

Bas Kreike

List of Publications by Year in descending order

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Version: 2024-02-01

22
papers

3,250
citations

331642

21
h-index

642715

23
g-index

23
all docs

23
docs citations

23
times ranked

5720
citing authors

#	ARTICLE	IF	CITATIONS
1	Search for a Gene Expression Signature of Breast Cancer Local Recurrence in Young Women. <i>Clinical Cancer Research</i> , 2012, 18, 1704-1715.	7.0	67
2	Functional characterization of the 19q12 amplicon in grade III breast cancers. <i>Breast Cancer Research</i> , 2012, 14, R53.	5.0	61
3	Divergent effects of insulin-like growth factor-1 receptor expression on prognosis of estrogen receptor positive versus triple negative invasive ductal breast carcinoma. <i>Breast Cancer Research and Treatment</i> , 2011, 129, 725-736.	2.5	53
4	Microarray-Based Class Discovery for Molecular Classification of Breast Cancer: Analysis of Interobserver Agreement. <i>Journal of the National Cancer Institute</i> , 2011, 103, 662-673.	6.3	121
5	Engagement of I-Branching β -1, 6-N-Acetylglucosaminyltransferase 2 in Breast Cancer Metastasis and TGF- β Signaling. <i>Cancer Research</i> , 2011, 71, 4846-4856.	0.9	73
6	MicroRNA Sequence and Expression Analysis in Breast Tumors by Deep Sequencing. <i>Cancer Research</i> , 2011, 71, 4443-4453.	0.9	331
7	Analysis of breast cancer related gene expression using natural splines and the Cox proportional hazard model to identify prognostic associations. <i>Breast Cancer Research and Treatment</i> , 2010, 122, 711-720.	2.5	22
8	The molecular underpinning of lobular histological growth pattern: a genome-wide transcriptomic analysis of invasive lobular carcinomas and grade- and molecular subtype-matched invasive ductal carcinomas of no special type. <i>Journal of Pathology</i> , 2010, 220, 45-57.	4.5	208
9	Transcriptomic analysis of tubular carcinomas of the breast reveals similarities and differences with molecular subtype-matched ductal and lobular carcinomas. <i>Journal of Pathology</i> , 2010, 222, 64-75.	4.5	33
10	The Snf1-related kinase, Hunk, is essential for mammary tumor metastasis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 15855-15860.	7.1	31
11	Local Recurrence after Breast-Conserving Therapy in Relation to Gene Expression Patterns in a Large Series of Patients. <i>Clinical Cancer Research</i> , 2009, 15, 4181-4190.	7.0	78
12	Metaplastic breast carcinomas are basal-like breast cancers: a genomic profiling analysis. <i>Breast Cancer Research and Treatment</i> , 2009, 117, 273-280.	2.5	208
13	Mucinous and neuroendocrine breast carcinomas are transcriptionally distinct from invasive ductal carcinomas of no special type. <i>Modern Pathology</i> , 2009, 22, 1401-1414.	5.5	110
14	Continuing Risk of Ipsilateral Breast Relapse After Breast-Conserving Therapy at Long-Term Follow-up. <i>International Journal of Radiation Oncology Biology Physics</i> , 2008, 71, 1014-1021.	0.8	90
15	An interferon-related gene signature for DNA damage resistance is a predictive marker for chemotherapy and radiation for breast cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 18490-18495.	7.1	484
16	The Chemokine Receptor CXCR6 and Its Ligand CXCL16 Are Expressed in Carcinomas and Inhibit Proliferation. <i>Cancer Research</i> , 2008, 68, 4701-4708.	0.9	47
17	Lung metastasis genes couple breast tumor size and metastatic spread. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 6740-6745.	7.1	331
18	Are triple-negative tumours and basal-like breast cancer synonymous? Authors' response. <i>Breast Cancer Research</i> , 2007, 9, .	5.0	9

#	ARTICLE	IF	CITATIONS
19	Gene expression profiling and histopathological characterization of triple-negative/basal-like breast carcinomas. <i>Breast Cancer Research</i> , 2007, 9, R65.	5.0	509
20	Predicting a local recurrence after breast-conserving therapy by gene expression profiling. <i>Breast Cancer Research</i> , 2006, 8, R62.	5.0	184
21	Classification of ductal carcinoma in situ by gene expression profiling. <i>Breast Cancer Research</i> , 2006, 8, R61.	5.0	142
22	Gene Expression Profiles of Primary Breast Carcinomas from Patients at High Risk for Local Recurrence after Breast-Conserving Therapy. <i>Clinical Cancer Research</i> , 2006, 12, 5705-5712.	7.0	56