

Martin Akerman

List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

2,000
citations

567281

15
h-index

940533

16
g-index

17
all docs

17
docs citations

17
times ranked

3798
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Differential Functions of Splicing Factors in Mammary Transformation and Breast Cancer Metastasis. Cell Reports, 2019, 29, 2672-2688.e7. | 6.4 | 70 |
| 2 | Differentiation of mammary tumors and reduction in metastasis upon <i>Malat1</i> lncRNA loss. Genes and Development, 2016, 30, 34-51. | 5.9 | 488 |
| 3 | Differential connectivity of splicing activators and repressors to the human spliceosome. Genome Biology, 2015, 16, 119. | 9.6 | 33 |
| 4 | SRSF1-Regulated Alternative Splicing in Breast Cancer. Molecular Cell, 2015, 60, 105-117. | 9.7 | 290 |
| 5 | Splicing-Factor Oncoprotein SRSF1 Stabilizes p53 via RPL5 and Induces Cellular Senescence. Molecular Cell, 2013, 50, 56-66. | 9.7 | 96 |
| 6 | Exon-centric regulation of pyruvate kinase M alternative splicing via mutually exclusive exons. Journal of Molecular Cell Biology, 2012, 4, 79-87. | 3.3 | 81 |
| 7 | The splicing factor SRSF1 regulates apoptosis and proliferation to promote mammary epithelial cell transformation. Nature Structural and Molecular Biology, 2012, 19, 220-228. | 8.2 | 342 |
| 8 | Oncogenic Splicing Factor SRSF1 Is a Critical Transcriptional Target of MYC. Cell Reports, 2012, 1, 110-117. | 6.4 | 169 |
| 9 | Ancient diversity of splicing motifs and protein surfaces in the wild emmer wheat (<i>Triticum</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 1 Pathology, 2012, 13, 276-287. | 4.2 | 45 |
| 10 | SpliceTrap: a method to quantify alternative splicing under single cellular conditions. Bioinformatics, 2011, 27, 3010-3016. | 4.1 | 86 |
| 11 | Heterogeneous Nuclear Ribonucleoprotein H1/H2-dependent Unsplicing of Thymidine Phosphorylase Results in Anticancer Drug Resistance. Journal of Biological Chemistry, 2011, 286, 3741-3754. | 3.4 | 45 |
| 12 | SFmap: a web server for motif analysis and prediction of splicing factor binding sites. Nucleic Acids Research, 2010, 38, W281-W285. | 14.5 | 89 |
| 13 | A computational approach for genome-wide mapping of splicing factor binding sites. Genome Biology, 2009, 10, R30. | 9.6 | 54 |
| 14 | Does distance matter? Variations in alternative splicing regulation. Nucleic Acids Research, 2007, 35, 5487-5498. | 14.5 | 16 |
| 15 | Alternative splicing regulation at tandem 3' splice sites. Nucleic Acids Research, 2006, 34, 23-31. | 14.5 | 68 |
| 16 | Novel motifs in amino acid permease genes from Leishmania. Biochemical and Biophysical Research Communications, 2004, 325, 353-366. | 2.1 | 27 |