

# Jrgen Veeck

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

39  
papers

2,408  
citations

26  
h-index

42  
g-index

42  
ext. papers

2,819  
ext. citations

8.1  
avg, IF

4.55  
L-index

#	Paper	IF	Citations
39	The trans-DATA study: aims and design of a translational breast cancer prognostic marker identification study. <i>Diagnostic and Prognostic Research</i> , <b>2019</b> , 3, 20	5.5	1
38	Analysis of DNA methylation in cancer: location revisited. <i>Nature Reviews Clinical Oncology</i> , <b>2018</b> , 15, 459-466	19.4	246
37	Promoter methylation of DNA damage repair (DDR) genes in human tumor entities: / is almost exclusively methylated in bladder cancer. <i>Clinical Epigenetics</i> , <b>2018</b> , 10, 15	7.7	20
36	ITIH5 mediates epigenetic reprogramming of breast cancer cells. <i>Molecular Cancer</i> , <b>2017</b> , 16, 44	42.1	15
35	Differential diagnosis of bladder versus colorectal adenocarcinoma: keratin 7 and GATA3 positivity in nuclear E-catenin-negative glandular tumours defines adenocarcinoma of the bladder. <i>Journal of Clinical Pathology</i> , <b>2016</b> , 69, 307-12	3.9	13
34	Epigenetic biomarker to support classification into pluripotent and non-pluripotent cells. <i>Scientific Reports</i> , <b>2015</b> , 5, 8973	4.9	28
33	Formalin-fixed, paraffin-embedded (FFPE) tissue epigenomics using Infinium HumanMethylation450 BeadChip assays. <i>Laboratory Investigation</i> , <b>2015</b> , 95, 833-42	5.9	33
32	Low expression of ITIH5 in adenocarcinoma of the lung is associated with unfavorable patientsT outcome. <i>Epigenetics</i> , <b>2015</b> , 10, 903-12	5.7	23
31	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> , <b>2015</b> , 1855, 1-16	11.2	44
30	Fibroblast growth factor receptor (FGFR) gene amplifications are rare events in bladder cancer. <i>Histopathology</i> , <b>2015</b> , 66, 639-49	7.3	34
29	Towards sustainable data management in professional biobanking. <i>Studies in Health Technology and Informatics</i> , <b>2015</b> , 212, 94-102	0.5	2
28	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> , <b>2013</b> , 15, R29	8.3	48
27	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> , <b>2013</b> , 15, R4	8.3	92
26	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> , <b>2013</b> , 39, 189-98	14.4	9
25	Targeting the Wnt pathway in cancer: the emerging role of Dickkopf-3. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , <b>2012</b> , 1825, 18-28	11.2	97
24	Taxane resistance in breast cancer: a closed HER2 circuit?. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , <b>2012</b> , 1825, 197-206	11.2	20
23	Genetics and epigenetics of cutaneous malignant melanoma: a concert out of tune. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , <b>2012</b> , 1826, 89-102	11.2	39

22	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> , <b>2012</b> , 65, 1013-8	3.9	4
21	Epigenetic changes in Basal Cell Carcinoma affect SHH and WNT signaling components. <i>PLoS ONE</i> , <b>2012</b> , 7, e51710	3.7	30
20	Characteristics of triple-negative breast cancer. <i>Journal of Cancer Research and Clinical Oncology</i> , <b>2011</b> , 137, 183-92	4.9	172
19	Paradox of sonic hedgehog (SHH) transcriptional regulation: Alternative transcription initiation overrides the effect of downstream promoter DNA methylation. <i>Epigenetics</i> , <b>2011</b> , 6, 465-77	5.7	8
18	BRCA1 CpG island hypermethylation predicts sensitivity to poly(adenosine diphosphate)-ribose polymerase inhibitors. <i>Journal of Clinical Oncology</i> , <b>2010</b> , 28, e563-4; author reply e565-6	2.2	129
17	Breast cancer epigenetics: from DNA methylation to microRNAs. <i>Journal of Mammary Gland Biology and Neoplasia</i> , <b>2010</b> , 15, 5-17	2.4	144
16	RNA expression analysis on formalin-fixed paraffin-embedded tissues in TMA format by RNA in situ hybridization. <i>Methods in Molecular Biology</i> , <b>2010</b> , 664, 135-50	1.4	8
15	Prognostic relevance of Wnt-inhibitory factor-1 (WIF1) and Dickkopf-3 (DKK3) promoter methylation in human breast cancer. <i>BMC Cancer</i> , <b>2009</b> , 9, 217	4.8	69
14	Frequent loss of endothelin-3 (EDN3) expression due to epigenetic inactivation in human breast cancer. <i>Breast Cancer Research</i> , <b>2009</b> , 11, R34	8.3	38
13	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> , <b>2008</b> , 27, 865-76 <sup>9.2</sup>	9.2	60
12	Promoter methylation-associated loss of ID4 expression is a marker of tumour recurrence in human breast cancer. <i>BMC Cancer</i> , <b>2008</b> , 8, 154	4.8	67
11	Frequent expression loss of Inter-alpha-trypsin inhibitor heavy chain (ITI) genes in multiple human solid tumors: a systematic expression analysis. <i>BMC Cancer</i> , <b>2008</b> , 8, 25	4.8	121
10	Tight correlation between expression of the Forkhead transcription factor FOXM1 and HER2 in human breast cancer. <i>BMC Cancer</i> , <b>2008</b> , 8, 42	4.8	114
9	Promoter hypermethylation of the SFRP2 gene is a high-frequent alteration and tumor-specific epigenetic marker in human breast cancer. <i>Molecular Cancer</i> , <b>2008</b> , 7, 83	42.1	63
8	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> , <b>2008</b> , 10, R82	8.3	70
7	The ubiquitin-like molecule interferon-stimulated gene 15 (ISG15) is a potential prognostic marker in human breast cancer. <i>Breast Cancer Research</i> , <b>2008</b> , 10, R58	8.3	80
6	Epigenetic inactivation of the secreted frizzled-related protein-5 (SFRP5) gene in human breast cancer is associated with unfavorable prognosis. <i>Carcinogenesis</i> , <b>2008</b> , 29, 991-8	4.6	80
5	Frequent loss of SFRP1 expression in multiple human solid tumours: association with aberrant promoter methylation in renal cell carcinoma. <i>Oncogene</i> , <b>2007</b> , 26, 5680-91	9.2	118

4	Aberrant methylation of the Wnt antagonist SFRP1 in breast cancer is associated with unfavourable prognosis. <i>Oncogene</i> , <b>2006</b> , 25, 3479-88	9.2	204
3	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> , <b>2004</b> , 31 Suppl 1, S7-15	14.5	1
2	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> , <b>1999</b> , 15, 385-96	3.4	30
1	Minimizing the exposure to UV light when extracting DNA from agarose gels. <i>BioTechniques</i> , <b>1998</b> , 25, 586	2.5	1