

Rui Ye

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4725242/publications.pdf>

Version: 2024-02-01

10
papers

1,969
citations

1162889

8
h-index

1372474

10
g-index

11
all docs

11
docs citations

11
times ranked

4830
citing authors

#	ARTICLE	IF	CITATIONS
1	Frequent mutations of chromatin remodeling genes in transitional cell carcinoma of the bladder. <i>Nature Genetics</i> , 2011, 43, 875-878.	9.4	638
2	Whole-genome sequencing identifies recurrent mutations in hepatocellular carcinoma. <i>Genome Research</i> , 2013, 23, 1422-1433.	2.4	457
3	Whole-genome and whole-exome sequencing of bladder cancer identifies frequent alterations in genes involved in sister chromatid cohesion and segregation. <i>Nature Genetics</i> , 2013, 45, 1459-1463.	9.4	400
4	Frequent mutations of genes encoding ubiquitin-mediated proteolysis pathway components in clear cell renal cell carcinoma. <i>Nature Genetics</i> , 2012, 44, 17-19.	9.4	295
5	Sex Differences in Genetic Associations With Longevity. <i>JAMA Network Open</i> , 2018, 1, e181670.	2.8	60
6	An epigenetic biomarker combination of PCDH17 and POU4F2 detects bladder cancer accurately by methylation analyses of urine sediment DNA in Han Chinese. <i>Oncotarget</i> , 2016, 7, 2754-2764.	0.8	53
7	The correlation of copy number variations with longevity in a genome-wide association study of Han Chinese. <i>Aging</i> , 2018, 10, 1206-1222.	1.4	25
8	Novel variants in <i>MLL</i> confer to bladder cancer recurrence identified by whole-exome sequencing. <i>Oncotarget</i> , 2016, 7, 2629-2645.	0.8	25
9	Case report of a "Fraumeni syndrome-like phenotype with a de novo mutation in CHEK2. <i>Medicine (United States)</i> , 2016, 95, e4251.	0.4	8
10	A random forest-based framework for genotyping and accuracy assessment of copy number variations. <i>NAR Genomics and Bioinformatics</i> , 2020, 2, lqaa071.	1.5	8