Lin Zhang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8137633/publications.pdf

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30	3,477	22	29
papers	citations	h-index	g-index
30	30	30	6598 citing authors
all docs	docs citations	times ranked	

#	Article	IF	CITATIONS
1	Systematic Pan-Cancer Characterization of Nuclear Receptors Identifies Potential Cancer Biomarkers and Therapeutic Targets. Cancer Research, 2022, 82, 46-59.	0.9	3
2	Systematic illumination of druggable genes in cancer genomes. Cell Reports, 2022, 38, 110400.	6.4	14
3	Endothelial plasticity drives aberrant vascularization and impedes cardiac repair after myocardial infarction., 2022, 1, 372-388.		9
4	Targeting PAK4 to reprogram the vascular microenvironment and improve CAR-T immunotherapy for glioblastoma. Nature Cancer, 2021, 2, 83-97.	13.2	56
5	Synergistic immunotherapy of glioblastoma by dual targeting of IL-6 and CD40. Nature Communications, 2021, 12, 3424.	12.8	74
6	Targeting the transcription cycle and RNA processing in cancer treatment. Current Opinion in Pharmacology, 2021, 58, 69-75.	3.5	1
7	The Cancer Surfaceome Atlas integrates genomic, functional and drug response data to identify actionable targets. Nature Cancer, 2021, 2, 1406-1422.	13.2	33
8	Integrated proteogenomic deep sequencing and analytics accurately identify non-canonical peptides in tumor immunopeptidomes. Nature Communications, 2020, 11 , 1293 .	12.8	196
9	Integrative comparison of the genomic and transcriptomic landscape between prostate cancer patients of predominantly African or European genetic ancestry. PLoS Genetics, 2020, 16, e1008641.	3.5	78
10	Vascular niche IL-6 induces alternative macrophage activation in glioblastoma through HIF-2α. Nature Communications, 2018, 9, 559.	12.8	176
11	Repression of BET activity sensitizes homologous recombination–proficient cancers to PARP inhibition. Science Translational Medicine, 2017, 9, .	12.4	180
12	Suppression of MicroRNA 200 Family Expression by Oncogenic KRAS Activation Promotes Cell Survival and Epithelial-Mesenchymal Transition in KRAS-Driven Cancer. Molecular and Cellular Biology, 2016, 36, 2742-2754.	2.3	42
13	Detection of Long Noncoding RNA Expression by Nonradioactive Northern Blots. Methods in Molecular Biology, 2016, 1402, 177-188.	0.9	5
14	Characterization of Long Noncoding RNA-Associated Proteins by RNA-Immunoprecipitation. Methods in Molecular Biology, 2016, 1402, 19-26.	0.9	16
15	Effects of BRCA1/2 on Ovarian and Breast Cancer Survivalâ€"Response. Clinical Cancer Research, 2015, 21, 3807-3807.	7.0	5
16	Comprehensive Genomic Characterization of Long Non-coding RNAs across Human Cancers. Cancer Cell, 2015, 28, 529-540.	16.8	601
17	Effects of <i>BRCA1</i> - and <i>BRCA2</i> -Related Mutations on Ovarian and Breast Cancer Survival: A Meta-analysis. Clinical Cancer Research, 2015, 21, 211-220.	7.0	165
18	A Functional Genomic Approach Identifies FAL1 as an Oncogenic Long Noncoding RNA that Associates with BMI1 and Represses p21 Expression in Cancer. Cancer Cell, 2014, 26, 344-357.	16.8	361

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19	Lin-28 Homologue A (LIN28A) Promotes Cell Cycle Progression via Regulation of Cyclin-dependent Kinase 2 (CDK2), Cyclin D1 (CCND1), and Cell Division Cycle 25 Homolog A (CDC25A) Expression in Cancer. Journal of Biological Chemistry, 2012, 287, 17386-17397.	3.4	60
20	Genomic DNA Copy-Number Alterations of the let-7 Family in Human Cancers. PLoS ONE, 2012, 7, e44399.	2.5	44
21	Distinct Expression Levels and Patterns of Stem Cell Marker, Aldehyde Dehydrogenase Isoform 1 (ALDH1), in Human Epithelial Cancers. PLoS ONE, 2010, 5, e10277.	2.5	362
22	Expression of Activated PIK3CA in Ovarian Surface Epithelium Results in Hyperplasia but Not Tumor Formation. PLoS ONE, 2009, 4, e4295.	2.5	30
23	Expression profile of microRNA in epithelial cancer: diagnosis, classification and prediction. Expert Opinion on Medical Diagnostics, 2009, 3, 25-36.	1.6	4
24	Therapeutic MicroRNA Strategies in Human Cancer. AAPS Journal, 2009, 11, 747-57.	4.4	153
25	MicroRNA epigenetic alterations in human cancer: One step forward in diagnosis and treatment. International Journal of Cancer, 2008, 122, 963-968.	5.1	84
26	Genomic and epigenetic alterations deregulate microRNA expression in human epithelial ovarian cancer. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 7004-7009.	7.1	491
27	Transcriptional Regulation of PIK3CA Oncogene by NF-κB in Ovarian Cancer Microenvironment. PLoS ONE, 2008, 3, e1758.	2.5	41
28	Integrative Genomic Analysis of Phosphatidylinositol 3′-Kinase Family Identifies PIK3R3 as a Potential Therapeutic Target in Epithelial Ovarian Cancer. Clinical Cancer Research, 2007, 13, 5314-5321.	7.0	111
29	miRNA genetic alterations in human cancers. Expert Opinion on Biological Therapy, 2007, 7, 1375-1386.	3.1	50
30	MicroRNAs: A New Insight into Cancer Genome. Cell Cycle, 2006, 5, 2216-2219.	2.6	32