## Tineke E Buffart

List of Publications by Year in descending order

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840776 794594 23 398 11 19 citations h-index g-index papers 23 23 23 860 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Development of a miRNAâ€based classifier for detection of colorectal cancer molecular subtypes. Molecular Oncology, 2022, 16, 2693-2709.	4.6	6
2	Time dependent effect of cold ischemia on the phosphoproteome and protein kinase activity in fresh-frozen colorectal cancer tissue obtained from patients. Clinical Proteomics, 2021, 18, 8.	2.1	2
3	Feasibility of whole-genome sequencing in routine clinical practice Journal of Clinical Oncology, 2021, 39, 3013-3013.	1.6	3
4	Perioperative Systemic Therapy vs Cytoreductive Surgery and Hyperthermic Intraperitoneal Chemotherapy Alone for Resectable Colorectal Peritoneal Metastases. JAMA Surgery, 2021, 156, 710-720.	4.3	34
5	Trends in Use and Perceptions About Triplet Chemotherapy Plus Bevacizumab for Metastatic Colorectal Cancer. JAMA Network Open, 2021, 4, e2124766.	5.9	2
6	Study protocol: Whole genome sequencing Implementation in standard Diagnostics for Every cancer patient (WIDE). BMC Medical Genomics, 2020, 13, 169.	1.5	30
7	Safety and Feasibility of Additional Tumor Debulking to First-Line Palliative Combination Chemotherapy for Patients with Multiorgan Metastatic Colorectal Cancer. Oncologist, 2020, 25, e1195-e1201.	3.7	7
8	Proteomic Analysis of miR-195 and miR-497 Replacement Reveals Potential Candidates that Increase Sensitivity to Oxaliplatin in MSI/P53wt Colorectal Cancer Cells. Cells, 2019, 8, 1111.	4.1	25
9	CD276-positive circulating endothelial cells in advanced colorectal cancer Journal of Clinical Oncology, 2019, 37, 572-572.	1.6	2
10	Comparison of phosphoproteomic profiles in left- and right-sided colorectal cancers Journal of Clinical Oncology, 2019, 37, 582-582.	1.6	0
11	Evaluation of several methodological challenges in circulating miRNA qPCR studies in patients with head and neck cancer. Experimental and Molecular Medicine, 2018, 50, e454-e454.	7.7	59
12	Combination of a six microRNA expression profile with four clinicopathological factors for response prediction of systemic treatment in patients with advanced colorectal cancer. PLoS ONE, 2018, 13, e0201809.	2.5	20
13	Safety and feasibility of adding tumor debulking to palliative chemotherapy in multi-organ metastatic colorectal cancer: The ORCHESTRA trial Journal of Clinical Oncology, 2017, 35, 3553-3553.	1.6	О
14	The ORCHESTRA trial: A phase III trial of adding tumor debulking to systemic therapy versus systemic therapy alone in multi-organ metastatic colorectal cancer (mCRC) Journal of Clinical Oncology, 2016, 34, TPS788-TPS788.	1.6	2
15	Comparison of deep sequencing miRNA expression analysis in primary colorectal cancer and paired metastases Journal of Clinical Oncology, 2015, 33, e14682-e14682.	1.6	0
16	The ORCHESTRA trial: A phase III trial of adding tumor debulking to systemic therapy versus systemic therapy alone in (mCRC) multi-organ metastatic colorectal cancer Journal of Clinical Oncology, 2015, 33, TPS3631-TPS3631.	1.6	0
17	Losses of Chromosome 5q and 14q Are Associated with Favorable Clinical Outcome of Patients with Gastric Cancer. Oncologist, 2012, 17, 653-662.	3.7	27
18	Gastric cancers of Western European and African patients show different patterns of genomic instability. BMC Medical Genomics, 2011, 4, 7.	1.5	30

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#	Article	IF	CITATION
19	High resolution analysis of DNA copy-number aberrations of chromosomes 8, 13, and 20 in gastric cancers. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2009, 455, 213-223.	2.8	51
20	NMD inhibition fails to identify tumour suppressor genes in microsatellite stable gastric cancer cell lines. BMC Medical Genomics, 2009, 2, 39.	1.5	4
21	Across array comparative genomic hybridization: A strategy to reduce reference channel hybridizations. Genes Chromosomes and Cancer, 2008, 47, 994-1004.	2.8	52
22	DNA copy number profiles of gastric cancer precursor lesions. BMC Genomics, 2007, 8, 345.	2.8	28
23	DNA Quality Assessment for Array CGH by Isothermal Whole Genome Amplification. Analytical Cellular Pathology, 2007, 29, 351-359.	1.4	14