Jun S Wei

List of Publications by Year in descending order

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LUN S WEI

#	Article	IF	CITATIONS
1	Classification and diagnostic prediction of cancers using gene expression profiling and artificial neural networks. Nature Medicine, 2001, 7, 673-679.	30.7	2,352
2	The genetic landscape of high-risk neuroblastoma. Nature Genetics, 2013, 45, 279-284.	21.4	990
3	Comprehensive Genomic Analysis of Rhabdomyosarcoma Reveals a Landscape of Alterations Affecting a Common Genetic Axis in Fusion-Positive and Fusion-Negative Tumors. Cancer Discovery, 2014, 4, 216-231.	9.4	596
4	Relapsed neuroblastomas show frequent RAS-MAPK pathway mutations. Nature Genetics, 2015, 47, 864-871.	21.4	451
5	The Genomic Landscape of the Ewing Sarcoma Family of Tumors Reveals Recurrent STAG2 Mutation. PLoS Genetics, 2014, 10, e1004475.	3.5	335
6	Genetic predisposition to neuroblastoma mediated by a LMO1 super-enhancer polymorphism. Nature, 2015, 528, 418-421.	27.8	263
7	PAX3–FOXO1 Establishes Myogenic Super Enhancers and Confers BET Bromodomain Vulnerability. Cancer Discovery, 2017, 7, 884-899.	9.4	221
8	Prediction of Clinical Outcome Using Gene Expression Profiling and Artificial Neural Networks for Patients with Neuroblastoma. Cancer Research, 2004, 64, 6883-6891.	0.9	183
9	EZH2 Mediates Epigenetic Silencing of Neuroblastoma Suppressor Genes <i>CASZ1</i> , <i>CLU</i> , <i>RUNX3</i> , and <i>NGFR</i> . Cancer Research, 2012, 72, 315-324.	0.9	161
10	Identification of GPC2 as an Oncoprotein and Candidate Immunotherapeutic Target in High-Risk Neuroblastoma. Cancer Cell, 2017, 32, 295-309.e12.	16.8	148
11	Cross-Cohort Analysis Identifies a TEAD4–MYCN Positive Feedback Loop as the Core Regulatory Element of High-Risk Neuroblastoma. Cancer Discovery, 2018, 8, 582-599.	9.4	119
12	Inherited predisposition to malignant mesothelioma and overall survival following platinum chemotherapy. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 9008-9013.	7.1	108
13	Therapeutic targeting of ATR yields durable regressions in small cell lung cancers with high replication stress. Cancer Cell, 2021, 39, 566-579.e7.	16.8	107
14	MEK inhibition induces MYOG and remodels super-enhancers in RAS-driven rhabdomyosarcoma. Science Translational Medicine, 2018, 10, .	12.4	104
15	Genomic Classification and Clinical Outcome in Rhabdomyosarcoma: A Report From an International Consortium. Journal of Clinical Oncology, 2021, 39, 2859-2871.	1.6	101
16	MultiDimensional ClinOmics for Precision Therapy of Children and Adolescent Young Adults with Relapsed and Refractory Cancer: A Report from the Center for Cancer Research. Clinical Cancer Research, 2016, 22, 3810-3820.	7.0	99
17	Clinically Relevant Cytotoxic Immune Cell Signatures and Clonal Expansion of T-Cell Receptors in High-Risk <i>MYCN</i> -Not-Amplified Human Neuroblastoma. Clinical Cancer Research, 2018, 24, 5673-5684.	7.0	92
18	Clonal Evolution and Heterogeneity of Osimertinib Acquired Resistance Mechanisms in EGFR Mutant Lung Cancer. Cell Reports Medicine, 2020, 1, 100007.	6.5	78

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19	Tremelimumab in Combination With Microwave Ablation in Patients With RefractoryÂBiliary Tract Cancer. Hepatology, 2019, 69, 2048-2060.	7.3	77
20	Notch signaling and efficacy of PD-1/PD-L1 blockade in relapsed small cell lung cancer. Nature Communications, 2021, 12, 3880.	12.8	71
21	Low mutation burden and frequent loss of CDKN2A/B and SMARCA2, but not PRC2, define premalignant neurofibromatosis type 1–associated atypical neurofibromas. Neuro-Oncology, 2019, 21, 981-992.	1.2	69
22	Frequent inactivating germline mutations in DNA repair genes in patients with Ewing sarcoma. Genetics in Medicine, 2017, 19, 955-958.	2.4	60
23	Clonality and Evolutionary History of Rhabdomyosarcoma. PLoS Genetics, 2015, 11, e1005075.	3.5	58
24	Aurora B kinase is a potent and selective target in MYCN-driven neuroblastoma. Oncotarget, 2015, 6, 35247-35262.	1.8	52
25	Identification of microRNAs specific for epithelial cell adhesion molecule–positive tumor cells in hepatocellular carcinoma. Hepatology, 2015, 62, 829-840.	7.3	51
26	microRNA Profiling Identifies Cancer-Specific and Prognostic Signatures in Pediatric Malignancies. Clinical Cancer Research, 2009, 15, 5560-5568.	7.0	49
27	MYCN controls an alternative RNA splicing program in high-risk metastatic neuroblastoma. Cancer Letters, 2016, 371, 214-224.	7.2	46
28	Whole-exome sequencing reveals germline-mutated small cell lung cancer subtype with favorable response to DNA repair–targeted therapies. Science Translational Medicine, 2021, 13, .	12.4	35
29	CASZ1 induces skeletal muscle and rhabdomyosarcoma differentiation through a feed-forward loop with MYOD and MYOC. Nature Communications, 2020, 11, 911.	12.8	32
30	Expression Quantitative Trait Loci and Receptor Pharmacology Implicate Arg1 and the GABA-A Receptor as Therapeutic Targets in Neuroblastoma. Cell Reports, 2014, 9, 1034-1046.	6.4	28
31	Pathogenic Germline Variants in Cancer Susceptibility Genes in Children and Young Adults With Rhabdomyosarcoma. JCO Precision Oncology, 2021, 5, 75-87.	3.0	27
32	Miswired Enhancer Logic Drives a Cancer of the Muscle Lineage. IScience, 2020, 23, 101103.	4.1	26
33	Immuno-transcriptomic profiling of extracranial pediatric solid malignancies. Cell Reports, 2021, 37, 110047.	6.4	26
34	BAF complexes drive proliferation and block myogenic differentiation in fusion-positive rhabdomyosarcoma. Nature Communications, 2021, 12, 6924.	12.8	25
35	Anaplastic Lymphoma Kinase Gene Rearrangement in Children and Young Adults With Mesothelioma. Journal of Thoracic Oncology, 2020, 15, 457-461.	1.1	24
36	Massively Parallel Sequencing Reveals an Accumulation of De Novo Mutations and an Activating Mutation of LPAR1 in a Patient with Metastatic Neuroblastoma. PLoS ONE, 2013, 8, e77731.	2.5	24

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37	Somatic structural variation targets neurodevelopmental genes and identifies <i>SHANK2</i> as a tumor suppressor in neuroblastoma. Genome Research, 2020, 30, 1228-1242.	5.5	20
38	Paired Expression Analysis of Tumor Cell Surface Antigens. Frontiers in Oncology, 2017, 7, 173.	2.8	16
39	Clinical and Genomic Characteristics of Small Cell Lung Cancer in Never Smokers. Chest, 2020, 158, 1723-1733.	0.8	16
40	Phase 2 Study of Olaparib in Malignant Mesothelioma and Correlation of Efficacy With Germline or Somatic Mutations in BAP1 Gene. JTO Clinical and Research Reports, 2021, 2, 100231.	1,1	16
41	Tumor Mutation Burden, Expressed Neoantigens and the Immune Microenvironment in Diffuse Gliomas. Cancers, 2021, 13, 6092.	3.7	14
42	Outcome-Related Signatures Identified by Whole Transcriptome Sequencing of Resectable Stage III/IV Melanoma Evaluated after Starting Hu14.18-IL2. Clinical Cancer Research, 2020, 26, 3296-3306.	7.0	12
43	Genomic and Transcriptomic Analysis of Relapsed and Refractory Childhood Solid Tumors Reveals a Diverse Molecular Landscape and Mechanisms of Immune Evasion. Cancer Research, 2021, 81, 5818-5832.	0.9	10
44	New technologies for diagnosing pediatric tumors. Expert Opinion on Medical Diagnostics, 2008, 2, 1205-1219.	1.6	5
45	Ataxia telangiectasia mutated germline pathogenic variant in adrenocortical carcinoma. Cancer Genetics, 2021, 256-257, 21-25.	0.4	4
46	Dynamics of genomic and immune responses during primary immunotherapy resistance in mismatch repair–deficient tumors. Journal of Physical Education and Sports Management, 2020, 6, a005678.	1.2	3
47	Report of Canonical <i>BCR</i> - <i>ABL1</i> Fusion in Glioblastoma. JCO Precision Oncology, 2021, 5, 1348-1353.	3.0	3
48	Detailed Multi-Method Analysis of Bone Marrow from Pediatric Pre-B-ALL Patients Prior to CD19-CAR-T Therapy Subsequently Evidencing Overt CAR-T Resistance. Blood, 2019, 134, 2744-2744.	1.4	3
49	Case Report: Single-Cell Transcriptomic Analysis of an Anaplastic Oligodendroglioma Post Immunotherapy. Frontiers in Oncology, 2020, 10, 601452.	2.8	1
50	Aneurysmal Fibrous Histiocytoma: A Large Soft Tissue Tumor with Metastases Treated with Palliative Radiation Therapy and Targeted Therapy. Case Reports in Oncology, 2021, 14, 17-23.	0.7	1
51	Dentithecamides A–H, Diacylated Zoanthoxanthin Derivatives with PAX3-FOXO1 Inhibitory Activity from the Hydroid <i>Dentitheca habereri</i> . Journal of Natural Products, 2022, 85, 1419-1427.	3.0	1
52	Initial Genomic Analysis of a Pure Erythroid Leukemia Developing in Association with Hydroyurea Treatment for Sickle Cell Anemia. Blood, 2012, 120, 3254-3254.	1.4	0
53	Clonal evolution and osimertinib resistance mechanisms identified by whole exome and transcriptome sequencing in EGFR mutant NSCLC Journal of Clinical Oncology, 2019, 37, 9049-9049.	1.6	0