

Mingming Jia

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

16
papers

15,301
citations

15
h-index

17
g-index

17
ext. papers

17,198
ext. citations

34.9
avg, IF

5.13
L-index

#	Paper	IF	Citations
16	COSMIC: exploring the world's knowledge of somatic mutations in human cancer. <i>Nucleic Acids Research</i> , 2015 , 43, D805-11	20.1	1795
15	COSMIC: mining complete cancer genomes in the Catalogue of Somatic Mutations in Cancer. <i>Nucleic Acids Research</i> , 2011 , 39, D945-50	20.1	1774
14	Massive genomic rearrangement acquired in a single catastrophic event during cancer development. <i>Cell</i> , 2011 , 144, 27-40	56.2	1628
13	A comprehensive catalogue of somatic mutations from a human cancer genome. <i>Nature</i> , 2010 , 463, 191-5	50.4	1303
12	COSMIC: somatic cancer genetics at high-resolution. <i>Nucleic Acids Research</i> , 2017 , 45, D777-D783	20.1	1279
11	The landscape of cancer genes and mutational processes in breast cancer. <i>Nature</i> , 2012 , 486, 400-4	50.4	1264
10	The patterns and dynamics of genomic instability in metastatic pancreatic cancer. <i>Nature</i> , 2010 , 467, 1109-13	50.4	1013
9	The life history of 21 breast cancers. <i>Cell</i> , 2012 , 149, 994-1007	56.2	979
8	Exome sequencing identifies frequent mutation of the SWI/SNF complex gene PBRM1 in renal carcinoma. <i>Nature</i> , 2011 , 469, 539-42	50.4	943
7	Systematic sequencing of renal carcinoma reveals inactivation of histone modifying genes. <i>Nature</i> , 2010 , 463, 360-3	50.4	927
6	A small-cell lung cancer genome with complex signatures of tobacco exposure. <i>Nature</i> , 2010 , 463, 184-9	50.4	852
5	Complex landscapes of somatic rearrangement in human breast cancer genomes. <i>Nature</i> , 2009 , 462, 1005-10	50.4	684
4	Somatic mutations of the histone H3K27 demethylase gene UTX in human cancer. <i>Nature Genetics</i> , 2009 , 41, 521-3	36.3	627
3	Whole exome sequencing of adenoid cystic carcinoma. <i>Journal of Clinical Investigation</i> , 2013 , 123, 2965-8	55.9	188
2	Data mining using the Catalogue of Somatic Mutations in Cancer BioMart. <i>Database: the Journal of Biological Databases and Curation</i> , 2011 , 2011, bar018	5	41
1	Genomic sequencing identifies a few mutations driving the independent origin of primary liver tumors in a chronic hepatitis murine model. <i>PLoS ONE</i> , 2017 , 12, e0187551	3.7	1