

Ulrich Lehmann

List of Publications by Year in descending order

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121
papers

5,092
citations

101543

36
h-index

95266

68
g-index

129
all docs

129
docs citations

129
times ranked

8600
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular diagnostics and therapies for gastrointestinal tumors: a real-world experience. <i>Journal of Cancer Research and Clinical Oncology</i> , 2022, 148, 2137-2144.	2.5	1
2	Differences in the MRI Signature and ADC Values of Diffuse Midline Gliomas with H3 K27M Mutation Compared to Midline Glioblastomas. <i>Cancers</i> , 2022, 14, 1397.	3.7	6
3	Interobserver agreement for the histological diagnosis of invasive lobular breast carcinoma. <i>Journal of Pathology: Clinical Research</i> , 2022, 8, 191-205.	3.0	19
4	ERBB2 mutation is associated with sustained tumor cell proliferation after short-term preoperative endocrine therapy in early lobular breast cancer. <i>Modern Pathology</i> , 2022, 35, 1804-1811.	5.5	4
5	Choledochal Cysts Resected during Childhood Show No Mutations of KRAS and BRAF as Early Markers of Malignancy in Cholangiocytes. <i>European Journal of Pediatric Surgery</i> , 2021, 31, 020-024.	1.3	1
6	The Hannover Unified Biobank (HUB) – Centralized Standardised Biobanking at Hannover Medical School. <i>Open Journal of Bioresources</i> , 2021, 8, .	1.5	1
7	Genome-wide DNA methylation profiling is able to identify prefibrotic PMF cases at risk for progression to myelofibrosis. <i>Clinical Epigenetics</i> , 2021, 13, 28.	4.1	1
8	BH3-only protein expression determines hepatocellular carcinoma response to sorafenib-based treatment. <i>Cell Death and Disease</i> , 2021, 12, 736.	6.3	10
9	Diffuse midline gliomas, H3 K27M-mutant are associated with less peritumoral edema and contrast enhancement in comparison to glioblastomas, H3 K27M-wildtype of midline structures. <i>PLoS ONE</i> , 2021, 16, e0249647.	2.5	14
10	Chromosome 2q gain and epigenetic silencing of <i>GATA3</i> in microglandular adenosis of the breast. <i>Journal of Pathology: Clinical Research</i> , 2021, 7, 220-232.	3.0	3
11	<i>TP53</i> mutations are associated with primary endocrine resistance in luminal early breast cancer. <i>Cancer Medicine</i> , 2021, 10, 8581-8594.	2.8	14
12	Mutations associated with age-related clonal hematopoiesis in PMF patients with rapid progression to myelofibrosis. <i>Leukemia</i> , 2020, 34, 1364-1372.	7.2	27
13	FGFR inhibitors in cholangiocarcinoma: what's now and what's next?. <i>Therapeutic Advances in Medical Oncology</i> , 2020, 12, 175883592095329.	3.2	33
14	Infratentorial IDH-mutant astrocytoma is a distinct subtype. <i>Acta Neuropathologica</i> , 2020, 140, 569-581.	7.7	45
15	E-cadherin to P-cadherin switching in lobular breast cancer with tubular elements. <i>Modern Pathology</i> , 2020, 33, 2483-2498.	5.5	26
16	Expression of Olig2, Nestin, NogoA and AQP4 have no impact on overall survival in IDH-wildtype glioblastoma. <i>PLoS ONE</i> , 2020, 15, e0229274.	2.5	9
17	Feasibility of Combined Detection of Gene Mutations and Fusion Transcripts in Bone Marrow Trepines from Leukemic Neoplasms. <i>Journal of Molecular Diagnostics</i> , 2020, 22, 591-598.	2.8	2
18	Coincidence of lymphomatoid granulomatosis, chronic myelomonocytic leukemia, and anaplastic T cell lymphoma after methotrexate therapy for rheumatoid arthritis. <i>Annals of Hematology</i> , 2019, 98, 515-517.	1.8	2

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19	The prognostic role of IDH mutations in homogeneously treated patients with anaplastic astrocytomas and glioblastomas. <i>Acta Neuropathologica Communications</i> , 2019, 7, 156.	5.2	47
20	LINE-1 hypomethylation in human hepatocellular carcinomas correlates with shorter overall survival and CIMP phenotype. <i>PLoS ONE</i> , 2019, 14, e0216374.	2.5	12
21	Variant classification in precision oncology. <i>International Journal of Cancer</i> , 2019, 145, 2996-3010.	5.1	76
22	Discordancy for a Villous Maturation Defect in a Dizygotic Twin Placenta. <i>Fetal and Pediatric Pathology</i> , 2019, 38, 432-436.	0.7	1
23	The detection of BKPyV genotypes II and IV after renal transplantation as a simple tool for risk assessment for PyVAN and transplant outcome already at early stages of BKPyV reactivation. <i>Journal of Clinical Virology</i> , 2019, 113, 14-19.	3.1	8
24	Comprehensive mutation profiling and mRNA expression analysis in atypical chronic myeloid leukemia in comparison with chronic myelomonocytic leukemia. <i>Cancer Medicine</i> , 2019, 8, 742-750.	2.8	19
25	<i>ERBB2</i> mutation frequency in lobular breast cancer with pleomorphic histology or high-risk characteristics by molecular expression profiling. <i>Genes Chromosomes and Cancer</i> , 2019, 58, 175-185.	2.8	27
26	Detection of Aberrant DNA Methylation Patterns in the RB1 Gene. <i>Methods in Molecular Biology</i> , 2018, 1726, 35-47.	0.9	5
27	Next-Generation Sequencing Analysis of Laser-Microdissected Formalin-Fixed and Paraffin-Embedded (FFPE) Tissue Specimens. <i>Methods in Molecular Biology</i> , 2018, 1723, 111-118.	0.9	6
28	<i>CDKN2A</i> loss and <i>PIK3CA</i> mutation in myoepithelial-like metaplastic breast cancer. <i>Journal of Pathology</i> , 2018, 245, 373-383.	4.5	28
29	Molecular defects in BRAF wild-type ameloblastomas and craniopharyngiomas—differences in mutation profiles in epithelial-derived oropharyngeal neoplasms. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2018, 472, 1055-1059.	2.8	14
30	Lobular carcinoma in situ and invasive lobular breast cancer are characterized by enhanced expression of transcription factor AP-2 β . <i>Laboratory Investigation</i> , 2018, 98, 117-129.	3.7	24
31	Estrogen receptor (ESR1) mutation in bone metastases from breast cancer. <i>Modern Pathology</i> , 2018, 31, 56-61.	5.5	29
32	Cerebellar glioblastoma: a clinical series with contemporary molecular analysis. <i>Acta Neurochirurgica</i> , 2018, 160, 2237-2248.	1.7	22
33	Regulation of Cellular Heterogeneity and Rates of Symmetric and Asymmetric Divisions in Triple-Negative Breast Cancer. <i>Cell Reports</i> , 2018, 24, 3237-3250.	6.4	31
34	Hotspot mutations in cancer genes may be missed in routine diagnostics due to neighbouring sequence variants. <i>Experimental and Molecular Pathology</i> , 2018, 105, 37-40.	2.1	2
35	Activating human epidermal growth factor receptor 2 (HER2) gene mutation in bone metastases from breast cancer. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2018, 473, 577-582.	2.8	14
36	Impact of Molecular Genetics on Outcome in Myelofibrosis Patients after Allogeneic Stem Cell Transplantation. <i>Biology of Blood and Marrow Transplantation</i> , 2017, 23, 1095-1101.	2.0	89

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37	Rare ADAR and RNASEH2B variants and a type I interferon signature in glioma and prostate carcinoma risk and tumorigenesis. <i>Acta Neuropathologica</i> , 2017, 134, 905-922.	7.7	12
38	EGFR T790M mutation testing of non-small cell lung cancer tissue and blood samples artificially spiked with circulating cell-free tumor DNA: results of a round robin trial. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2017, 471, 509-520.	2.8	29
39	Molecular Analysis of Circulating Cell-Free DNA from Lung Cancer Patients in Routine Laboratory Practice. <i>Journal of Molecular Diagnostics</i> , 2017, 19, 722-732.	2.8	37
40	A novel germline POLE mutation causes an early onset cancer prone syndrome mimicking constitutional mismatch repair deficiency. <i>Familial Cancer</i> , 2017, 16, 67-71.	1.9	52
41	A liver nodule in a patient transplanted for primary sclerosing cholangitis: an interdisciplinary diagnostic approach. <i>Zeitschrift Fur Gastroenterologie</i> , 2017, 55, 56-62.	0.5	3
42	Transposable Elements in Human Cancer: Causes and Consequences of Deregulation. <i>International Journal of Molecular Sciences</i> , 2017, 18, 974.	4.1	128
43	Absence of MGMT promoter methylation in diffuse midline glioma, H3 K27M-mutant. <i>Acta Neuropathologica Communications</i> , 2017, 5, 98.	5.2	35
44	<i>hsa-mir-183</i> is frequently methylated and related to poor survival in human hepatocellular carcinoma. <i>World Journal of Gastroenterology</i> , 2017, 23, 1568.	3.3	17
45	Prognostic factors in the myoepithelial-like spindle cell type of metaplastic breast cancer. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2016, 469, 191-201.	2.8	12
46	NGS-based BRCA1/2 mutation testing of high-grade serous ovarian cancer tissue: results and conclusions of the first international round robin trial. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2016, 468, 697-705.	2.8	24
47	Precise <i>ERBB2</i> copy number assessment in breast cancer by means of molecular inversion probe array analysis. <i>Oncotarget</i> , 2016, 7, 82733-82740.	1.8	11
48	Routine clinical mutation profiling using next generation sequencing and a customized gene panel improves diagnostic precision in myeloid neoplasms. <i>Oncotarget</i> , 2016, 7, 30084-30093.	1.8	42
49	A child with Li-Fraumeni syndrome: Modes to inactivate the second allele of <i>TP53</i> in three different malignancies. <i>Pediatric Blood and Cancer</i> , 2015, 62, 1481-1484.	1.5	22
50	Loss of DNA methylation at imprinted loci is a frequent event in hepatocellular carcinoma and identifies patients with shortened survival. <i>Clinical Epigenetics</i> , 2015, 7, 110.	4.1	14
51	MicroRNAs: Emerging Novel Clinical Biomarkers for Hepatocellular Carcinomas. <i>Journal of Clinical Medicine</i> , 2015, 4, 1631-1650.	2.4	53
52	Breast Cancer Anti-Estrogen Resistance 4 (BCAR4) Drives Proliferation of IPH-926 lobular Carcinoma Cells. <i>PLoS ONE</i> , 2015, 10, e0136845.	2.5	30
53	Lobular breast cancer - the most common special subtype or a most special common subtype?. <i>Breast Cancer Research</i> , 2015, 17, 99.	5.0	10
54	Testosterone-receptor positive hepatocellular carcinoma in a 29-year old bodybuilder with a history of anabolic androgenic steroid abuse: a case report. <i>BMC Gastroenterology</i> , 2015, 15, 60.	2.0	42

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55	The CpG island methylator phenotype in breast cancer is associated with the lobular subtype. <i>Epigenomics</i> , 2015, 7, 187-199.	2.1	26
56	Quantitative DNA Methylation Analysis by Pyrosequencing®. <i>Methods in Molecular Biology</i> , 2015, 1315, 175-188.	0.9	12
57	Quantitative Validation and Quality Control of Pyrosequencing® Assays. <i>Methods in Molecular Biology</i> , 2015, 1315, 39-46.	0.9	6
58	Analysis of Mutational Hotspots in Routinely Processed Bone Marrow Trepines by Pyrosequencing®. <i>Methods in Molecular Biology</i> , 2015, 1315, 103-114.	0.9	11
59	Impact of Molecular Genetics on Disease-Free Survival in Myelofibrosis Patients Following Allogeneic Stem Cell Transplantation. <i>Blood</i> , 2015, 126, 352-352.	1.4	2
60	Comprehensive Molecular Profiling of Archival Bone Marrow Trepines Using a Commercially Available Leukemia Panel and Semiconductor-Based Targeted Resequencing. <i>PLoS ONE</i> , 2015, 10, e0133930.	2.5	14
61	DNA methylation, microRNAs, and their crosstalk as potential biomarkers in hepatocellular carcinoma. <i>World Journal of Gastroenterology</i> , 2014, 20, 7894.	3.3	74
62	Aberrant DNA methylation of microRNA genes in human breast cancer – a critical appraisal. <i>Cell and Tissue Research</i> , 2014, 356, 657-664.	2.9	23
63	Deregulation of <i>Rb1</i> expression by loss of imprinting in human hepatocellular carcinoma. <i>Journal of Pathology</i> , 2014, 233, 392-401.	4.5	38
64	Prognostic Impact of Splicing Factor Mutations in Patients with Myelofibrosis Undergoing Allogeneic Hematopoietic Stem Cell Transplantation. <i>Blood</i> , 2014, 124, 3171-3171.	1.4	1
65	Concordant hypermethylation of intergenic microRNA genes in human hepatocellular carcinoma as new diagnostic and prognostic marker. <i>International Journal of Cancer</i> , 2013, 133, 660-670.	5.1	68
66	Oncogenic <i>PIK3CA</i> mutations in lobular breast cancer progression. <i>Genes Chromosomes and Cancer</i> , 2013, 52, 69-80.	2.8	36
67	Archival bone marrow trephines are suitable for high-throughput mutation analysis using next generation sequencing technology. <i>Haematologica</i> , 2013, 98, e115-e116.	3.5	9
68	High intrahepatic HHV-6 virus loads but neither CMV nor EBV are associated with decreased graft survival after diagnosis of graft hepatitis. <i>Journal of Hepatology</i> , 2012, 56, 1063-1069.	3.7	35
69	IPH-926 lobular breast cancer cells harbor a p53 mutant with temperature-sensitive functional activity and allow for profiling of p53-responsive genes. <i>Laboratory Investigation</i> , 2012, 92, 1635-1647.	3.7	26
70	Limited Value of KAI1/CD82 Protein Expression as a Prognostic Marker in Human Gastric Cancer. <i>Disease Markers</i> , 2012, 32, 337-342.	1.3	5
71	microRNA Expression Profiling in Archival Tissue Specimens: Methods and Data Processing. <i>Molecular Biotechnology</i> , 2012, 50, 159-169.	2.4	9
72	Loss of Imprinting and Allelic Switching at the DLK1-MEG3 Locus in Human Hepatocellular Carcinoma. <i>PLoS ONE</i> , 2012, 7, e49462.	2.5	119

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73	High-Resolution Quantitative Methylation Analysis of MicroRNA Genes Using Pyrosequencing. <i>Methods in Molecular Biology</i> , 2012, 878, 229-240.	0.9	1
74	Plexiform Lesions in Pulmonary Arterial Hypertension. <i>American Journal of Pathology</i> , 2011, 179, 167-179.	3.8	144
75	MicroRNA profiles of healthy basal and luminal mammary epithelial cells are distinct and reflected in different breast cancer subtypes. <i>Breast Cancer Research and Treatment</i> , 2011, 130, 735-745.	2.5	117
76	Systematic cross-validation of 454 sequencing and pyrosequencing for the exact quantification of DNA methylation patterns with single CpG resolution. <i>BMC Biotechnology</i> , 2011, 11, 6.	3.3	39
77	MicroRNA Profiling Using Fluorescence-Labeled Beads: Data Acquisition and Processing. <i>Methods in Molecular Biology</i> , 2011, 676, 253-268.	0.9	6
78	Identification of differentially expressed microRNAs in human male breast cancer. <i>BMC Cancer</i> , 2010, 10, 109.	2.6	119
79	Epigenetic inactivation of tumour suppressor gene <i>KLF11</i> in myelodysplastic syndromes*. <i>European Journal of Haematology</i> , 2010, 84, 298-303.	2.2	30
80	Frequent and Distinct Aberrations of DNA Methylation Patterns in Fibrolamellar Carcinoma of the Liver. <i>PLoS ONE</i> , 2010, 5, e13688.	2.5	34
81	MicroRNA-Profilng in Formalin-Fixed Paraffin-Embedded Specimens. <i>Methods in Molecular Biology</i> , 2010, 667, 113-125.	0.9	11
82	Complete cytogenetic remission after decitabine treatment in a patient with secondary AML harbouring high p15 INK4b gene methylation and high global DNA methylation. <i>Annals of Hematology</i> , 2009, 88, 275-277.	1.8	2
83	Comprehensive genetic and functional characterization of IPH926: a novel <i>CDH1</i> null tumour cell line from human lobular breast cancer. <i>Journal of Pathology</i> , 2009, 217, 620-632.	4.5	32
84	Tissue Procurement for Molecular Studies Using Laser-Assisted Microdissection. <i>Methods in Molecular Biology</i> , 2009, 506, 299-310.	0.9	2
85	Reliable microRNA profiling in routinely processed formalin-fixed paraffin-embedded breast cancer specimens using fluorescence labelled bead technology. <i>BMC Biotechnology</i> , 2008, 8, 90.	3.3	43
86	Re: Rahner et al. Coexisting somatic promoter hypermethylation and pathogenic <i>MLH1</i> germline mutation in Lynch syndrome. <i>J Pathol</i> 2008; 214: 10-16. <i>Journal of Pathology</i> , 2008, 215, 97-97.	4.5	0
87	KAI1/CD82 is a novel target of estrogen receptor-mediated gene repression and downregulated in primary human breast cancer. <i>International Journal of Cancer</i> , 2008, 123, 2239-2246.	5.1	33
88	Persistence of Occult Hepatitis B after Removal of the Hepatitis B Virus-Infected Liver. <i>Journal of Infectious Diseases</i> , 2008, 197, 355-360.	4.0	30
89	Epigenetic defects of hepatocellular carcinoma are already found in non-neoplastic liver cells from patients with hereditary haemochromatosis. <i>Human Molecular Genetics</i> , 2007, 16, 1335-1342.	2.9	45
90	Quantitative High-Resolution CpG Island Mapping with Pyrosequencing Reveals Disease-Specific Methylation Patterns of the <i>CDKN2B</i> Gene in Myelodysplastic Syndrome and Myeloid Leukemia. <i>Clinical Chemistry</i> , 2007, 53, 17-23.	3.2	69

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91	Recipient-Derived Neoangiogenesis of Arterioles and Lymphatics in Quilty Lesions of Cardiac Allografts. <i>Transplantation</i> , 2007, 84, 1335-1342.	1.0	26
92	Laser-Assisted Microdissection and Isolation of DNA and RNA. , 2006, 120, 65-76.		5
93	Bone marrow infiltration pattern in B-cell chronic lymphocytic leukemia is related to immunoglobulin heavy-chain variable region mutation status and expression of 70-kd $\hat{\imath}$ -associated protein (ZAP-70). <i>Human Pathology</i> , 2006, 37, 1153-1161.	2.0	24
94	Testing the importance of p27 degradation by the SCFskp2 pathway in murine models of lung and colon cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 14009-14014.	7.1	62
95	High-Resolution Genomic Profiling Reveals Association of Chromosomal Aberrations on 1q and 16p with Histologic and Genetic Subgroups of Invasive Breast Cancer. <i>Clinical Cancer Research</i> , 2006, 12, 345-352.	7.0	85
96	LEF-1 is crucial for neutrophil granulocytopoiesis and its expression is severely reduced in congenital neutropenia. <i>Nature Medicine</i> , 2006, 12, 1191-1197.	30.7	1,015
97	Up-regulation of DNA methyltransferases DNMT1, 3A, and 3B in myelodysplastic syndrome. <i>Leukemia Research</i> , 2005, 29, 325-329.	0.8	40
98	Absence of p21CIP1, p27KIP1 and p57KIP2 methylation in MDS and AML. <i>Leukemia Research</i> , 2005, 29, 1357-1360.	0.8	39
99	Hypermethylation of the suppressor of cytokine signalling-1 (SOCS-1) in myelodysplastic syndrome. <i>British Journal of Haematology</i> , 2005, 130, 209-217.	2.5	57
100	Down-regulation of the IGF-2/H19 locus during normal and malignant hematopoiesis is independent of the imprinting pattern. <i>International Journal of Oncology</i> , 2005, 26, 499.	3.3	7
101	Distinct Methylation Patterns of Benign and Malignant Liver Tumors Revealed by Quantitative Methylation Profiling. <i>Clinical Cancer Research</i> , 2005, 11, 3654-3660.	7.0	60
102	Down-regulation of the IGF-2/H19 locus during normal and malignant hematopoiesis is independent of the imprinting pattern. <i>International Journal of Oncology</i> , 2005, 26, 499-507.	3.3	16
103	Combined laser-assisted microdissection and short tandem repeat analysis for detection of in situ microchimerism after solid organ transplantation. <i>Methods in Molecular Biology</i> , 2005, 293, 113-23.	0.9	3
104	Role of epigenetic changes in hematological malignancies. <i>Annals of Hematology</i> , 2004, 83, 137-152.	1.8	35
105	Overexpression of delta-like (Dlk) in a subset of myelodysplastic syndrome bone marrow trephines. <i>Leukemia Research</i> , 2004, 28, 1081-1083.	0.8	20
106	Real-Time PCR-Based Assay for Quantitative Determination of Methylation Status. , 2004, 287, 207-218.		12
107	Low level of DAP-kinase DNA methylation in myelodysplastic syndrome. <i>Blood</i> , 2004, 104, 1586-1588.	1.4	13
108	Quantitative Intra-Individual Monitoring of BCR-ABL Transcript Levels in Archival Bone Marrow Trepines of Patients with Chronic Myeloid Leukemia. <i>Journal of Molecular Diagnostics</i> , 2003, 5, 54-60.	2.8	24

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109	Megakaryocytes from chronic myeloproliferative disorders show enhanced nuclear bFGF expression. <i>Blood</i> , 2002, 100, 2274-2275.	1.4	26
110	Quantitative Assessment of Promoter Hypermethylation during Breast Cancer Development. <i>American Journal of Pathology</i> , 2002, 160, 605-612.	3.8	210
111	Promoter hypermethylation of the death-associated protein kinase gene in breast cancer is associated with the invasive lobular subtype. <i>Cancer Research</i> , 2002, 62, 6634-8.	0.9	58
112	Real-Time PCR Analysis of DNA and RNA Extracted from Formalin-Fixed and Paraffin-Embedded Biopsies. <i>Methods</i> , 2001, 25, 409-418.	3.8	336
113	Demonstration of Light Chain Restricted Clonal B-Lymphoid Infiltrates in Archival Bone Marrow Trephines by Quantitative Real-Time Polymerase Chain Reaction. <i>American Journal of Pathology</i> , 2001, 159, 2023-2029.	3.8	19
114	One-Step Extraction of RNA from Archival Biopsies. <i>Analytical Biochemistry</i> , 2001, 295, 116-117.	2.4	44
115	Amplification of Growth Regulatory Genes in Intraductal Breast Cancer Is Associated with Higher Nuclear Grade but Not with the Progression to Invasiveness. <i>Laboratory Investigation</i> , 2001, 81, 565-571.	3.7	35
116	Quantitative Analysis of Promoter Hypermethylation in Laser-Microdissected Archival Specimens. <i>Laboratory Investigation</i> , 2001, 81, 635-637.	3.7	45
117	Positive Display of Methylated Sites: A Novel Method for the Detection of Promoter Methylation. <i>Diagnostic Molecular Pathology</i> , 2000, 9, 165-171.	2.1	2
118	Common and epithelioid variants of hepatic angiomyolipoma exhibit clonal growth and share a distinctive immunophenotype. <i>Hepatology</i> , 2000, 32, 213-217.	7.3	74
119	Quantitative Molecular Analysis of Laser-Microdissected Paraffin- Embedded Human Tissues. <i>Pathobiology</i> , 2000, 68, 202-208.	3.8	41
120	Detection of Gene Amplification in Archival Breast Cancer Specimens by Laser-Assisted Microdissection and Quantitative Real-Time Polymerase Chain Reaction. <i>American Journal of Pathology</i> , 2000, 156, 1855-1864.	3.8	138
121	Polymerase chain reaction-based analyses of nucleic acids from archival material. , 0, , 254-261.		0