JÃ¼rgen Veeck

List of Publications by Year
in descending order

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39 papers	3,157 citations	201658 27 h-index	265191 42 g-index
42 all docs	42 docs citations	42 times ranked	5638 citing authors

#	Article	IF	Citations
1	The trans-DATA study: aims and design of a translational breast cancer prognostic marker identification study. Diagnostic and Prognostic Research, 2019, 3, 20.	1.8	1
2	Analysis of DNA methylation in cancer: location revisited. Nature Reviews Clinical Oncology, 2018, 15, 459-466.	27.6	486
3	Promoter methylation of DNA damage repair (DDR) genes in human tumor entities: RBBP8/CtIP is almost exclusively methylated in bladder cancer. Clinical Epigenetics, 2018, 10, 15.	4.1	32
4	ITIH5 mediates epigenetic reprogramming of breast cancer cells. Molecular Cancer, 2017, 16, 44.	19.2	29
5	Differential diagnosis of bladder versus colorectal adenocarcinoma: keratin 7 and GATA3 positivity in nuclear ÄŸ-catenin-negative glandular tumours defines adenocarcinoma of the bladder. Journal of Clinical Pathology, 2016, 69, 307-312.	2.0	19
6	Fibroblast growth factor receptor (FGFR) gene amplifications are rare events in bladder cancer. Histopathology, 2015, 66, 639-649.	2.9	38
7	Epigenetic Biomarker to Support Classification into Pluripotent and Non-Pluripotent Cells. Scientific Reports, 2015, 5, 8973.	3.3	49
8	Formalin-fixed, paraffin-embedded (FFPE) tissue epigenomics using Infinium HumanMethylation450 BeadChip assays. Laboratory Investigation, 2015, 95, 833-842.	3.7	40
9	Low expression of ITIH5 in adenocarcinoma of the lung is associated with unfavorable patients' outcome. Epigenetics, 2015, 10, 903-912.	2.7	30
10	Resistance to sunitinib in renal cell carcinoma: From molecular mechanisms to predictive markers and future perspectives. Biochimica Et Biophysica Acta: Reviews on Cancer, 2015, 1855, 1-16.	7.4	73
11	Towards sustainable data management in professional biobanking. Studies in Health Technology and Informatics, 2015, 212, 94-102.	0.3	2
12	A randomised controlled phase II trial of pre-operative celecoxib treatment reveals anti-tumour transcriptional response in primary breast cancer. Breast Cancer Research, 2013, 15, R29.	5.0	55
13	Promoter hypermethylation of the tumor-suppressor genes ITIH5, DKK3, and RASSF1A as novel biomarkers for blood-based breast cancer screening. Breast Cancer Research, 2013, 15, R4.	5.0	113
14	If there is no overall survival benefit in metastatic breast cancer: Does it imply lack of efficacy? Taxanes as an example. Cancer Treatment Reviews, 2013, 39, 189-198.	7.7	9
15	Post-mortem analysis of bone marrow osteoclasts using tartrate-resistant acid phosphatase staining: does histochemistry work and correlate with time since death?. Journal of Clinical Pathology, 2012, 65, 1013-1018.	2.0	7
16	Epigenetic Changes in Basal Cell Carcinoma Affect SHH and WNT Signaling Components. PLoS ONE, 2012, 7, e51710.	2.5	38
17	Targeting the Wnt pathway in cancer: The emerging role of Dickkopf-3. Biochimica Et Biophysica Acta: Reviews on Cancer, 2012, 1825, 18-28.	7.4	132
18	Taxane resistance in breast cancer: A closed HER2 circuit?. Biochimica Et Biophysica Acta: Reviews on Cancer, 2012, 1825, 197-206.	7.4	22

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19	Genetics and epigenetics of cutaneous malignant melanoma: A concert out of tune. Biochimica Et Biophysica Acta: Reviews on Cancer, 2012, 1826, 89-102.	7.4	46
20	Characteristics of triple-negative breast cancer. Journal of Cancer Research and Clinical Oncology, 2011, 137, 183-192.	2.5	225
21	Paradox of sonic hedgehog (SHH) transcriptional regulation: Alternative transcription initiation overrides the effect of downstream promoter DNA methylation. Epigenetics, 2011, 6, 465-477.	2.7	10
22	Breast Cancer Epigenetics: From DNA Methylation to microRNAs. Journal of Mammary Gland Biology and Neoplasia, 2010, 15, 5-17.	2.7	167
23	BRCA1 CpG Island Hypermethylation Predicts Sensitivity to Poly(Adenosine Diphosphate)- Ribose Polymerase Inhibitors. Journal of Clinical Oncology, 2010, 28, e563-e564.	1.6	152
24	RNA Expression Analysis on Formalin-Fixed Paraffin-Embedded Tissues in TMA Format by RNA In Situ Hybridization. Methods in Molecular Biology, 2010, 664, 135-150.	0.9	8
25	Prognostic relevance of Wnt-inhibitory factor-1 (WIF1) and Dickkopf-3 (DKK3) promoter methylation in human breast cancer. BMC Cancer, 2009, 9, 217.	2.6	81
26	Frequent loss of endothelin-3 (EDN3) expression due to epigenetic inactivation in human breast cancer. Breast Cancer Research, 2009, 11, R34.	5.0	46
27	The extracellular matrix protein ITIH5 is a novel prognostic marker in invasive node-negative breast cancer and its aberrant expression is caused by promoter hypermethylation. Oncogene, 2008, 27, 865-876.	5.9	75
28	Promoter methylation-associated loss of ID4expression is a marker of tumour recurrence in human breast cancer. BMC Cancer, 2008, 8, 154.	2.6	72
29	Frequent expression loss of Inter-alpha-trypsin inhibitor heavy chain (ITIH) genes in multiple human solid tumors: A systematic expression analysis. BMC Cancer, 2008, 8, 25.	2.6	179
30	Tight correlation between expression of the Forkhead transcription factor FOXM1 and HER2 in human breast cancer. BMC Cancer, 2008, 8, 42.	2.6	139
31	Promoter hypermethylation of the SFRP2 gene is a high-frequent alteration and tumor-specific epigenetic marker in human breast cancer. Molecular Cancer, 2008, 7, 83.	19.2	77
32	Wnt signalling in human breast cancer: expression of the putative Wnt inhibitor Dickkopf-3 (DKK3) is frequently suppressed by promoter hypermethylation in mammary tumours. Breast Cancer Research, 2008, 10, R82.	5.0	86
33	The ubiquitin-like molecule interferon-stimulated gene 15 (ISG15) is a potential prognostic marker in human breast cancer. Breast Cancer Research, 2008, 10, R58.	5.0	95
34	Epigenetic inactivation of the secreted frizzled-related protein-5 (SFRP5) gene in human breast cancer is associated with unfavorable prognosis. Carcinogenesis, 2008, 29, 991-998.	2.8	89
35	Frequent loss of SFRP1 expression in multiple human solid tumours: association with aberrant promoter methylation in renal cell carcinoma. Oncogene, 2007, 26, 5680-5691.	5.9	127
36	Aberrant methylation of the Wnt antagonist SFRP1 in breast cancer is associated with unfavourable prognosis. Oncogene, 2006, 25, 3479-3488.	5.9	234

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37	Virologic therapy response significantly correlates with the number of active drugs as evaluated using a LiPA HIV-1 resistance scoring system. Journal of Clinical Virology, 2004, 31, 7-15.	3.1	1
38	Biosynthesis of phytochelatins in the fission yeast. Phytochelatin synthesis: a second role for the glutathione synthetase gene ofSchizosaccharomyces pombe. Yeast, 1999, 15, 385-396.	1.7	31
39	Minimizing the Exposure to UV Light When Extracting DNA from Agarose Gels. BioTechniques, 1998, 25, 586.	1.8	1