

# JÃ¼rgen Veeck

## List of Publications by Year in descending order

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39  
papers

3,157  
citations

201674

27  
h-index

265206

42  
g-index

42  
all docs

42  
docs citations

42  
times ranked

5638  
citing authors

#	ARTICLE	IF	CITATIONS
1	Analysis of DNA methylation in cancer: location revisited. <i>Nature Reviews Clinical Oncology</i> , 2018, 15, 459-466.	27.6	486
2	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
3	Characteristics of triple-negative breast cancer. <i>Journal of Cancer Research and Clinical Oncology</i> , 2011, 137, 183-192.	2.5	225
4	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
5	Breast Cancer Epigenetics: From DNA Methylation to microRNAs. <i>Journal of Mammary Gland Biology and Neoplasia</i> , 2010, 15, 5-17.	2.7	167
6	<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
7	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
8	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
9	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
10	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
11	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
12	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
13	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
14	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
15	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
16	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
17	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
18	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

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19	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
20	Epigenetic Biomarker to Support Classification into Pluripotent and Non-Pluripotent Cells. <i>Scientific Reports</i> , 2015, 5, 8973.	3.3	49
21	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
22	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
23	Formalin-fixed, paraffin-embedded (FFPE) tissue epigenomics using Infinium HumanMethylation450 BeadChip assays. <i>Laboratory Investigation</i> , 2015, 95, 833-842.	3.7	40
24	Epigenetic Changes in Basal Cell Carcinoma Affect SHH and WNT Signaling Components. <i>PLoS ONE</i> , 2012, 7, e51710.	2.5	38
25	Fibroblast growth factor receptor (FGFR) gene amplifications are rare events in bladder cancer. <i>Histopathology</i> , 2015, 66, 639-649.	2.9	38
26	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
27	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
28	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
29	ITIH5 mediates epigenetic reprogramming of breast cancer cells. <i>Molecular Cancer</i> , 2017, 16, 44.	19.2	29
30	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
31	Differential diagnosis of bladder versus colorectal adenocarcinoma: keratin 7 and GATA3 positivity in nuclear $\beta$ -catenin-negative glandular tumours defines adenocarcinoma of the bladder. <i>Journal of Clinical Pathology</i> , 2016, 69, 307-312.	2.0	19
32	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
33	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
34	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
35	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
36	Towards sustainable data management in professional biobanking. <i>Studies in Health Technology and Informatics</i> , 2015, 212, 94-102.	0.3	2

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37	Minimizing the Exposure to UV Light When Extracting DNA from Agarose Gels. <i>BioTechniques</i> , 1998, 25, 586.	1.8	1
38	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
39	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