Shifu Chen

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/382585/publications.pdf

Version: 2024-02-01

40 papers

11,917 citations

623734 14 h-index 330143 37 g-index

43 all docs 43 docs citations

43 times ranked

13550 citing authors

#	Article	IF	CITATIONS
1	fastp: an ultra-fast all-in-one FASTQ preprocessor. Bioinformatics, 2018, 34, i884-i890.	4.1	11,005
2	AfterQC: automatic filtering, trimming, error removing and quality control for fastq data. BMC Bioinformatics, 2017, 18, 80.	2.6	286
3	Gencore: an efficient tool to generate consensus reads for error suppressing and duplicate removing of NGS data. BMC Bioinformatics, 2019, 20, 606.	2.6	43
4	E-health in China: Challenges, Initial Directions, and Experience. Telemedicine Journal and E-Health, 2010, 16, 344-349.	2.8	41
5	GeneFuse: detection and visualization of target gene fusions from DNA sequencing data. International Journal of Biological Sciences, 2018, 14, 843-848.	6.4	36
6	Detection of Urothelial Bladder Carcinoma via Microfluidic Immunoassay and Single-Cell DNA Copy-Number Alteration Analysis of Captured Urinary-Exfoliated Tumor Cells. Cancer Research, 2018, 78, 4073-4085.	0.9	34
7	Comprehensive analysis of POLE and POLD1 Gene Variations identifies cancer patients potentially benefit from immunotherapy in Chinese population. Scientific Reports, 2019, 9, 15767.	3.3	34
8	A computational toolset for rapid identification of SARS-CoV-2, other viruses and microorganisms from sequencing data. Briefings in Bioinformatics, 2021, 22, 924-935.	6.5	34
9	The contribution of hereditary cancer-related germline mutations to lung cancer susceptibility. Translational Lung Cancer Research, 2020, 9, 646-658.	2.8	33
10	Monitoring colorectal cancer following surgery using plasma circulating tumor DNA. Oncology Letters, 2018, 15, 4365-4375.	1.8	24
11	The Genomic Characteristics of ALK Fusion Positive Tumors in Chinese NSCLC Patients. Frontiers in Oncology, 2020, 10, 726.	2.8	21
12	Comprehensive profiling of 1015 patients' exomes reveals genomic-clinical associations in colorectal cancer. Nature Communications, 2022, 13, 2342.	12.8	21
13	An adaptive kernel regression method for 3D ultrasound reconstruction using speckle prior and parallel GPU implementation. Neurocomputing, 2018, 275, 208-223.	5.9	17
14	ctDNA Concentration, MIKI67 Mutations and Hyper-Progressive Disease Related Gene Mutations Are Prognostic Markers for Camrelizumab and Apatinib Combined Multiline Treatment in Advanced NSCLC. Frontiers in Oncology, 2020, 10, 1706.	2.8	17
15	Genomic temporal heterogeneity of circulating tumour DNA in unresectable metastatic colorectal cancer under first-line treatment. Gut, 2022, 71, 1340-1349.	12.1	17
16	Segmentation of abdomen MR images using kernel graph cuts with shape priors. BioMedical Engineering OnLine, 2013, 12, 124.	2.7	15
17	Bioinformatics Analysis for Cell-Free Tumor DNA Sequencing Data. Methods in Molecular Biology, 2018, 1754, 67-95.	0.9	15
18	MutScan: fast detection and visualization of target mutations by scanning FASTQ data. BMC Bioinformatics, 2018, 19, 16.	2.6	15

#	Article	IF	CITATIONS
19	Mutational Profile Evaluates Response and Survival to Firstâ€Line Chemotherapy in Lung Cancer. Advanced Science, 2021, 8, 2003263.	11.2	14
20	A Fast and Flexible Sorting Algorithm with CUDA. Lecture Notes in Computer Science, 2009, , 281-290.	1.3	12
21	Development and Recent Achievements of Telemedicine in China. Telemedicine Journal and E-Health, 2010, 16, 634-638.	2.8	11
22	Targeted deep sequencing helps distinguish independent primary tumors from intrapulmonary metastasis for lung cancer diagnosis. Journal of Cancer Research and Clinical Oncology, 2020, 146, 2359-2367.	2.5	9
23	Accelerating Algebraic Reconstruction Using CUDA-Enabled GPU., 2009, , .		7
24	CUDA-based acceleration and algorithm refinement for volume image registration. , 2009, , .		7
25	An efficient sorting algorithm with CUDA. Journal of the Chinese Institute of Engineers, Transactions of the Chinese Institute of Engineers, Series A/Chung-kuo Kung Ch'eng Hsuch K'an, 2009, 32, 915-921.	1.1	7
26	Sequencing of circulating tumor DNA for dynamic monitoring of gene mutations in advanced non-small cell lung cancer. Oncology Letters, 2018, 15, 3726-3734.	1.8	6
27	Metastatic Low-Grade Sarcoma with <i>CARS</i> - <i>ALK</i> Fusion Dramatically Responded to Multiple ALK Tyrosine Kinase Inhibitors: A Case Report with Comprehensive Genomic Analysis. Oncologist, 2021, 26, e524-e529.	3.7	6
28	Identification of a novel germline frameshift mutation p.D300fs of PMS1 in a patient with hepatocellular carcinoma. Medicine (United States), 2020, 99, e19076.	1.0	6
29	E-Health–Oriented Community Health Information System in China: Our Challenges, Solution, and Experience. Telemedicine Journal and E-Health, 2011, 17, 584-588.	2.8	5
30	Circulating tumor DNA sequencing for colorectal cancers: A comparative analysis of colon cancer and rectal cancer data. Cancer Biomarkers, 2019, 26, 313-322.	1.7	5
31	A Study of Cell-free DNA Fragmentation Pattern and Its Application in DNA Sample Type Classification. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2018, , 1-1.	3.0	4
32	The Design and Implementation of a Rescue Terminal with Vital Signs Telemonitoring Based on Beidou 1 Navigation Satellite System. Telemedicine Journal and E-Health, 2011, 17, 76-79.	2.8	2
33	The Introduction and Clinical Application of Cell-Free Tumor DNA. Methods in Molecular Biology, 2018, 1754, 45-65.	0.9	2
34	A rare multiple primary sarcomatoid carcinoma (SCA) of small intestine harboring driver gene mutations: a case report and a literature review. Translational Cancer Research, 2021, 10, 1150-1161.	1.0	2
35	The genomic characteristics of RET fusion positive tumors in Chinese non-small cell lung cancer (NSCLC) patients. Journal of Cancer Research and Clinical Oncology, 2023, 149, 1019-1028.	2.5	2
36	A Comparison of Two Data Storage Strategies for Implementing e-Health in China. Telemedicine Journal and E-Health, 2010, 16, 364-369.	2.8	1

SHIFU CHEN

#	Article	IF	CITATIONS
37	Case Report: A Chronological Combination Treatment of Icotinib, Osimertinib, and Crizotinib on Lung Adenocarcinoma Guided by Serial Genetic Tests of Circulating Tumor DNA and Sediment Cell Genomic DNA From Pleural Effusion. Frontiers in Oncology, 2020, 10, 561341.	2.8	1
38	Identification of Genomic Alterations in Sporadic Pancreatic Neurogenic Tumors. Pancreas, 2020, 49, 1393-1397.	1.1	1
39	Detect and visualize tumor microsatellite instability status from next-generation sequencing data by simulating PCR techniques Journal of Clinical Oncology, 2019, 37, e13052-e13052.	1.6	1
40	SeqMaker: A next generation sequencing simulator with variations, sequencing errors and amplification bias integrated. , 2016, , .		0