

Richard John Byers

List of Publications by Year in descending order

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25
papers

623
citations

759233
12
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24
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27
all docs

27
docs citations

27
times ranked

941
citing authors

#	ARTICLE	IF	CITATIONS
1	The impact of obesity and bariatric surgery on the immune microenvironment of the endometrium. International Journal of Obesity, 2022, 46, 605-612.	3.4	20
2	Immune infiltrate diversity confers a good prognosis in follicular lymphoma. Cancer Immunology, Immunotherapy, 2021, 70, 3573-3585.	4.2	8
3	Spatial proximity between T and PD-L1 expressing cells as a prognostic biomarker for oropharyngeal squamous cell carcinoma. British Journal of Cancer, 2020, 122, 539-544.	6.4	35
4	Measurement of indicator genes using global complementary DNA (cDNA) amplification, by polyadenylic acid reverse transcriptase polymerase chain reaction (poly A RT-PCR): A feasibility study using paired samples from tissue and ductal juice in patients undergoing pancreatoduodenectomy. Pancreatology, 2018, 18, 458-462.	1.1	2
5	Single-cell analysis of K562 cells: An imatinib-resistant subpopulation is adherent and has upregulated expression of BCR-ABL mRNA and protein. Experimental Hematology, 2014, 42, 183-191.e5.	0.4	20
6	Guidelines and considerations for conducting experiments using tissue microarrays. Histopathology, 2013, 62, 827-839.	2.9	57
7	Detection of IDH1 R132H Mutation in Acute Myeloid Leukemia by Mutation-specific Immunohistochemistry. Applied Immunohistochemistry and Molecular Morphology, 2012, 20, 37-40.	1.2	8
8	Microarray Gene Expression Analysis of Fixed Archival Tissue Permits Molecular Classification and Identification of Potential Therapeutic Targets in Diffuse Large B-Cell Lymphoma. Journal of Molecular Diagnostics, 2012, 14, 223-232.	2.8	24
9	Quantitative multiplex quantum dot in-situ hybridisation based gene expression profiling in tissue microarrays identifies prognostic genes in acute myeloid leukaemia. Biochemical and Biophysical Research Communications, 2012, 425, 333-339.	2.1	12
10	A comparative study of quantitative immunohistochemistry and quantum dot immunohistochemistry for mutation carrier identification in Lynch syndrome. Journal of Clinical Pathology, 2011, 64, 208-214.	2.0	15
11	Impact of gene expression profiling in lymphoma diagnosis and prognosis. Histopathology, 2011, 58, 106-127.	2.9	12
12	Effect of prolonged formalin fixation on immunohistochemical staining for the proliferation marker Ki67. Histopathology, 2011, 59, 1261-1263.	2.9	11
13	Nodal B-cell lymphoma and Hodgkin lymphoma: current WHO classification and differential diagnostic dilemmas. Diagnostic Histopathology, 2010, 16, 54-68.	0.4	6
14	Massive haemorrhagic adrenal metastases leading to sudden death: a case report. BMJ Case Reports, 2009, 2009, bcr0620080190-bcr0620080190.	0.5	10
15	Molecular diagnostics in haematopathology. Diagnostic Histopathology, 2008, 14, 223-235.	0.4	4
16	Molecular diagnosis in pancreatic cancer. Diagnostic Histopathology, 2008, 14, 214-222.	0.4	5
17	Blocks to paraffin get the CHOP, +R. Blood, 2008, 112, 3006-3007.	1.4	1
18	Clinical quantitation of immune signature in follicular lymphoma by RT-PCR-based gene expression profiling. Blood, 2008, 111, 4764-4770.	1.4	84

#	ARTICLE	IF	CITATIONS
19	Clinical quantitation of diagnostic and predictive gene expression levels in follicular and diffuse large B-cell lymphoma by RT-PCR gene expression profiling. Blood, 2007, 109, 3922-3928.	1.4	83
20	Imaging of multiple mRNA targets using quantum dot based in situ hybridization and spectral deconvolution in clinical biopsies. Biochemical and Biophysical Research Communications, 2006, 348, 628-636.	2.1	73
21	Gene expression analysis of myeloid and lymphoid lineage markers during mouse haematopoiesis. British Journal of Haematology, 2006, 135, 105-116.	2.5	8
22	Routine expression profiling of microarray gene signatures in acute leukaemia by real-time PCR of human bone marrow*. British Journal of Haematology, 2005, 130, 233-248.	2.5	33
23	Quantum Dot Based Duplex In Situ Hybridisation for Gene Expression Profiling.. Blood, 2005, 106, 3265-3265.	1.4	0
24	Notch Oncoproteins Depend on β -Secretase/Presenilin Activity for Processing and Function. Journal of Biological Chemistry, 2004, 279, 30771-30780.	3.4	67
25	PolyA PCR Amplification of cDNA from RNA Extracted from Formalin-Fixed Paraffin-Embedded Tissue. Diagnostic Molecular Pathology, 2004, 13, 144-150.	2.1	22