

JÃ¼rgen Veeck

List of Publications by Year in descending order

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Version: 2024-02-01

39
papers

3,157
citations

201674

27
h-index

265206

42
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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. <i>Diagnostic and Prognostic Research</i> , 2019, 3, 20.	1.8	1
2	Analysis of DNA methylation in cancer: location revisited. <i>Nature Reviews Clinical Oncology</i> , 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. <i>Clinical Epigenetics</i> , 2018, 10, 15.	4.1	32
4	ITIH5 mediates epigenetic reprogramming of breast cancer cells. <i>Molecular Cancer</i> , 2017, 16, 44.	19.2	29
5	Differential diagnosis of bladder versus colorectal adenocarcinoma: keratin 7 and GATA3 positivity in nuclear β -catenin-negative glandular tumours defines adenocarcinoma of the bladder. <i>Journal of Clinical Pathology</i> , 2016, 69, 307-312.	2.0	19
6	Fibroblast growth factor receptor (FGFR) gene amplifications are rare events in bladder cancer. <i>Histopathology</i> , 2015, 66, 639-649.	2.9	38
7	Epigenetic Biomarker to Support Classification into Pluripotent and Non-Pluripotent Cells. <i>Scientific Reports</i> , 2015, 5, 8973.	3.3	49
8	Formalin-fixed, paraffin-embedded (FFPE) tissue epigenomics using Infinium HumanMethylation450 BeadChip assays. <i>Laboratory Investigation</i> , 2015, 95, 833-842.	3.7	40
9	Low expression of ITIH5 in adenocarcinoma of the lung is associated with unfavorable patients' outcome. <i>Epigenetics</i> , 2015, 10, 903-912.	2.7	30
10	Resistance to sunitinib in renal cell carcinoma: From molecular mechanisms to predictive markers and future perspectives. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2015, 1855, 1-16.	7.4	73
11	Towards sustainable data management in professional biobanking. <i>Studies in Health Technology and Informatics</i> , 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. <i>Breast Cancer Research</i> , 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. <i>Breast Cancer Research</i> , 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. <i>Cancer Treatment Reviews</i> , 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?. <i>Journal of Clinical Pathology</i> , 2012, 65, 1013-1018.	2.0	7
16	Epigenetic Changes in Basal Cell Carcinoma Affect SHH and WNT Signaling Components. <i>PLoS ONE</i> , 2012, 7, e51710.	2.5	38
17	Targeting the Wnt pathway in cancer: The emerging role of Dickkopf-3. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2012, 1825, 18-28.	7.4	132
18	Taxane resistance in breast cancer: A closed HER2 circuit?. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2012, 1825, 197-206.	7.4	22

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19	Genetics and epigenetics of cutaneous malignant melanoma: A concert out of tune. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2012, 1826, 89-102.	7.4	46
20	Characteristics of triple-negative breast cancer. <i>Journal of Cancer Research and Clinical Oncology</i> , 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. <i>Epigenetics</i> , 2011, 6, 465-477.	2.7	10
22	Breast Cancer Epigenetics: From DNA Methylation to microRNAs. <i>Journal of Mammary Gland Biology and Neoplasia</i> , 2010, 15, 5-17.	2.7	167
23	<i>BRCA1</i> CpG Island Hypermethylation Predicts Sensitivity to Poly(Adenosine Diphosphate)-Ribose Polymerase Inhibitors. <i>Journal of Clinical Oncology</i> , 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. <i>Methods in Molecular Biology</i> , 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. <i>BMC Cancer</i> , 2009, 9, 217.	2.6	81
26	Frequent loss of endothelin-3 (EDN3) expression due to epigenetic inactivation in human breast cancer. <i>Breast Cancer Research</i> , 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. <i>Oncogene</i> , 2008, 27, 865-876.	5.9	75
28	Promoter methylation-associated loss of ID4 expression is a marker of tumour recurrence in human breast cancer. <i>BMC Cancer</i> , 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. <i>BMC Cancer</i> , 2008, 8, 25.	2.6	179
30	Tight correlation between expression of the Forkhead transcription factor FOXM1 and HER2 in human breast cancer. <i>BMC Cancer</i> , 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. <i>Molecular Cancer</i> , 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. <i>Breast Cancer Research</i> , 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. <i>Breast Cancer Research</i> , 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. <i>Carcinogenesis</i> , 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. <i>Oncogene</i> , 2007, 26, 5680-5691.	5.9	127
36	Aberrant methylation of the Wnt antagonist SFRP1 in breast cancer is associated with unfavourable prognosis. <i>Oncogene</i> , 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. <i>Journal of Clinical Virology</i> , 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 of <i>Schizosaccharomyces pombe</i> . <i>Yeast</i> , 1999, 15, 385-396.	1.7	31
39	Minimizing the Exposure to UV Light When Extracting DNA from Agarose Gels. <i>BioTechniques</i> , 1998, 25, 586.	1.8	1