

# Katie L Meehan

## 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.

27  
papers

4,048  
citations

12  
h-index

29  
g-index

29  
ext. papers

5,893  
ext. citations

5.3  
avg, IF

4.06  
L-index

#	Paper	IF	Citations
27	Dynamic Landscape of Extracellular Vesicle-Associated Proteins Is Related to Treatment Response of Patients with Metastatic Breast Cancer. <i>Membranes</i> , <b>2021</b> , 11,	3.8	1
26	Restoration of the Oral Microbiota After Surgery for Head and Neck Squamous Cell Carcinoma Is Associated With Patient Outcomes. <i>Frontiers in Oncology</i> , <b>2021</b> , 11, 737843	5.3	0
25	Low-coverage whole-genome sequencing of extracellular vesicle-associated DNA in patients with metastatic cancer. <i>Scientific Reports</i> , <b>2021</b> , 11, 4016	4.9	3
24	Proteomic Analysis of Circulating Extracellular Vesicles Identifies Potential Biomarkers for Lymph Node Metastasis in Oral Tongue Squamous Cell Carcinoma. <i>Cells</i> , <b>2021</b> , 10,	7.9	5
23	Profiling of extracellular vesicles in oral cancer, from transcriptomics to proteomics. <i>Seminars in Cancer Biology</i> , <b>2021</b> , 74, 3-23	12.7	6
22	Extracellular Vesicles in Head and Neck Cancer: A Potential New Trend in Diagnosis, Prognosis, and Treatment. <i>International Journal of Molecular Sciences</i> , <b>2020</b> , 21,	6.3	8
21	Characterization of the immune profile of oral tongue squamous cell carcinomas with advancing disease. <i>Cancer Medicine</i> , <b>2020</b> , 9, 4791-4807	4.8	12
20	A comparative study of extracellular vesicle-associated and cell-free DNA and RNA for HPV detection in oropharyngeal squamous cell carcinoma. <i>Scientific Reports</i> , <b>2020</b> , 10, 6083	4.9	12
19	Detection of splicing variants in plasma-derived cell-free nucleic acids and extracellular vesicles of melanoma patients failing targeted therapy therapies. <i>Oncotarget</i> , <b>2020</b> , 11, 4016-4027	3.3	1
18	Comparison of Circulating Tumour DNA and Extracellular Vesicle DNA by Low-Pass Whole-Genome Sequencing Reveals Molecular Drivers of Disease in a Breast Cancer Patient. <i>Biomedicines</i> , <b>2020</b> , 9,	4.8	5
17	The Intersection between Oral Microbiota, Host Gene Methylation and Patient Outcomes in Head and Neck Squamous Cell Carcinoma. <i>Cancers</i> , <b>2020</b> , 12,	6.6	11
16	Laboratory validation studies in Ki-67 digital image analysis of breast carcinoma: a pathway to routine quality assurance. <i>Pathology</i> , <b>2019</b> , 51, 246-252	1.6	5
15	The influence of hydration on the architectural rearrangement of normal and neoplastic human breast tissues. <i>Heliyon</i> , <b>2019</b> , 5, e01219	3.6	3
14	Tumor-Derived Extracellular Vesicles as a Novel Source of Protein Biomarkers for Cancer Diagnosis and Monitoring. <i>Proteomics</i> , <b>2019</b> , 19, e1800155	4.8	29
13	Interobserver variation in the diagnosis of fibroepithelial lesions of the breast: a multicentre audit by digital pathology. <i>Journal of Clinical Pathology</i> , <b>2018</b> , 71, 672-679	3.9	14
12	Genetic characterisation of molecular targets in carcinoma of unknown primary. <i>Journal of Translational Medicine</i> , <b>2018</b> , 16, 185	8.5	13
11	Automated enumeration of lymphoid and plasma cells in bone marrow to establish normal reference ranges. <i>Journal of Clinical Pathology</i> , <b>2018</b> , 71, 916-925	3.9	4

10	Minimal information for studies of extracellular vesicles 2018 (MISEV2018): a position statement of the International Society for Extracellular Vesicles and update of the MISEV2014 guidelines. <i>Journal of Extracellular Vesicles</i> , <b>2018</b> , 7, 1535750	16.4	3642
9	HER2 mRNA transcript quantitation in breast cancer. <i>Clinical and Translational Oncology</i> , <b>2017</b> , 19, 606-615	3.5	4
8	Megakaryocytic hyperplasia in myeloproliferative neoplasms is driven by disordered proliferative, apoptotic and epigenetic mechanisms. <i>Journal of Clinical Pathology</i> , <b>2016</b> , 69, 155-63	3.9	17
7	The contribution of tumour-derived exosomes to the hallmarks of cancer. <i>Critical Reviews in Clinical Laboratory Sciences</i> , <b>2016</b> , 53, 121-31	9.4	74
6	Mutational Analysis of BRAF Inhibitor-Associated Squamoproliferative Lesions. <i>Journal of Molecular Diagnostics</i> , <b>2015</b> , 17, 644-51	5.1	7
5	Decidual-secreted factors alter invasive trophoblast membrane and secreted proteins implying a role for decidual cell regulation of placentation. <i>PLoS ONE</i> , <b>2012</b> , 7, e31418	3.7	35
4	Combination of hydrogel nanoparticles and proteomics to reveal secreted proteins associated with decidualization of human uterine stromal cells. <i>Proteome Science</i> , <b>2011</b> , 9, 50	2.6	5
3	Proteomics and the search for biomarkers of female reproductive diseases. <i>Reproduction</i> , <b>2010</b> , 140, 505-19	3.8	17
2	An optimized procedure for the capture, fractionation and proteomic analysis of proteins using hydrogel nanoparticles. <i>Proteomics</i> , <b>2010</b> , 10, 332-6	4.8	12
1	Proteomic analysis of normal and malignant prostate tissue to identify novel proteins lost in cancer. <i>Prostate</i> , <b>2002</b> , 50, 54-63	4.2	103