

Andrew J Oldfield

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4982026/publications.pdf>

Version: 2024-02-01

17
papers

967
citations

759233

12
h-index

940533

16
g-index

22
all docs

22
docs citations

22
times ranked

1571
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular coupling of Tsix regulation and pluripotency. Nature, 2010, 468, 457-460.	27.8	176
2	Histone-Fold Domain Protein NF-Y Promotes Chromatin Accessibility for Cell Type-Specific Master Transcription Factors. Molecular Cell, 2014, 55, 708-722.	9.7	149
3	Multi-omic Profiling Reveals Dynamics of the Phased Progression of Pluripotency. Cell Systems, 2019, 8, 427-445.e10.	6.2	111
4	A prominent and conserved role for YY1 in Xist transcriptional activation. Nature Communications, 2014, 5, 4878.	12.8	100
5	Intragenic Enhancers Attenuate Host Gene Expression. Molecular Cell, 2017, 68, 104-117.e6.	9.7	85
6	The Central Role of PDR1 in the Foundation of Yeast Drug Resistance. Journal of Biological Chemistry, 2007, 282, 5063-5074.	3.4	64
7	NF-Y controls fidelity of transcription initiation at gene promoters through maintenance of the nucleosome-depleted region. Nature Communications, 2019, 10, 3072.	12.8	53
8	Decoding the function of bivalent chromatin in development and cancer. Genome Research, 2021, 31, 2170-2184.	5.5	48
9	Splicing-associated chromatin signatures: a combinatorial and position-dependent role for histone marks in splicing definition. Nature Communications, 2021, 12, 682.	12.8	43
10	Integrative analysis identifies co-dependent gene expression regulation of BRG1 and CHD7 at distal regulatory sites in embryonic stem cells. Bioinformatics, 2017, 33, 1916-1920.	4.1	31
11	Integrative framework for identification of key cell identity genes uncovers determinants of ES cell identity and homeostasis. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E1581-90.	7.1	26
12	IRFinder-S: a comprehensive suite to discover and explore intron retention. Genome Biology, 2021, 22, 307.	8.8	19
13	Histone marks regulate the epithelial-to-mesenchymal transition via alternative splicing. Cell Reports, 2022, 38, 110357.	6.4	15
14	Transcriptional network dynamics during the progression of pluripotency revealed by integrative statistical learning. Nucleic Acids Research, 2020, 48, 1828-1842.	14.5	14
15	TALC: Transcript-level Aware Long-read Correction. Bioinformatics, 2020, 36, 5000-5006.	4.1	14
16	A cell-to-patient machine learning transfer approach uncovers novel basal-like breast cancer prognostic markers amongst alternative splice variants. BMC Biology, 2021, 19, 70.	3.8	13
17	A prominent and conserved role for YY1 in Xist transcriptional activation. Epigenetics and Chromatin, 2013, 6, .	3.9	0