

Jrgen Veeck

List of Publications by Citations

Source: <https://exaly.com/author-pdf/3809404/jjrgen-veeck-publications-by-citations.pdf>

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

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	Analysis of DNA methylation in cancer: location revisited. <i>Nature Reviews Clinical Oncology</i> , 2018 , 15, 459-466	19.4	246
38	Aberrant methylation of the Wnt antagonist SFRP1 in breast cancer is associated with unfavourable prognosis. <i>Oncogene</i> , 2006 , 25, 3479-88	9.2	204
37	Characteristics of triple-negative breast cancer. <i>Journal of Cancer Research and Clinical Oncology</i> , 2011 , 137, 183-92	4.9	172
36	Breast cancer epigenetics: from DNA methylation to microRNAs. <i>Journal of Mammary Gland Biology and Neoplasia</i> , 2010 , 15, 5-17	2.4	144
35	BRCA1 CpG island hypermethylation predicts sensitivity to poly(adenosine diphosphate)-ribose polymerase inhibitors. <i>Journal of Clinical Oncology</i> , 2010 , 28, e563-4; author reply e565-6	2.2	129
34	Frequent expression loss of Inter-alpha-trypsin inhibitor heavy chain (ITI _H) genes in multiple human solid tumors: a systematic expression analysis. <i>BMC Cancer</i> , 2008 , 8, 25	4.8	121
33	Frequent loss of SFRP1 expression in multiple human solid tumours: association with aberrant promoter methylation in renal cell carcinoma. <i>Oncogene</i> , 2007 , 26, 5680-91	9.2	118
32	Tight correlation between expression of the Forkhead transcription factor FOXM1 and HER2 in human breast cancer. <i>BMC Cancer</i> , 2008 , 8, 42	4.8	114
31	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	11.2	97
30	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	8.3	92
29	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	8.3	80
28	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, 991-8	4.6	80
27	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	8.3	70
26	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	4.8	69
25	Promoter methylation-associated loss of ID4 expression is a marker of tumour recurrence in human breast cancer. <i>BMC Cancer</i> , 2008 , 8, 154	4.8	67
24	Promoter hypermethylation of the SFRP2 gene is a high-frequency alteration and tumor-specific epigenetic marker in human breast cancer. <i>Molecular Cancer</i> , 2008 , 7, 83	42.1	63
23	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, 865-76	9.2	60

22	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	8.3	48
21	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	11.2	44
20	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	11.2	39
19	Frequent loss of endothelin-3 (EDN3) expression due to epigenetic inactivation in human breast cancer. <i>Breast Cancer Research</i> , 2009 , 11, R34	8.3	38
18	Fibroblast growth factor receptor (FGFR) gene amplifications are rare events in bladder cancer. <i>Histopathology</i> , 2015 , 66, 639-49	7.3	34
17	Formalin-fixed, paraffin-embedded (FFPE) tissue epigenomics using Infinium HumanMethylation450 BeadChip assays. <i>Laboratory Investigation</i> , 2015 , 95, 833-42	5.9	33
16	Epigenetic changes in Basal Cell Carcinoma affect SHH and WNT signaling components. <i>PLoS ONE</i> , 2012 , 7, e51710	3.7	30
15	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, 385-96	3.4	30
14	Epigenetic biomarker to support classification into pluripotent and non-pluripotent cells. <i>Scientific Reports</i> , 2015 , 5, 8973	4.9	28
13	Low expression of ITIH5 in adenocarcinoma of the lung is associated with unfavorable patientsT outcome. <i>Epigenetics</i> , 2015 , 10, 903-12	5.7	23
12	Promoter methylation of DNA damage repair (DDR) genes in human tumor entities: / is almost exclusively methylated in bladder cancer. <i>Clinical Epigenetics</i> , 2018 , 10, 15	7.7	20
11	Taxane resistance in breast cancer: a closed HER2 circuit?. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2012 , 1825, 197-206	11.2	20
10	ITIH5 mediates epigenetic reprogramming of breast cancer cells. <i>Molecular Cancer</i> , 2017 , 16, 44	42.1	15
9	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> , 2016 , 69, 307-12	3.9	13
8	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, 189-98	14.4	9
7	Paradox of sonic hedgehog (SHH) transcriptional regulation: Alternative transcription initiation overrides the effect of downstream promoter DNA methylation. <i>Epigenetics</i> , 2011 , 6, 465-77	5.7	8
6	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, 135-50	1.4	8
5	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, 1013-8	3.9	4

4	Towards sustainable data management in professional biobanking. <i>Studies in Health Technology and Informatics</i> , 2015 , 212, 94-102	0.5	2
3	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	5.5	1
2	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 Suppl 1, S7-15	14.5	1
1	Minimizing the exposure to UV light when extracting DNA from agarose gels. <i>BioTechniques</i> , 1998 , 25, 586	2.5	1