANEOUS CANINE AND HUMAN GLIOMA. <i>Neuro-Oncology</i> , 2019, 21, vi110-vi110.	0.6	0
13	Longitudinal molecular trajectories of diffuse glioma in adults. <i>Nature</i> , 2019, 576, 112-120.	13.7	320
14	Linked-read Sequencing Analysis Reveals Tumor-specific Genome Variation Landscapes in Neurofibromatosis Type 2 (NF2) Patients. <i>Otology and Neurotology</i> , 2019, 40, e150-e159.	0.7	3
15	Linking secondary metabolites to gene clusters through genome sequencing of six diverse <i>Aspergillus</i> species. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E753-E761.	3.3	126
16	Picky comprehensively detects high-resolution structural variants in nanopore long reads. <i>Nature Methods</i> , 2018, 15, 455-460.	9.0	80
17	<i>Sox2</i> conditional mutation in mouse causes ataxic symptoms, cerebellar vermis hypoplasia, and postnatal defects of <i>Bergmann</i> glia. <i>Glia</i> , 2018, 66, 1929-1946.	2.5	28
18	Comparative genomics reveals high biological diversity and specific adaptations in the industrially and medically important fungal genus <i>Aspergillus</i> . <i>Genome Biology</i> , 2017, 18, 28.	3.8	417

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19	<scp>DNA</scp> methylation and gene expression regulation associated with vascularization in <i>Sorghum bicolor</i>. <i>New Phytologist</i> , 2017, 214, 1213-1229.	3.5	47
20	Complete Genome Sequence of <i>Nitrosomonas cryotolerans</i> ATCC 49181, a Phylogenetically Distinct Ammonia-Oxidizing Bacterium Isolated from Arctic Waters. <i>Genome Announcements</i> , 2017, 5, .	0.8	3
21	Expression of <i>Aspergillus niger</i> CAZymes is determined by compositional changes in wheat straw generated by hydrothermal or ionic liquid pretreatments. <i>Biotechnology for Biofuels</i> , 2017, 10, 35.	6.2	18
22	Genome Sequence of <i>Roseovarius</i> sp. Strain MCTG156(2b) Isolated from a Phytoplankton Net Trawl on the Scottish West Coast. <i>Genome Announcements</i> , 2017, 5, .	0.8	1
23	Genome Sequence of <i>Oceanicola</i> sp. Strain MCTG156(1a), Isolated from a Scottish Coastal Phytoplankton Net Sample. <i>Genome Announcements</i> , 2017, 5, .	0.8	3
24	Genome Sequence of <i>Arenibacter algicola</i> Strain TG409, a Hydrocarbon-Degrading Bacterium Associated with Marine Eukaryotic Phytoplankton. <i>Genome Announcements</i> , 2016, 4, .	0.8	8
25	High-Quality Draft Genomes from <i>Thermus caliditerrae</i> YIM 77777 and <i>T.Âtengchongensis</i> YIM 77401, Isolates from Tengchong, China. <i>Genome Announcements</i> , 2016, 4, .	0.8	5
26	Application of Long Sequence Reads To Improve Genomes for <i>Clostridium thermocellum</i> AD2, <i>Clostridium thermocellum</i> LQRI, and <i>Pelosinus fermentans</i> R7. <i>Genome Announcements</i> , 2016, 4, .	0.8	2
27	Permanent draft genome of <i>Thermithiobacillus tepidarius</i> DSM 3134T, a moderately thermophilic, obligately chemolithoautotrophic member of the Acidithiobacillia. <i>Standards in Genomic Sciences</i> , 2016, 11, 74.	1.5	15
28	Genome Sequence of <i>Marinobacter</i> sp. Strain MCTG268 Isolated from the Cosmopolitan Marine Diatom <i>Skeletonema costatum</i> . <i>Genome Announcements</i> , 2016, 4, .	0.8	1
29	Complete genome of <i>Nitrosospora briensis</i> C-128, an ammonia-oxidizing bacterium from agricultural soil. <i>Standards in Genomic Sciences</i> , 2016, 11, 46.	1.5	22
30	Integrative epigenomic analysis reveals unique epigenetic signatures involved in unipotency of mouse female germline stem cells. <i>Genome Biology</i> , 2016, 17, 162.	3.8	61
31	Near-Complete Genome Sequence of <i>Thalassospora</i> sp. Strain KO164 Isolated from a Lignin-Enriched Marine Sediment Microcosm. <i>Genome Announcements</i> , 2016, 4, .	0.8	1
32	High-Quality Draft Genome Sequence of <i>Thermocrinis jamiesonii</i> GBS1 ^T Isolated from Great Boiling Spring, Nevada. <i>Genome Announcements</i> , 2016, 4, .	0.8	0
33	High-quality draft genome sequence of the <i>Thermus amyloliquefaciens</i> type strain YIM 77409T with an incomplete denitrification pathway. <i>Standards in Genomic Sciences</i> , 2016, 11, 20.	1.5	7
34	Diversity and population structure of northern switchgrass as revealed through exome capture sequencing. <i>Plant Journal</i> , 2015, 84, 800-815.	2.8	47
35	Lineage-specific chromatin signatures reveal a regulator of lipid metabolism in microalgae. <i>Nature Plants</i> , 2015, 1, 15107.	4.7	89
36	Impact of library preparation protocols and template quantity on the metagenomic reconstruction of a mock microbial community. <i>BMC Genomics</i> , 2015, 16, 856.	1.2	79

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37	High-Quality Draft Genome Sequence of <i>Kallotenucia paperylyticum</i> JKG1 T Reveals Broad Heterotrophic Capacity Focused on Carbohydrate and Amino Acid Metabolism. <i>Genome Announcements</i> , 2015, 3, .	0.8	4
38	Phylogenomic Analyses Indicate that Early Fungi Evolved Digesting Cell Walls of Algal Ancestors of Land Plants. <i>Genome Biology and Evolution</i> , 2015, 7, 1590-1601.	1.1	175
39	Genome Sequence of <i>Polycyclovorans algicola</i> Strain TG408, an Obligate Polycyclic Aromatic Hydrocarbon-Degrading Bacterium Associated with Marine Eukaryotic Phytoplankton. <i>Genome Announcements</i> , 2015, 3, .	0.8	4
40	Genome Sequence of <i>Porticoccus hydrocarbonoclasticus</i> Strain MCTG13d, an Obligate Polycyclic Aromatic Hydrocarbon-Degrading Bacterium Associated with Marine Eukaryotic Phytoplankton. <i>Genome Announcements</i> , 2015, 3, .	0.8	9
41	Genome Sequence of <i>Halomonas</i> sp. Strain MCTG39a, a Hydrocarbon-Degrading and Exopolymeric Substance-Producing Bacterium. <i>Genome Announcements</i> , 2015, 3, .	0.8	6
42	Convergent losses of decay mechanisms and rapid turnover of symbiosis genes in mycorrhizal mutualists. <i>Nature Genetics</i> , 2015, 47, 410-415.	9.4	870
43	Strand-Specific RNA-Seq Analyses of Fruiting Body Development in <i>Coprinopsis cinerea</i> . <i>PLoS ONE</i> , 2015, 10, e0141586.	1.1	95
44	Genome sequencing of four <i>Aureobasidium pullulans</i> varieties: biotechnological potential, stress tolerance, and description of new species. <i>BMC Genomics</i> , 2014, 15, 549.	1.2	262
45	Marine algae and land plants share conserved phytochrome signaling systems. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 15827-15832.	3.3	108
46	Chromatin connectivity maps reveal dynamic promoter-enhancer long-range associations. <i>Nature</i> , 2013, 504, 306-310.	13.7	405
47	Inhibition of angiopoietin 2 attenuates lumen formation of tumour-associated vessels in vivo. <i>International Journal of Oncology</i> , 2013, 43, 1447-1455.	1.4	9
48	CTCF-mediated functional chromatin interactome in pluripotent cells. <i>Nature Genetics</i> , 2011, 43, 630-638.	9.4	567
49	Cancer cells survive with survivin. <i>Cancer Science</i> , 2008, 99, 1709-1714.	1.7	150
50	Overexpression of Tyrosine Kinase B Protein as a Predictor for Distant Metastases and Prognosis in Gastric Carcinoma. <i>Oncology</i> , 2008, 75, 17-26.	0.9	31
51	Aberrant Expression of Connexin 26 Is Associated with Lung Metastasis of Colorectal Cancer. <i>Clinical Cancer Research</i> , 2008, 14, 677-684.	3.2	72
52	CDC25A inhibition suppresses the growth and invasion of human hepatocellular carcinoma cells. <i>International Journal of Molecular Medicine</i> , 2008, 21, 145-52.	1.8	14
53	Overexpression of connexin 26 in carcinoma of the pancreas. <i>Oncology Reports</i> , 2008, 19, 627-31.	1.2	18
54	Stromal Myofibroblasts Predict Disease Recurrence for Colorectal Cancer. <i>Clinical Cancer Research</i> , 2007, 13, 2082-2090.	3.2	305

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55	A multivariate analysis of adhesion molecules expression in assessment of colorectal cancer. <i>Journal of Surgical Oncology</i> , 2007, 95, 652-662.	0.8	75
56	Quantitative evaluation of vimentin expression in tumour stroma of colorectal cancer. <i>British Journal of Cancer</i> , 2007, 96, 986-992.	2.9	123
57	Oxaliplatin induces mitotic catastrophe and apoptosis in esophageal cancer cells. <i>Cancer Science</i> , 2007, 99, 071019192917002-???	1.7	46
58	Expression of PPAR γ in multistage carcinogenesis of the colorectum: implications of malignant cancer morphology. <i>British Journal of Cancer</i> , 2006, 95, 889-895.	2.9	62
59	Construction of a novel DNA decoy that inhibits the oncogenic β -catenin/T-cell factor pathway. <i>Molecular Cancer Therapeutics</i> , 2006, 5, 985-994.	1.9	19
60	Low-Dose Oxaliplatin Enhances the Antitumor Efficacy of Paclitaxel in Human Gastric Cancer Cell Lines. <i>Digestion</i> , 2006, 74, 19-27.	1.2	17
61	Antisense to Cyclin D1 Inhibits Vascular Endothelial Growth Factor- α -Stimulated Growth of Vascular Endothelial Cells: Implication of Tumor Vascularization. <i>Clinical Cancer Research</i> , 2006, 12, 4720-4729.	3.2	60
62	Hypoxia-induced up-regulation of angiopoietin-2 in colorectal cancer. <i>Oncology Reports</i> , 2006, 15, 779-83.	1.2	18
63	Oxaliplatin, a Potent Inhibitor of Survivin, Enhances Paclitaxel-induced Apoptosis and Mitotic Catastrophe in Colon Cancer Cells. <i>Japanese Journal of Clinical Oncology</i> , 2005, 35, 453-463.	0.6	63
64	Role of p21waf1/cip1 in effects of oxaliplatin in colorectal cancer cells. <i>Molecular Cancer Therapeutics</i> , 2005, 4, 1585-1594.	1.9	37
65	Hepatic expression of ANG2 RNA in metastatic colorectal cancer. <i>Hepatology</i> , 2004, 39, 528-539.	3.6	46
66	Overexpression of CDC25A phosphatase is associated with hypergrowth activity and poor prognosis of human hepatocellular carcinomas. <i>Clinical Cancer Research</i> , 2003, 9, 1764-72.	3.2	68