

# Bernhard F Radlwimmer

## List of Publications by Year in descending order

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122  
papers

19,838  
citations

15504

65  
h-index

22832

112  
g-index

128  
all docs

128  
docs citations

128  
times ranked

29361  
citing authors

#	ARTICLE	IF	CITATIONS
1	International network of cancer genome projects. <i>Nature</i> , 2010, 464, 993-998.	27.8	2,114
2	DNA methylation-based classification of central nervous system tumours. <i>Nature</i> , 2018, 555, 469-474.	27.8	1,872
3	Hotspot Mutations in H3F3A and IDH1 Define Distinct Epigenetic and Biological Subgroups of Glioblastoma. <i>Cancer Cell</i> , 2012, 22, 425-437.	16.8	1,551
4	An endogenous tumour-promoting ligand of the human aryl hydrocarbon receptor. <i>Nature</i> , 2011, 478, 197-203.	27.8	1,514
5	Reduced H3K27me3 and DNA Hypomethylation Are Major Drivers of Gene Expression in K27M Mutant Pediatric High-Grade Gliomas. <i>Cancer Cell</i> , 2013, 24, 660-672.	16.8	633
6	Stem Cell Marker CD133 Affects Clinical Outcome in Glioma Patients. <i>Clinical Cancer Research</i> , 2008, 14, 123-129.	7.0	540
7	A Chromosome 8 Gene-Cluster Polymorphism with Low Human Beta-Defensin 2 Gene Copy Number Predisposes to Crohn Disease of the Colon. <i>American Journal of Human Genetics</i> , 2006, 79, 439-448.	6.2	487
8	Atypical Teratoid/Rhabdoid Tumors Are Comprised of Three Epigenetic Subgroups with Distinct Enhancer Landscapes. <i>Cancer Cell</i> , 2016, 29, 379-393.	16.8	438
9	BRAF gene duplication constitutes a mechanism of MAPK pathway activation in low-grade astrocytomas. <i>Journal of Clinical Investigation</i> , 2008, 118, 1739-1749.	8.2	437
10	Recurrent mutation of the ID3 gene in Burkitt lymphoma identified by integrated genome, exome and transcriptome sequencing. <i>Nature Genetics</i> , 2012, 44, 1316-1320.	21.4	389
11	BCAT1 promotes cell proliferation through amino acid catabolism in gliomas carrying wild-type IDH1. <i>Nature Medicine</i> , 2013, 19, 901-908.	30.7	388
12	Decoding the regulatory landscape of medulloblastoma using DNA methylation sequencing. <i>Nature</i> , 2014, 510, 537-541.	27.8	378
13	Longitudinal molecular trajectories of diffuse glioma in adults. <i>Nature</i> , 2019, 576, 112-120.	27.8	320
14	Differentiation Therapy Exerts Antitumor Effects on Stem-like Glioma Cells. <i>Clinical Cancer Research</i> , 2010, 16, 2715-2728.	7.0	279
15	MYC High Level Gene Amplification Is a Distinctive Feature of Angiosarcomas after Irradiation or Chronic Lymphedema. <i>American Journal of Pathology</i> , 2010, 176, 34-39.	3.8	276
16	Outcome Prediction in Pediatric Medulloblastoma Based on DNA Copy-Number Aberrations of Chromosomes 6q and 17q and the <i>MYC</i> and <i>MYCN</i> Loci. <i>Journal of Clinical Oncology</i> , 2009, 27, 1627-1636.	1.6	274
17	BCAT1 restricts $\hat{\pm}$ KG levels in AML stem cells leading to IDHmut-like DNA hypermethylation. <i>Nature</i> , 2017, 551, 384-388.	27.8	261
18	Molecular classification of diffuse cerebral WHO grade II/III gliomas using genome- and transcriptome-wide profiling improves stratification of prognostically distinct patient groups. <i>Acta Neuropathologica</i> , 2015, 129, 679-693.	7.7	254

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19	Automated array-based genomic profiling in chronic lymphocytic leukemia: Development of a clinical tool and discovery of recurrent genomic alterations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 1039-1044.	7.1	221
20	Identification of Gains on 1q and Epidermal Growth Factor Receptor Overexpression as Independent Prognostic Markers in Intracranial Ependymoma. <i>Clinical Cancer Research</i> , 2006, 12, 2070-2079.	7.0	212
21	Genomic and Protein Expression Profiling Identifies CDK6 As Novel Independent Prognostic Marker in Medulloblastoma. <i>Journal of Clinical Oncology</i> , 2005, 23, 8853-8862.	1.6	207
22	Apoptosis-based treatment of glioblastomas with ABT-737, a novel small molecule inhibitor of Bcl-2 family proteins. <i>Oncogene</i> , 2008, 27, 6646-6656.	5.9	188
23	Integrated DNA methylation and copy-number profiling identify three clinically and biologically relevant groups of anaplastic glioma. <i>Acta Neuropathologica</i> , 2014, 128, 561-571.	7.7	176
24	Etiology-dependent molecular mechanisms in human hepatocarcinogenesis. <i>Hepatology</i> , 2008, 47, 511-520.	7.3	173
25	Evolutionary Trajectories of IDHWT Glioblastomas Reveal a Common Path of Early Tumorigenesis Instigated Years ahead of Initial Diagnosis. <i>Cancer Cell</i> , 2019, 35, 692-704.e12.	16.8	172
26	The "five-sites" rule and the evolution of red and green color vision in mammals. <i>Molecular Biology and Evolution</i> , 1998, 15, 560-567.	8.9	162
27	Tagmentation-based whole-genome bisulfite sequencing. <i>Nature Protocols</i> , 2013, 8, 2022-2032.	12.0	161
28	Genomic and Expression Profiling of Glioblastoma Stem Cell-Like Spheroid Cultures Identifies Novel Tumor-Relevant Genes Associated with Survival. <i>Clinical Cancer Research</i> , 2009, 15, 6541-6550.	7.0	158
29	Adaptive evolution of color vision of the Comoran coelacanth ( <i>Latimeria chalumnae</i> ). <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1999, 96, 6279-6284.	7.1	147
30	Molecular genetics and the evolution of ultraviolet vision in vertebrates. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 11731-11736.	7.1	147
31	Recurrent FGFR1 amplification and high FGFR1 protein expression in oral squamous cell carcinoma (OSCC). <i>Oral Oncology</i> , 2007, 43, 60-66.	1.5	147
32	De-repression of CTGF via the miR-17-92 cluster upon differentiation of human glioblastoma spheroid cultures. <i>Oncogene</i> , 2010, 29, 3411-3422.	5.9	142
33	Disclosure of Candidate Genes in Acute Myeloid Leukemia With Complex Karyotypes Using Microarray-Based Molecular Characterization. <i>Journal of Clinical Oncology</i> , 2006, 24, 3887-3894.	1.6	141
34	Genomic DNA-Chip Hybridization Reveals a Higher Incidence of Genomic Amplifications in Pancreatic Cancer than Conventional Comparative Genomic Hybridization and Leads to the Identification of Novel Candidate Genes. <i>Cancer Research</i> , 2004, 64, 4428-4433.	0.9	140
35	Severe mental retardation with breathing abnormalities (Pitt-Hopkins syndrome) is caused by haploinsufficiency of the neuronal bHLH transcription factor TCF4. <i>Human Molecular Genetics</i> , 2007, 16, 1488-1494.	2.9	137
36	Methylome analysis and integrative profiling of human HCCs identify novel protumorigenic factors. <i>Hepatology</i> , 2012, 56, 1817-1827.	7.3	136

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37	Epigenetic Upregulation of lncRNAs at 13q14.3 in Leukemia Is Linked to the In Cis Downregulation of a Gene Cluster That Targets NF- $\kappa$ B. <i>PLoS Genetics</i> , 2013, 9, e1003373.	3.5	134
38	Comprehensive genomic analysis of desmoplastic medulloblastomas: identification of novel amplified genes and separate evaluation of the different histological components. <i>Journal of Pathology</i> , 2006, 208, 554-563.	4.5	129
39	The Endogenous Tryptophan Metabolite and NAD <sup>+</sup> Precursor Quinolinic Acid Confers Resistance of Gliomas to Oxidative Stress. <i>Cancer Research</i> , 2013, 73, 3225-3234.	0.9	126
40	RNAi screening in glioma stem-like cells identifies PFKFB4 as a key molecule important for cancer cell survival. <i>Oncogene</i> , 2012, 31, 3235-3243.	5.9	123
41	Targeting Self-Renewal in High-Grade Brain Tumors Leads to Loss of Brain Tumor Stem Cells and Prolonged Survival. <i>Cell Stem Cell</i> , 2014, 15, 185-198.	11.1	123
42	Ultraviolet pigments in birds evolved from violet pigments by a single amino acid change. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000, 97, 7366-7371.	7.1	121
43	Genomic DNA-chip hybridization in t(11;14)-positive mantle cell lymphomas shows a high frequency of aberrations and allows a refined characterization of consensus regions. <i>Blood</i> , 2004, 104, 795-801.	1.4	121
44	The nuclear receptor <i>tailless</i> induces long-term neural stem cell expansion and brain tumor initiation. <i>Genes and Development</i> , 2010, 24, 683-695.	5.9	121
45	DNA methylome analysis in Burkitt and follicular lymphomas identifies differentially methylated regions linked to somatic mutation and transcriptional control. <i>Nature Genetics</i> , 2015, 47, 1316-1325.	21.4	119
46	Molecular characterization of long-term survivors of glioblastoma using genome- and transcriptome-wide profiling. <i>International Journal of Cancer</i> , 2014, 135, 1822-1831.	5.1	117
47	Angiocentric Glioma. <i>American Journal of Surgical Pathology</i> , 2007, 31, 1709-1718.	3.7	110
48	Recurrent copy number gain of transcription factor <i>SOX2</i> and corresponding high protein expression in oral squamous cell carcinoma. <i>Genes Chromosomes and Cancer</i> , 2010, 49, 9-16.	2.8	106
49	High-resolution genomic profiling of childhood T-ALL reveals frequent copy-number alterations affecting the TGF- $\beta$ 2 and PI3K-AKT pathways and deletions at 6q15-16.1 as a genomic marker for unfavorable early treatment response. <i>Blood</i> , 2009, 114, 1053-1062.	1.4	105
50	Detection of chromosomal imbalances in retinoblastoma by matrix-based comparative genomic hybridization. <i>Genes Chromosomes and Cancer</i> , 2005, 43, 294-301.	2.8	101
51	Genomic and transcriptomic changes complement each other in the pathogenesis of sporadic Burkitt lymphoma. <i>Nature Communications</i> , 2019, 10, 1459.	12.8	99
52	Frequent loss of chromosome 9, homozygous CDKN2A/p14ARF/CDKN2B deletion and low TSC1 mRNA expression in pleomorphic xanthoastrocytomas. <i>Oncogene</i> , 2007, 26, 1088-1097.	5.9	98
53	Candidate genes in breast cancer revealed by microarray-based comparative genomic hybridization of archived tissue. <i>Cancer Research</i> , 2005, 65, 439-47.	0.9	98
54	MINCR is a MYC-induced lncRNA able to modulate MYC's transcriptional network in Burkitt lymphoma cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E5261-70.	7.1	91

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55	Regeneration of ultraviolet pigments of vertebrates. <i>FEBS Letters</i> , 1998, 423, 155-158.	2.8	88
56	Expression of an ASCL2 related stem cell signature and IGF2 in colorectal cancer liver metastases with 11p15.5 gain. <i>Gut</i> , 2010, 59, 1236-1244.	12.1	88
57	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
58	DNA microarray analysis identifies candidate regions and genes in unexplained mental retardation. <i>Neurology</i> , 2007, 68, 743-750.	1.1	82
59	The Wnt secretion protein Evi/Gpr177 promotes glioma tumourigenesis. <i>EMBO Molecular Medicine</i> , 2012, 4, 38-51.	6.9	81
60	<sc>LGR5</sc> is a Marker of Poor Prognosis in Glioblastoma and is Required for Survival of Brain Cancer Stemâ€Like Cells. <i>Brain Pathology</i> , 2013, 23, 60-72.	4.1	80
61	DNA dot-blot hybridization implicates human papillomavirus type 11-DNA in epithelioma cuniculatum. <i>Journal of Medical Virology</i> , 1989, 29, 33-37.	5.0	77
62	Supratentorial primitive neuroectodermal tumors of the central nervous system frequently harbor deletions of theCDKN2A locus and other genomic aberrations distinct from medulloblastomas. <i>Genes Chromosomes and Cancer</i> , 2007, 46, 839-851.	2.8	76
63	Molecular signatures classify astrocytic gliomas by <i>IDH1</i> mutation status. <i>International Journal of Cancer</i> , 2011, 128, 1095-1103.	5.1	75
64	EEF1A2 inactivates p53 by way of PI3K/AKT/mTOR-dependent stabilization of MDM4 in hepatocellular carcinoma. <i>Hepatology</i> , 2014, 59, 1886-1899.	7.3	74
65	Branchedâ€chain ketoacids secreted by glioblastoma cells via <sc>MCT</sc> 1 modulate macrophage phenotype. <i>EMBO Reports</i> , 2017, 18, 2172-2185.	4.5	74
66	Co-localization of CENP-C and CENP-H to discontinuous domains of CENP-A chromatin at human neocentromeres. <i>Genome Biology</i> , 2007, 8, R148.	9.6	73
67	Microglia isolated from patients with glioma gain antitumor activities on poly (I:C) stimulation. <i>Neuro-Oncology</i> , 2012, 14, 64-78.	1.2	66
68	GPD1 Specifically Marks Dormant Glioma Stem Cells with a Distinct Metabolic Profile. <i>Cell Stem Cell</i> , 2019, 25, 241-257.e8.	11.1	66
69	Recurrent coamplification of cytoskeletonâ€associated genes <i>EMS1</i> and <i>SHANK2</i> with <i>CCND1</i> in oral squamous cell carcinoma. <i>Genes Chromosomes and Cancer</i> , 2006, 45, 118-125.	2.8	60
70	Mapping candidate regions and genes for congenital anomalies of the kidneys and urinary tract (CAKUT) by array-based comparative genomic hybridization. <i>Nephrology Dialysis Transplantation</i> , 2011, 26, 136-143.	0.7	60
71	The branched-chain amino acid transaminase 1 sustains growth of antiestrogen-resistant and ERÎ±-negative breast cancer. <i>Oncogene</i> , 2017, 36, 4124-4134.	5.9	60
72	Array-based profiling of reference-independent methylation status (aPRIMES) identifies frequent promoter methylation and consecutive downregulation of ZIC2 in pediatric medulloblastoma. <i>Nucleic Acids Research</i> , 2007, 35, e51.	14.5	58

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73	Comprehensive Characterization of Genomic Aberrations in Gangliogliomas by CGH, Array-based CGH and Interphase FISH. <i>Brain Pathology</i> , 2008, 18, 326-337.	4.1	58
74	Expression of podoplanin in human astrocytic brain tumors is controlled by the PI3K-AKT-AP-1 signaling pathway and promoter methylation. <i>Neuro-Oncology</i> , 2012, 14, 426-439.	1.2	55
75	Frequent amplifications and abundant expression of TRIO, NKD2, and IRX2 in soft tissue sarcomas. <i>Genes Chromosomes and Cancer</i> , 2006, 45, 829-838.	2.8	52
76	The genomic and transcriptional landscape of primary central nervous system lymphoma. <i>Nature Communications</i> , 2022, 13, 2558.	12.8	52
77	Mutagenesis and reconstitution of middle-to-long-wave-sensitive visual pigments of New World monkeys for testing the tuning effect of residues at sites 229 and 233. <i>Vision Research</i> , 2004, 44, 2225-2231.	1.4	49
78	Glioblastoma epigenome profiling identifies SOX10 as a master regulator of molecular tumour subtype. <i>Nature Communications</i> , 2020, 11, 6434.	12.8	48
79	Primary glioblastoma cultures: can profiling of stem cell markers predict radiotherapy sensitivity?. <i>Journal of Neurochemistry</i> , 2014, 131, 251-264.	3.9	47
80	Differential Retinoic Acid Signaling in Tumors of Long- and Short-term Glioblastoma Survivors. <i>Journal of the National Cancer Institute</i> , 2011, 103, 598-601.	6.3	46
81	Towards mapping phenotypical traits in 18p <sup>+</sup> syndrome by array-based comparative genomic hybridisation and fluorescent in situ hybridisation. <i>European Journal of Human Genetics</i> , 2007, 15, 35-44.	2.8	42
82	Isochromosome breakpoints on 17p in medulloblastoma are flanked by different classes of DNA sequence repeats. <i>Genes Chromosomes and Cancer</i> , 2006, 45, 401-410.	2.8	41
83	Establishment, characterization and drug sensitivity testing in primary cultures of human thymoma and thymic carcinoma. <i>International Journal of Cancer</i> , 2008, 122, 2719-2725.	5.1	41
84	Loss of FUBP1 expression in gliomas predicts FUBP1 mutation and is associated with oligodendroglial differentiation, IDH1 mutation and 1p/19q loss of heterozygosity. <i>Neuropathology and Applied Neurobiology</i> , 2014, 40, 205-216.	3.2	41
85	<i>FARP2</i> , <i>HDLBP</i> and <i>PASK</i> are downregulated in a patient with autism and 2q37.3 deletion syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2009, 149A, 952-959.	1.2	40
86	Molecular evolution of color vision of zebra finch. <i>Gene</i> , 2000, 259, 17-24.	2.2	37
87	Genetic analyses of the green visual pigments of rabbit ( <i>Oryctolagus cuniculus</i> ) and rat ( <i>Rattus</i> ) Tj ETQq1 1 0.784314 rgBT / Overlock 10	2.2	36
88	Molecular classification of human gliomas using matrix-based comparative genomic hybridization. <i>International Journal of Cancer</i> , 2005, 117, 95-103.	5.1	36
89	Genomic and spectral analyses of long to middle wavelength-sensitive visual pigments of common marmoset ( <i>Callithrix jacchus</i> ). <i>Gene</i> , 2001, 269, 45-51.	2.2	34
90	The onset of p53 loss of heterozygosity is differentially induced in various stem cell types and may involve the loss of either allele. <i>Cell Death and Differentiation</i> , 2014, 21, 1419-1431.	11.2	34

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91	Mutational mechanisms shaping the coding and noncoding genome of germinal center derived B-cell lymphomas. <i>Leukemia</i> , 2021, 35, 2002-2016.	7.2	34
92	Expansive growth of two glioblastoma stem-like cell lines is mediated by bFGF and not by EGF. <i>Radiology and Oncology</i> , 2013, 47, 330-337.	1.7	29
93	PII Protein-Derived FRET Sensors for Quantification and Live-Cell Imaging of 2-Oxoglutarate. <i>Scientific Reports</i> , 2017, 7, 1437.	3.3	29
94	Gilvocarcin V exhibits both equilibrium DNA binding and UV light induced DNA adduct formation which is sequence context dependent. <i>Nucleic Acids Research</i> , 1992, 20, 4553-4557.	14.5	22
95	Copy number alterations in childhood acute lymphoblastic leukemia and their association with minimal residual disease. <i>Genes Chromosomes and Cancer</i> , 2008, 47, 471-480.	2.8	21
96	Sensitivity and resistance towards isoliquiritigenin, doxorubicin and methotrexate in T cell acute lymphoblastic leukaemia cell lines by pharmacogenomics. <i>Naunyn-Schmiedeberg's Archives of Pharmacology</i> , 2010, 382, 221-234.	3.0	20
97	Genomic profiling reveals distinctive molecular relapse patterns in <i>IDH1/2</i> wild-type glioblastoma. <i>Genes Chromosomes and Cancer</i> , 2014, 53, 589-605.	2.8	18
98	Matrix-comparative genomic hybridization from multicenter formalin-fixed paraffin-embedded colorectal cancer tissue blocks. <i>BMC Cancer</i> , 2007, 7, 58.	2.6	17
99	Leucine and branched-chain amino acid metabolism contribute to the growth of bone sarcomas by regulating AMPK and mTORC1 signaling. <i>Biochemical Journal</i> , 2020, 477, 1579-1599.	3.7	17
100	A gene signature distinguishing CD133hi from CD133- colorectal cancer cells: essential role for EGR1 and downstream factors. <i>Pathology</i> , 2011, 43, 220-227.	0.6	16
101	Cloning and expression of the red visual pigment gene of goat ( <i>Capra hircus</i> ). <i>Gene</i> , 1997, 198, 211-215.	2.2	15
102	A Set of Cell Lines Derived from a Genetic Murine Glioblastoma Model Recapitulates Molecular and Morphological Characteristics of Human Tumors. <i>Cancers</i> , 2021, 13, 230.	3.7	13
103	Genome-wide analysis for micro-aberrations in familial exstrophy of the bladder using array-based comparative genomic hybridization. <i>BJU International</i> , 2007, 100, 646-650.	2.5	12
104	Rosetted glioneuronal tumor of the spine with overtly anaplastic histological features. <i>Acta Neuropathologica</i> , 2009, 117, 591-593.	7.7	10
105	Pilocytic astrocytoma demethylation and transcriptional landscapes link bZIP transcription factors to immune response. <i>Neuro-Oncology</i> , 2020, 22, 1327-1338.	1.2	10
106	Decreased expression of the mitochondrial BCAT protein correlates with improved patient survival in IDH-wt gliomas. <i>Brain Pathology</i> , 2016, 26, 789-791.	4.1	8
107	Array-CGH in unclear syndromic nephropathies identifies a microdeletion in Xq22.3-q23. <i>Pediatric Nephrology</i> , 2009, 24, 1673-1681.	1.7	7
108	Gilvocarcin V exhibits both equilibrium DNA binding and UV light induced DNA adduct formation which is sequence context dependent. <i>Nucleic Acids Research</i> , 1993, 21, 3920-3920.	14.5	4

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109	Focal structural variants revealed by whole genome sequencing disrupt the histone demethylase KDM4C in B-cell lymphomas. <i>Haematologica</i> , 2023, 108, 543-554.	3.5	2
110	Human papillomavirus in cervix carcinoma and condylomata acuminata-identification of HPV-DNA by improved dot-blot-hybridization. <i>Clinical and Experimental Dermatology</i> , 1992, 17, 392-396.	1.3	1
111	348: BCAT1 promotes cell proliferation through amino acid catabolism in gliomas carrying wild-type IDH1. <i>European Journal of Cancer</i> , 2014, 50, S83.	2.8	1
112	Abstract 3240: The Wnt secretion factor Evi/Wls/Gpr177 promotes glioma tumorigenesis. , 2012, , .		1
113	P42: High resolution genomic profiling reveals association of chromosomal aberrations on 1q and 16p with histological and genetic subgroups of invasive breast cancer. <i>European Journal of Medical Genetics</i> , 2005, 48, 506-507.	1.3	0
114	P45: Comprehensive genomic analysis of desmoplastic medulloblastomas reveals novel amplified genes and supports a monoclonal origin of the different histologic components. <i>European Journal of Medical Genetics</i> , 2005, 48, 509.	1.3	0
115	32LBA High level gene amplification of MYC characterizes radiation-induced angiosarcomas. <i>European Journal of Cancer, Supplement</i> , 2009, 7, 16.	2.2	0
116	MOLECULAR GENETIC DETERMINANTS OF LONG-TERM SURVIVAL WITH GLIOBLASTOMA. <i>Neuro-Oncology</i> , 2014, 16, iii46-iii47.	1.2	0
117	294: In vitro modulation of CITED4 gene expression in a colorectal cancer cell line. <i>European Journal of Cancer</i> , 2014, 50, S69-S70.	2.8	0
118	COMPREHENSIVE EPIGENETIC AND TRANSCRIPTIONAL SURVEY OF THE "IMPRINTOME" IN NORMAL B-CELLS AND GERMINAL CENTER DERIVED B-CELL LYMPHOMAS OF THE MMML AND ICGC MMML-SEQ NETWORKS. <i>Hematological Oncology</i> , 2017, 35, 155-156.	1.7	0
119	Characterization of Secondary Alterations in Mantle Cell Lymphomas Using Matrix-/Array CGH.. <i>Blood</i> , 2004, 104, 4349-4349.	1.4	0
120	Automated Screening for Genomic Imbalances in Multiple Myeloma Using Microarray-Based Comparative Genomic Hybridization (mCGH).. <i>Blood</i> , 2005, 106, 499-499.	1.4	0
121	Molecular risk stratification in pediatric medulloblastoma. <i>Journal of Clinical Oncology</i> , 2007, 25, 9506-9506.	1.6	0
122	Abstract 3797: Regulation and function of the mucin-like glycoprotein podoplanin in glioma.. , 2013, , .		0