Wayne O Miles

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3379415/publications.pdf

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		687363	526287
27	803	13	27
papers	citations	h-index	g-index
29	29	29	1658
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Pumilio facilitates miRNA regulation of the E2F3 oncogene. Genes and Development, 2012, 26, 356-368.	5.9	140
2	Modeling tumor invasion and metastasis in <i>Drosophila</i> . DMM Disease Models and Mechanisms, 2011, 4, 753-761.	2.4	99
3	Proteomic analysis of pRb loss highlights a signature of decreased mitochondrial oxidative phosphorylation. Genes and Development, 2015, 29, 1875-1889.	5.9	76
4	Inactivation of <i>Fbxw7</i> Impairs dsRNA Sensing and Confers Resistance to PD-1 Blockade. Cancer Discovery, 2020, 10, 1296-1311.	9.4	49
5	Beyond CLIP: advances and opportunities to measure RBP–RNA and RNA–RNA interactions. Nucleic Acids Research, 2019, 47, 5490-5501.	14.5	48
6	Alternative Polyadenylation in Triple-Negative Breast Tumors Allows NRAS and c-JUN to Bypass PUMILIO Posttranscriptional Regulation. Cancer Research, 2016, 76, 7231-7241.	0.9	47
7	Destabilization of NOXA mRNA as a common resistance mechanism to targeted therapies. Nature Communications, 2019, 10, 5157.	12.8	46
8	Medea SUMOylation restricts the signaling range of the Dpp morphogen in the <i>Drosophila</i> embryo. Genes and Development, 2008, 22, 2578-2590.	5.9	45
9	Thermostability, Tunability, and Tenacity of RNA as Rubbery Anionic Polymeric Materials in Nanotechnology and Nanomedicine—Specific Cancer Targeting with Undetectable Toxicity. Chemical Reviews, 2021, 121, 7398-7467.	47.7	45
10	Non-coding and Coding Transcriptional Profiles Are Significantly Altered in Pediatric Retinoblastoma Tumors. Frontiers in Oncology, 2019, 9, 221.	2.8	27
11	Postâ€transcriptional gene expression control by <scp>NANOS</scp> is upâ€regulated and functionally important in <scp>pR</scp> bâ€deficient cells. EMBO Journal, 2014, 33, 2201-2215.	7.8	25
12	Transcriptional targeting of oncogene addiction in medullary thyroid cancer. JCI Insight, 2018, 3, .	5.0	19
13	Clinical Utility of CDK4/6 Inhibitors in Sarcoma: Successes and Future Challenges. JCO Precision Oncology, 2022, 6, e2100211.	3.0	19
14	Proteogenomic Analysis of Surgically Resected Lung Adenocarcinoma. Journal of Thoracic Oncology, 2018, 13, 1519-1529.	1.1	17
15	Comprehensive Analysis of MEN1 Mutations and Their Role in Cancer. Cancers, 2020, 12, 2616.	3.7	16
16	<i>In Vivo</i> Regulation of E2F1 by Polycomb Group Genes in <i>Drosophila</i> . G3: Genes, Genomes, Genetics, 2012, 2, 1651-1660.	1.8	14
17	The LSD1 Family of Histone Demethylases and the Pumilio Posttranscriptional Repressor Function in a Complex Regulatory Feedback Loop. Molecular and Cellular Biology, 2015, 35, 4199-4211.	2.3	12
18	Metabolite systems profiling identifies exploitable weaknesses in retinoblastoma. FEBS Letters, 2019, 593, 23-41.	2.8	11

#	Article	IF	CITATIONS
19	RNA-Sequencing of Primary Retinoblastoma Tumors Provides New Insights and Challenges Into Tumor Development. Frontiers in Genetics, 2018, 9, 170.	2.3	10
20	Reduced RBPMS Levels Promote Cell Proliferation and Decrease Cisplatin Sensitivity in Ovarian Cancer Cells. International Journal of Molecular Sciences, 2022, 23, 535.	4.1	6
21	Drivers of genomic loss of heterozygosity in leiomyosarcoma are distinct from carcinomas. Npj Precision Oncology, 2022, 6, 29.	5.4	6
22	Pumilio and nanos RNA-binding proteins counterbalance the transcriptional consequences of RB1 inactivation. Molecular and Cellular Oncology, 2014, 1, e968074.	0.7	3
23	RB-loss puts focus on Myc. Nature Cell Biology, 2015, 17, 968-969.	10.3	3
24	Slit2/Robo1 signaling inhibits smallâ€cell lung cancer by targeting βâ€catenin signaling in tumor cells and macrophages. Molecular Oncology, 2023, 17, 839-856.	4.6	3
25	RNA editing signatures identify melanoma patients who respond to Pembrolizumab or Nivolumab treatment. Translational Oncology, 2021, 14, 101197.	3.7	2
26	Mass COVID-19 patient screening using UvsX and UvsY mediated DNA recombination and high throughput parallel sequencing. Scientific Reports, 2022, 12, 4082.	3.3	2
27	Integrated multi-omics analysis of RB-loss identifies widespread cellular programming and synthetic weaknesses. Communications Biology, 2021, 4, 977.	4.4	1