## Jacqueline A James

List of Publications by Year in descending order

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172207 133063 5,848 60 29 59 citations h-index g-index papers 61 61 61 11417 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Swarm learning for decentralized artificial intelligence in cancer histopathology. Nature Medicine, 2022, 28, 1232-1239.	15.2	77
2	A biobank perspective on use of tissue samples donated by trial participants. Lancet Oncology, The, 2022, 23, e205.	5.1	2
3	General Roadmap and Core Steps for the Development of Al Tools in Digital Pathology. Diagnostics, 2022, 12, 1272.	1.3	4
4	Digital pathology and artificial intelligence will be key to supporting clinical and academic cellular pathology through COVID-19 and future crises: the PathLAKE consortium perspective. Journal of Clinical Pathology, 2021, 74, 443-447.	1.0	49
5	Identifying mismatch repairâ€deficient colon cancer: nearâ€perfect concordance between immunohistochemistry and microsatellite instability testing in a large, populationâ€based series. Histopathology, 2021, 78, 401-413.	1.6	55
6	Evolutionary genetic algorithm identifies <i>IL2RB</i> as a potential predictive biomarker for immune-checkpoint therapy in colorectal cancer. NAR Genomics and Bioinformatics, 2021, 3, Iqab016.	1.5	10
7	Colonic epithelial cathelicidin ( <scp>LL</scp> â€37) expression intensity is associated with progression of colorectal cancer and presence of <scp>CD8</scp> <sup>+</sup> T cell infiltrate. Journal of Pathology: Clinical Research, 2021, 7, 495-506.	1.3	8
8	A Means of Assessing Deep Learning-Based Detection of ICOS Protein Expression in Colon Cancer. Cancers, 2021, 13, 3825.	1.7	17
9	The Potential of Digital Image Analysis to Determine Tumor Cell Content in Biobanked Formalin-Fixed, Paraffin-Embedded Tissue Samples. Biopreservation and Biobanking, 2021, 19, 324-331.	0.5	5
10	Orthogonal <i>MET</i> analysis in a populationâ€representative stage llâ€"III colon cancer cohort: prognostic and potential therapeutic implications. Molecular Oncology, 2021, 15, 3317-3328.	2.1	3
11	Metastasis and Immune Evasion from Extracellular cGAMP Hydrolysis. Cancer Discovery, 2021, 11, 1212-1227.	7.7	139
12	PD-L1 Multiplex and Quantitative Image Analysis for Molecular Diagnostics. Cancers, 2021, 13, 29.	1.7	11
13	Fusobacterium nucleatum and oral cancer: a critical review. BMC Cancer, 2021, 21, 1212.	1.1	50
14	Comparison of Molecular Assays for HPV Testing in Oropharyngeal Squamous Cell Carcinomas: A Population-Based Study in Northern Ireland. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 31-38.	1.1	14
15	Alcohol intake, tobacco smoking, and esophageal adenocarcinoma survival: a molecular pathology epidemiology cohort study. Cancer Causes and Control, 2020, 31, 1-11.	0.8	16
16	Immune status is prognostic for poor survival in colorectal cancer patients and is associated with tumour hypoxia. British Journal of Cancer, 2020, 123, 1280-1288.	2.9	45
17	Improving the Diagnostic Accuracy of the PD-L1 Test with Image Analysis and Multiplex Hybridization. Cancers, 2020, 12, 1114.	1.7	34
18	The adaptive immune and immune checkpoint landscape of neoadjuvant treated esophageal adenocarcinoma using digital pathology quantitation. BMC Cancer, 2020, 20, 500.	1.1	20

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19	MAPKAPK2 (MK2) inhibition mediates radiation-induced inflammatory cytokine production and tumor growth in head and neck squamous cell carcinoma. Oncogene, 2019, 38, 7329-7341.	2.6	15
20	Recommendations for determining HPV status in patients with oropharyngeal cancers under TNM8 guidelines: a two-tier approach. British Journal of Cancer, 2019, 120, 827-833.	2.9	51
21	Immune activation by DNA damage predicts response to chemotherapy and survival in oesophageal adenocarcinoma. Gut, 2019, 68, 1918-1927.	6.1	18
22	Quality assurance guidance for scoring and reporting for pathologists and laboratories undertaking clinical trial work. Journal of Pathology: Clinical Research, 2019, 5, 91-99.	1.3	21
23	High PTGS2 expression in postâ€neoadjuvant chemotherapyâ€treated oesophageal adenocarcinoma is associated with improved survival: a populationâ€based cohort study. Histopathology, 2019, 74, 587-596.	1.6	1
24	Critical Appraisal of Programmed Death Ligand 1 Reflex Diagnostic Testing: Current Standards and Future Opportunities. Journal of Thoracic Oncology, 2019, 14, 45-53.	0.5	42
25	Punctate <scp>MLH</scp> 1 mismatch repair immunostaining in colorectal cancer. Histopathology, 2019, 74, 795-797.	1.6	11
26	Practical guide for the comparison of two next-generation sequencing systems for solid tumour analysis in a universal healthcare system. Journal of Clinical Pathology, 2019, 72, 225-231.	1.0	7
27	Validation of the systematic scoring of immunohistochemically stained tumour tissue microarrays using <i>QuPath</i> digital image analysis. Histopathology, 2018, 73, 327-338.	1.6	63
28	Integrated tumor identification and automated scoring minimizes pathologist involvement and provides new insights to key biomarkers in breast cancer. Laboratory Investigation, 2018, 98, 15-26.	1.7	81
29	Automated Tumour Recognition and Digital Pathology Scoring Unravels New Role for PD-L1 in Predicting Good Outcome in ER-/HER2+ Breast Cancer. Journal of Oncology, 2018, 2018, 1-14.	0.6	44
30	î"Np63î³/SRC/Slug Signaling Axis Promotes Epithelial-to-Mesenchymal Transition in Squamous Cancers. Clinical Cancer Research, 2018, 24, 3917-3927.	3.2	19
31	Glucose transporter 1 expression as a marker of prognosis in oesophageal adenocarcinoma. Oncotarget, 2018, 9, 18518-18528.	0.8	13
32	Vitamin D receptor as a marker of prognosis in oesophageal adenocarcinoma: a prospective cohort study. Oncotarget, 2018, 9, 34347-34356.	0.8	7
33	Sex hormone receptor expression and survival in esophageal adenocarcinoma: a prospective cohort study. Oncotarget, 2018, 9, 35300-35312.	0.8	6
34	The Northern Ireland Biobank: A Cancer Focused Repository of Science. Open Journal of Bioresources, 2018, 5, .	1.5	30
35	Tissue-based next generation sequencing: application in a universal healthcare system. British Journal of Cancer, 2017, 116, 553-560.	2.9	38
36	Evaluation of PTGS2 Expression, PIK3CA Mutation, Aspirin Use and Colon Cancer Survival in a Population-Based Cohort Study. Clinical and Translational Gastroenterology, 2017, 8, e91.	1.3	56

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37	Statin use, candidate mevalonate pathway biomarkers, and colon cancer survival in a population-based cohort study. British Journal of Cancer, 2017, 116, 1652-1659.	2.9	37
38	Molecular profiling of signet ring cell colorectal cancer provides a strong rationale for genomic targeted and immune checkpoint inhibitor therapies. British Journal of Cancer, 2017, 117, 203-209.	2.9	38
39	QuPath: Open source software for digital pathology image analysis. Scientific Reports, 2017, 7, 16878.	1.6	3,854
40	RNAscope <i>in situ</i> hybridization confirms mRNA integrity in formalin-fixed, paraffin-embedded cancer tissue samples. Oncotarget, 2017, 8, 93392-93403.	0.8	41
41	PD-L1 expression and response to neo-adjuvant chemotherapy in esophageal adenocarcinoma Journal of Clinical Oncology, 2017, 35, 4023-4023.	0.8	1
42	p16 as a prognostic indicator in ovarian/tubal highâ€grade serous carcinoma. Histopathology, 2016, 68, 615-618.	1.6	8
43	Quantification of HER2 heterogeneity in breast cancer–implications for identification of sub-dominant clones for personalised treatment. Scientific Reports, 2016, 6, 23383.	1.6	38
44	Building a †Repository of Science': The importance ofÂintegrating biobanks within molecular pathology programmes. European Journal of Cancer, 2016, 67, 191-199.	1.3	31
45	Delivering a researchâ€enabled multistakeholder partnership for enhanced patient care at a population level: The Northern Ireland Comprehensive Cancer Program. Cancer, 2016, 122, 664-673.	2.0	5
46	HPV-Related Oropharynx Cancer in the United Kingdom: An Evolution in the Understanding of Disease Etiology. Cancer Research, 2016, 76, 6598-6606.	0.4	128
47	PTEN mRNA detection by chromogenic, RNA in situ technologies: a reliable alternative to PTEN immunohistochemistry. Human Pathology, 2016, 47, 95-103.	1.1	17
48	Transcriptional upregulation of c-MET is associated with invasion and tumor budding in colorectal cancer. Oncotarget, 2016, 7, 78932-78945.	0.8	36
49	Automated tumor analysis for molecular profiling in lung cancer. Oncotarget, 2015, 6, 27938-27952.	0.8	43
50	PICan: An integromics framework for dynamic cancer biomarker discovery. Molecular Oncology, 2015, 9, 1234-1240.	2.1	15
51	Analysis of wntless (WLS) expression in gastric, ovarian, and breast cancers reveals a strong association with HER2 overexpression. Modern Pathology, 2015, 28, 428-436.	2.9	27
52	Comprehensive molecular pathology analysis of small bowel adenocarcinoma reveals novel targets with potential for clinical utility. Oncotarget, 2015, 6, 20863-20874.	0.8	41
53	Molecular classification of non-invasive breast lesions for personalised therapy and chemoprevention. Oncotarget, 2015, 6, 43244-43254.	0.8	8
54	Identification and Validation of an Anthracycline/Cyclophosphamide–Based Chemotherapy Response Assay in Breast Cancer. Journal of the National Cancer Institute, 2014, 106, djt335.	3.0	91

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55	The prognostic significance of the aberrant extremes of p53 immunophenotypes in breast cancer. Histopathology, 2014, 65, 340-352.	1.6	59
56	Digital pathology and image analysis in tissue biomarker research. Methods, 2014, 70, 59-73.	1.9	162
57	Molecular pathology – The value of an integrative approach. Molecular Oncology, 2014, 8, 1163-1168.	2.1	32
58	Association of a DNA damage response deficiency (DDRD) assay and prognosis in early-stage esophageal adenocarcinoma Journal of Clinical Oncology, 2014, 32, 4015-4015.	0.8	3
59	POU2F1 activity regulates HOXD10 and HOXD11 promoting a proliferative and invasive phenotype in Head and Neck cancer. Oncotarget, 2014, 5, 8803-8815.	0.8	43
60	Ultra-Fast Processing of Gigapixel Tissue MicroArray Images Using High Performance Computing. Analytical Cellular Pathology, 2010, 33, 271-285.	0.7	4