## Wenbin Wei

## List of Publications by Year in descending order

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Version: 2024-02-01

109137 4,532 82 35 h-index citations papers

65 g-index 83 83 83 7842 docs citations times ranked citing authors all docs

106150

#	Article	IF	CITATIONS
1	Novel markers for differentiation of lobular and ductal invasive breast carcinomas by laser microdissection and microarray analysis. BMC Cancer, 2007, 7, 55.	1.1	341
2	Cytomegalovirus-seropositivity has a profound influence on the magnitude of major lymphoid subsets within healthy individuals. Clinical and Experimental Immunology, 2009, 155, 423-432.	1.1	225
3	Meta-Analysis of the Alzheimer's Disease Human Brain Transcriptome and Functional Dissection in Mouse Models. Cell Reports, 2020, 32, 107908.	2.9	199
4	Identification of serum biomarkers for colon cancer by proteomic analysis. British Journal of Cancer, 2006, 94, 1898-1905.	2.9	198
5	The Epstein–Barr virus and the pathogenesis of lymphoma. Journal of Pathology, 2015, 235, 312-322.	2.1	184
6	Expression of the Epstein-Barr Virus-Encoded Epstein-Barr Virus Nuclear Antigen 1 in Hodgkin's Lymphoma Cells Mediates Up-Regulation of CCL20 and the Migration of Regulatory T Cells. American Journal of Pathology, 2008, 173, 195-204.	1.9	162
7	The H3K27me3 demethylase, KDM6B, is induced by Epstein–Barr virus and over-expressed in Hodgkin's Lymphoma. Oncogene, 2011, 30, 2037-2043.	2.6	133
8	Tagging genomic sequences that direct transgene expression by activation of a promoter trap in plants. Transgenic Research, 1993, 2, 33-47.	1.3	132
9	Should grade 3 endometrioid endometrial carcinoma be considered a type 2 cancer—A clinical and pathological evaluation. Gynecologic Oncology, 2012, 124, 15-20.	0.6	132
10	A novel CDK inhibitor, CYC202 (R-roscovitine), overcomes the defect in p53-dependent apoptosis in B-CLL by down-regulation of genes involved in transcription regulation and survival. Blood, 2005, 105, 4484-4491.	0.6	129
11	Epstein–Barr virus-encoded EBNA1 regulates cellular gene transcription and modulates the STAT1 and TGFβ signaling pathways. Oncogene, 2007, 26, 4135-4147.	2.6	114
12	DNA methylation profiles of long- and short-term glioblastoma survivors. Epigenetics, 2013, 8, 149-156.	1.3	108
13	Induction of autotaxin by the Epstein-Barr virus promotes the growth and survival of Hodgkin lymphoma cells. Blood, 2005, 106, 2138-2146.	0.6	101
14	Down-regulation of the TGF-beta target gene, PTPRK, by the Epstein-Barr virus–encoded EBNA1 contributes to the growth and survival of Hodgkin lymphoma cells. Blood, 2008, 111, 292-301.	0.6	96
15	The EBV-encoded latent membrane proteins, LMP2A and LMP2B, limit the actions of interferon by targeting interferon receptors for degradation. Oncogene, 2009, 28, 3903-3914.	2.6	94
16	Bmi-1 is induced by the Epstein-Barr virus oncogene LMP1 and regulates the expression of viral target genes in Hodgkin lymphoma cells. Blood, 2007, 109, 2597-2603.	0.6	89
17	Down-regulation of BLIMP1 $\hat{l}\pm$ by the EBV oncogene, LMP-1, disrupts the plasma cell differentiation program and prevents viral replication in B cells: implications for the pathogenesis of EBV-associated B-cell lymphomas. Blood, 2011, 117, 5907-5917.	0.6	86
18	Fibroblast gene expression profile reflects the stage of tumour progression in oral squamous cell carcinoma. Journal of Pathology, 2011, 223, 459-469.	2.1	84

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19	The ATM tumour suppressor gene is downâ€regulated in EBVâ€associated nasopharyngeal carcinoma. Journal of Pathology, 2009, 217, 345-352.	2.1	83
20	Liver Tumors in Wild Flatfish: A Histopathological, Proteomic, and Metabolomic Study. OMICS A Journal of Integrative Biology, 2005, 9, 281-299.	1.0	82
21	Epigenetic and Transcriptional Changes Which Follow Epstein-Barr Virus Infection of Germinal Center B Cells and Their Relevance to the Pathogenesis of Hodgkin's Lymphoma. Journal of Virology, 2011, 85, 9568-9577.	1.5	81
22	Oncogenic human papillomavirus imposes an instructive pattern of DNA methylation changes which parallel the natural history of cervical HPV infection in young women. Carcinogenesis, 2012, 33, 1286-1293.	1.3	79
23	Upregulation of Eps8 in oral squamous cell carcinoma promotes cell migration and invasion through integrin-dependent Rac1 activation. Oncogene, 2009, 28, 2524-2534.	2.6	77
24	Epstein-Barr virus induction of the Hedgehog signalling pathway imposes a stem cell phenotype on human epithelial cells. Journal of Pathology, 2013, 231, 367-377.	2.1	65
25	A data-driven approach links microglia to pathology and prognosis in amyotrophic lateral sclerosis. Acta Neuropathologica Communications, 2017, 5, 23.	2.4	63
26	Changes in the serum proteome associated with the development of hepatocellular carcinoma in hepatitis C-related cirrhosis. British Journal of Cancer, 2006, 94, 287-292.	2.9	62
27	High-frequency transformation ofArabidopsis thaliana byAgrobacterium tumefaciens. Plant Molecular Biology Reporter, 1992, 10, 178-189.	1.0	61
28	Proteomic profiling of urine for the detection of colon cancer. Proteome Science, 2008, 6, 19.	0.7	56
29	Apoptotic resistance to ionizing radiation in pediatric B-precursor acute lymphoblastic leukemia frequently involves increased NF-κB survival pathway signaling. Blood, 2004, 104, 1465-1473.	0.6	52
30	Tetraspanin CD151 is a novel prognostic marker in poor outcome endometrial cancer. British Journal of Cancer, 2011, 104, 1611-1618.	2.9	50
31	A Global View of the Oncogenic Landscape in Nasopharyngeal Carcinoma: An Integrated Analysis at the Genetic and Expression Levels. PLoS ONE, 2012, 7, e41055.	1.1	49
32	Identification of macrophage migration inhibitory factor and human neutrophil peptides 1–3 as potential biomarkers for gastric cancer. British Journal of Cancer, 2009, 101, 295-302.	2.9	45
33	Different Patterns of Epstein-Barr Virus Latency in Endemic Burkitt Lymphoma (BL) Lead to Distinct Variants within the BL-Associated Gene Expression Signature. Journal of Virology, 2013, 87, 2882-2894.	1.5	45
34	Combined proteome and transcriptome analyses for the discovery of urinary biomarkers for urothelial carcinoma. British Journal of Cancer, 2013, 108, 1854-1861.	2.9	41
35	The Pathway Coexpression Network: Revealing pathway relationships. PLoS Computational Biology, 2018, 14, e1006042.	1.5	41
36	A novel nucleic acid helicase gene identified by promoter trapping in Arabidopsis. Plant Journal, 1997, 11, 1307-1314.	2.8	39

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37	Long non-coding RNA Neat1 regulates adaptive behavioural response to stress in mice. Translational Psychiatry, 2020, 10, 171.	2.4	38
38	Collagen Induces a More Proliferative, Migratory and Chemoresistant Phenotype in Head and Neck Cancer via DDR1. Cancers, 2019, 11, 1766.	1.7	36
39	Deep phenotyping of peripheral tissue facilitates mechanistic disease stratification in sporadic Parkinson's disease. Progress in Neurobiology, 2020, 187, 101772.	2.8	35
40	The EBV-Encoded Oncoprotein, LMP1, Induces an Epithelial-to-Mesenchymal Transition (EMT) via Its CTAR1 Domain through Integrin-Mediated ERK-MAPK Signalling. Cancers, 2018, 10, 130.	1.7	34
41	Epstein–Barr virus-encoded LMP1 induces a hyperproliferative and inflammatory gene expression programme in cultured keratinocytes. Journal of General Virology, 2008, 89, 2806-2820.	1.3	33
42	Methylation profiling and evaluation of demethylating therapy in renal cell carcinoma. Clinical Epigenetics, 2013, 5, 16.	1.8	33
43	Oncogenic <scp>S1P</scp> signalling in <scp>EBV</scp> â€associated nasopharyngeal carcinoma activates <scp>AKT</scp> and promotes cell migration through <scp>S1P</scp> receptor 3. Journal of Pathology, 2017, 242, 62-72.	2.1	33
44	Suppression of the <scp>LMP2A</scp> target gene, <i><scp>EGR</scp>â€1</i> , protects Hodgkin's lymphoma cells from entry to the <scp>EBV</scp> lytic cycle. Journal of Pathology, 2013, 230, 399-409.	2.1	31
45	Plasma Proteome Analysis Reveals the Geographical Origin and Liver Tumor Status of Dab (Limanda) Tj ETQq $1\ 1$	0.784314 4.6	rgBT  Overlo
46	Investigation of chromosome 1q reveals differential expression of members of the S100 family in clinical subgroups of intracranial paediatric ependymoma. British Journal of Cancer, 2008, 99, 1136-1143.	2.9	30
47	Assessment of highâ€throughput highâ€resolution MALDIâ€TOFâ€MS of urinary peptides for the detection of muscleâ€invasive bladder cancer. Proteomics - Clinical Applications, 2011, 5, 493-503.	0.8	29
48	Three-dimensional culturing of the Hodgkin lymphoma cell-line L1236 induces a HL tissue-like gene expression pattern. Leukemia and Lymphoma, 2007, 48, 2042-2053.	0.6	28
49	Preclinical and post-treatment changes in the HCC-associated serum proteome. British Journal of Cancer, 2006, 95, 1379-1383.	2.9	27
50	Sphingosine-1-phosphate signalling drives an angiogenic transcriptional programme in diffuse large B cell lymphoma. Leukemia, 2019, 33, 2884-2897.	3.3	26
51	HOPX functions as a tumour suppressor in head and neck cancer. Scientific Reports, 2016, 6, 38758.	1.6	25
52	Human Paralogs of KIAA0187 Were Created through Independent Pericentromeric-Directed and Chromosome-Specific Duplication Mechanisms. Genome Research, 2002, 12, 67-80.	2.4	24
53	Assessment of novel combinations of biomarkers for the detection of colorectal cancer. Cancer Biomarkers, 2011, 7, 123-132.	0.8	23
54	Epigenetic Silencing of a Proapoptotic Cell Adhesion Molecule, the Immunoglobulin Superfamily Member IGSF4, by Promoter CpG Methylation Protects Hodgkin Lymphoma Cells from Apoptosis. American Journal of Pathology, 2010, 177, 1480-1490.	1.9	22

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55	Coordinated repression of BIM and PUMA by Epstein–Barr virus latent genes maintains the survival of Burkitt lymphoma cells. Cell Death and Differentiation, 2018, 25, 241-254.	5.0	20
56	Therapeutic Targeting the Loss of the Birt-Hogg-Dub $\tilde{A}$ $\mathbb O$ Suppressor Gene. Molecular Cancer Therapeutics, 2011, 10, 80-89.	1.9	18
57	Inflammation and tissue repair markers distinguish the nodular sclerosis and mixed cellularity subtypes of classical Hodgkin's lymphoma. British Journal of Cancer, 2009, 101, 1393-1401.	2.9	17
58	Copy number profiling in von hippelâ€lindau disease renal cell carcinoma. Genes Chromosomes and Cancer, 2011, 50, 479-488.	1,5	17
59	Chemoradiation in Advanced Vulval Carcinoma. International Journal of Gynecological Cancer, 2009, 19, 745-751.	1.2	16
60	Oncogenic effects of WNT5A in Epstein-Barr virus-associated nasopharyngeal carcinoma. International Journal of Oncology, 2014, 44, 1774-1780.	1.4	16
61	MALDI profiles of proteins and lipids for the rapid characterisation of upper GI-tract cancers. Journal of Proteomics, 2013, 80, 207-215.	1.2	15
62	Downâ€regulation of <scp>LPA</scp> receptor 5 contributes to aberrant <scp>LPA</scp> signalling in <scp>EBV</scp> â€associated nasopharyngeal carcinoma. Journal of Pathology, 2015, 235, 456-465.	2.1	15
63	Co-Expression of the Epstein-Barr Virus-Encoded Latent Membrane Proteins and the Pathogenesis of Classic Hodgkin Lymphoma. Cancers, 2018, 10, 285.	1.7	15
64	Macrophage migration inhibitory factor and DJ-1 in gastric cancer: differences between high-incidence and low-incidence areas. British Journal of Cancer, 2012, 107, 1595-1601.	2.9	14
65	Targeting the Ataxia Telangiectasia Mutated-null phenotype in chronic lymphocytic leukemia with pro-oxidants. Haematologica, 2015, 100, 1076-85.	1.7	13
66	CD40L membrane retention enhances the immunostimulatory effects of CD40 ligation. Scientific Reports, 2020, 10, 342.	1.6	13
67	Vesicle Transport in Plants: A Revised Phylogeny of SNARE Proteins. Evolutionary Bioinformatics, 2020, 16, 117693432095657.	0.6	12
68	Induction of Interferon-Stimulated Genes on the IL-4 Response Axis by Epstein-Barr Virus Infected Human B Cells; Relevance to Cellular Transformation. PLoS ONE, 2013, 8, e64868.	1.1	12
69	Evidence for a pathophysiological role of cysteinyl leukotrienes in classical Hodgkin lymphoma. International Journal of Cancer, 2008, 123, 2285-2293.	2.3	11
70	Wnt signalling in adenomas of familial adenomatous polyposis patients. British Journal of Cancer, 2010, 103, 910-917.	2.9	11
71	GCH1 Deficiency Activates Brain Innate Immune Response and Impairs Tyrosine Hydroxylase Homeostasis. Journal of Neuroscience, 2022, 42, 702-716.	1.7	10
72	Connective Tissue Growth Factor Is Expressed in Malignant Cells of Hodgkin Lymphoma but Not in Other Mature B-Cell Lymphomas. American Journal of Clinical Pathology, 2010, 133, 271-280.	0.4	8

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73	Synergistic action of dual IGF1/R and MEK inhibition sensitizes childhood acute lymphoblastic leukemia (ALL) cells to cytotoxic agents and involves downregulation of STAT6 and PDAP1. Experimental Hematology, 2018, 63, 52-63.e5.	0.2	8
74	Regulation of S1PR2 by the EBV oncogene LMP1 in aggressive ABCâ€subtype diffuse large Bâ€cell lymphoma. Journal of Pathology, 2019, 248, 142-154.	2.1	8
75	Low Expression and Promoter Hypermethylation of the Tumour Suppressor SLIT2, are Associated with Adverse Patient Outcomes in Diffuse Large B Cell Lymphoma. Pathology and Oncology Research, 2019, 25, 1223-1231.	0.9	8
76	Agrobacterium-Mediated Transformation of Arabidopsis thaliana: Application in T-DNA Tagging. , $1995$ , $49$ , $63-76$ .		7
77	Detection of pancreatic adenocarcinoma using circulating fragments of fibrinogen. European Journal of Gastroenterology and Hepatology, 2010, 22, 1358-1363.	0.8	7
78	The Quantitative Genetics of Flowering Traits in Wide Crosses of Chickpea. Agriculture (Switzerland), 2022, 12, 486.	1.4	7
79	10 Years of SELDI: What Have we Learnt?. Current Proteomics, 2010, 7, 15-25.	0.1	6
80	Use of Aleuria alantia Lectin Affinity Chromatography to Enrich Candidate Biomarkers from the Urine of Patients with Bladder Cancer. Proteomes, 2015, 3, 266-282.	1.7	5
81	Gene expression and protein array studies of folliculin-regulated pathways. Anticancer Research, 2012, 32, 4663-70.	0.5	5
82	Confounding Effects of Benign Lung Diseases on Non-Small Cell Lung Cancer Serum Biomarker Discovery. Clinical Proteomics, 2009, 5, 148-155.	1.1	2