

Mingming Jia

List of Publications by Year in Descending Order

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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: somatic cancer genetics at high-resolution. <i>Nucleic Acids Research</i> , 2017 , 45, D777-D783 | 20.1 | 1279 |
| 15 | 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 |
| 14 | COSMIC: exploring the world's knowledge of somatic mutations in human cancer. <i>Nucleic Acids Research</i> , 2015 , 43, D805-11 | 20.1 | 1795 |
| 13 | Whole exome sequencing of adenoid cystic carcinoma. <i>Journal of Clinical Investigation</i> , 2013 , 123, 2965-8 | 5.9 | 188 |
| 12 | The landscape of cancer genes and mutational processes in breast cancer. <i>Nature</i> , 2012 , 486, 400-4 | 50.4 | 1264 |
| 11 | The life history of 21 breast cancers. <i>Cell</i> , 2012 , 149, 994-1007 | 56.2 | 979 |
| 10 | Massive genomic rearrangement acquired in a single catastrophic event during cancer development. <i>Cell</i> , 2011 , 144, 27-40 | 56.2 | 1628 |
| 9 | 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 |
| 8 | 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 |
| 7 | 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 |
| 6 | A small-cell lung cancer genome with complex signatures of tobacco exposure. <i>Nature</i> , 2010 , 463, 184-90 | 50.4 | 852 |
| 5 | A comprehensive catalogue of somatic mutations from a human cancer genome. <i>Nature</i> , 2010 , 463, 191-5 | 50.4 | 1303 |
| 4 | Systematic sequencing of renal carcinoma reveals inactivation of histone modifying genes. <i>Nature</i> , 2010 , 463, 360-3 | 50.4 | 927 |
| 3 | The patterns and dynamics of genomic instability in metastatic pancreatic cancer. <i>Nature</i> , 2010 , 467, 1109-13 | 50.4 | 1013 |
| 2 | Complex landscapes of somatic rearrangement in human breast cancer genomes. <i>Nature</i> , 2009 , 462, 1005-10 | 50.4 | 684 |
| 1 | Somatic mutations of the histone H3K27 demethylase gene UTX in human cancer. <i>Nature Genetics</i> , 2009 , 41, 521-3 | 36.3 | 627 |