Chuanpeng Dong

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9643624/publications.pdf

Version: 2024-02-01

22 papers 356 citations

933447 10 h-index 17 g-index

25 all docs

25 docs citations

25 times ranked

440 citing authors

#	Article	IF	Citations
1	Diagnostic Evidence GAuge of Single cells (DEGAS): a flexible deep transfer learning framework for prioritizing cells in relation to disease. Genome Medicine, 2022, 14, 11.	8.2	10
2	Intron-Retention Neoantigen Load Predicts Favorable Prognosis in Pancreatic Cancer. JCO Clinical Cancer Informatics, 2022, 6, e2100124.	2.1	6
3	PLCG2 is associated with the inflammatory response and is induced by amyloid plaques in Alzheimer's disease. Genome Medicine, 2022, 14, 17.	8.2	34
4	Generation of the tumor-suppressive secretome from tumor cells. Theranostics, 2021, 11, 8517-8534.	10.0	20
5	Development and validation of a novel mRNA signature for predicting early relapse in non-small cell lung cancer. Japanese Journal of Clinical Oncology, 2021, 51, 1277-1286.	1.3	1
6	INPP5D expression is associated with risk for Alzheimer's disease and induced by plaque-associated microglia. Neurobiology of Disease, 2021, 153, 105303.	4.4	63
7	Intron retention-induced neoantigen load correlates with unfavorable prognosis in multiple myeloma. Oncogene, 2021, 40, 6130-6138.	5. 9	21
8	Tumor-Infiltrating Immune-Related Long Non-Coding RNAs Indicate Prognoses and Response to PD-1 Blockade in Head and Neck Squamous Cell Carcinoma. Frontiers in Immunology, 2021, 12, 692079.	4.8	7
9	Impact of <i>PLCG2</i> expression on Microglial Biology and Disease Pathogenesis in Alzheimer's Disease. Alzheimer's and Dementia, 2021, 17, e058740.	0.8	2
10	PLCG2 expression is associated with plaque-associated microglia in Alzheimer's disease Alzheimer's and Dementia, 2021, 17 Suppl 3, e054755.	0.8	0
11	Sestrin 3 Protects Against Dietâ€Induced Nonalcoholic Steatohepatitis in Mice Through Suppression of Transforming Growth Factor β Signal Transduction. Hepatology, 2020, 71, 76-92.	7.3	44
12	Highly robust model of transcription regulator activity predicts breast cancer overall survival. BMC Medical Genomics, 2020, 13, 49.	1.5	3
13	Upregulated KPNA2 promotes hepatocellular carcinoma progression and indicates prognostic significance across human cancer types. Acta Biochimica Et Biophysica Sinica, 2019, 51, 285-292.	2.0	29
14	Development of a Novel Deep Transfer Learning Framework to Characterize Inter- and Intra-Tumor Heterogeneity in Myeloma Patients. Blood, 2019, 134, 3075-3075.	1.4	0
15	A Highly Robust Model for Predicting Outcome of Multiple Myeloma Patients By Inferring Patient-Specific Transcription Factor Activity. Blood, 2019, 134, 1785-1785.	1.4	O
16	Seven Genes Based Novel Signature Predicts Clinical Outcome and Platinum Sensitivity of High Grade Illc Serous Ovarian Carcinoma. International Journal of Biological Sciences, 2018, 14, 2012-2022.	6.4	10
17	Cancer stem cell associated eight geneâ€'based signature predicts clinical outcomes of colorectal cancer. Oncology Letters, 2018, 17, 442-449.	1.8	5
18	A B7â€CD28 family based signature demonstrates significantly different prognoses and tumor immune landscapes in lung adenocarcinoma. International Journal of Cancer, 2018, 143, 2592-2601.	5.1	21

#	Article	IF	CITATIONS
19	Development of a novel prognostic signature of long non-coding RNAs in lung adenocarcinoma. Journal of Cancer Research and Clinical Oncology, 2017, 143, 1649-1657.	2.5	30
20	Panel of seven long noncoding RNA as a candidate prognostic biomarker for ovarian cancer. OncoTargets and Therapy, 2017, Volume 10, 2805-2813.	2.0	13
21	Regulatory activity based risk model identifies survival of stage II and III colorectal carcinoma. Oncotarget, 2017, 8, 98360-98370.	1.8	4
22	Integrated Multiple "-omics―Data Reveal Subtypes of Hepatocellular Carcinoma. PLoS ONE, 2016, 11, e0165457.	2.5	28