Yong Tao

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11431025/publications.pdf

Version: 2024-02-01

17 papers	1,511 citations	14 h-index	940416 16 g-index
19	19	19	3047
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Epigenetic therapy inhibits metastases by disrupting premetastatic niches. Nature, 2020, 579, 284-290.	13.7	213
2	Rab8 GTPase regulates Klotho-mediated inhibition of cell growth and progression by directly modulating its surface expression in human non-small cell lung cancer. EBioMedicine, 2019, 49, 118-132.	2.7	14
3	Defining UHRF1 Domains that Support Maintenance of Human Colon Cancer DNA Methylation and Oncogenic Properties. Cancer Cell, 2019, 35, 633-648.e7.	7.7	89
4	Aging-like Spontaneous Epigenetic Silencing Facilitates Wnt Activation, Stemness, and BrafV600E-Induced Tumorigenesis. Cancer Cell, 2019, 35, 315-328.e6.	7.7	107
5	Genetic Load and Potential Mutational Meltdown in Cancer Cell Populations. Molecular Biology and Evolution, 2019, 36, 541-552.	3.5	14
6	DNA Methylation Patterns Separate Senescence from Transformation Potential and Indicate Cancer Risk. Cancer Cell, 2018, 33, 309-321.e5.	7.7	84
7	Is the evolution in tumors Darwinian or non-Darwinian?. National Science Review, 2018, 5, 15-17.	4.6	24
8	New Insights into the Evolutionary Rate of Hepatitis B Virus at Different Biological Scales. Journal of Virology, 2015, 89, 3512-3522.	1.5	33
9	Extremely high genetic diversity in a single tumor points to prevalence of non-Darwinian cell evolution. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, E6496-505.	3.3	313
10	Excavation of Pid3 Orthologs with Differential Resistance Spectra to Magnaporthe oryzae in Rice Resource. PLoS ONE, 2014, 9, e93275.	1.1	23
11	Transcriptional Characteristics of Xa21-mediated Defense Responses in Rice. Journal of Integrative Plant Biology, 2011, 53, 300-311.	4.1	21
12	Rapid growth of a hepatocellular carcinoma and the driving mutations revealed by cell-population genetic analysis of whole-genome data. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 12042-12047.	3.3	117
13	Novel Autoimmune Hepatitis-Specific Autoantigens Identified Using Protein Microarray Technology. Journal of Proteome Research, 2010, 9, 30-39.	1.8	71
14	A transcriptomic analysis of superhybrid rice <i>LYP9</i> and its parents. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 7695-7701.	3.3	184
15	Identification of a New Rice Blast Resistance Gene, <i>Pid3</i> , by Genomewide Comparison of Paired Nucleotide-Binding Site–Leucine-Rich Repeat Genes and Their Pseudogene Alleles Between the Two Sequenced Rice Genomes. Genetics, 2009, 182, 1303-1311.	1.2	176
16	A protein chip approach for high-throughput antigen identification and characterization. Proteomics, 2007, 7, 2151-2161.	1.3	24
17	A recombination-based strategy for assembling open-reading frames. Plant Molecular Biology Reporter, 2005, 23, 397-403.	1.0	O