Amy Ellen McCart Reed

List of Publications by Year
in descending order

61 papers	3,094 citations	257357 24 h-index	175177 52 g-index
68 all docs	68 docs citations	68 times ranked	6966 citing authors

#	Article	IF	Citations
1	Urothelial Carcinoma and Prostate-specific Membrane Antigen: Cellular, Imaging, and Prognostic Implications. European Urology Focus, 2022, 8, 1256-1269.	1.6	4
2	Blood-Derived Extracellular Vesicle-Associated miR-3182 Detects Non-Small Cell Lung Cancer Patients. Cancers, 2022, 14, 257.	1.7	11
3	Epigenome erosion and SOX10 drive neural crest phenotypic mimicry in triple-negative breast cancer. Npj Breast Cancer, 2022, 8, 57.	2.3	11
4	Invasive lobular carcinoma of the breast: the increasing importance of this special subtype. Breast Cancer Research, 2021, 23, 6.	2.2	64
5	An Update on the Molecular Pathology of Metaplastic Breast Cancer. Breast Cancer: Targets and Therapy, 2021, Volume 13, 161-170.	1.0	7
6	The Genomic Landscape of Lobular Breast Cancer. Cancers, 2021, 13, 1950.	1.7	13
7	Tumor Signature Analysis Implicates Hereditary Cancer Genes in Endometrial Cancer Development. Cancers, 2021, 13, 1762.	1.7	5
8	Characterization of Immune Cell Subsets of Tumor Infiltrating Lymphocytes in Brain Metastases. Biology, 2021, 10, 425.	1.3	6
9	Integrin alpha-2 and beta-1 expression increases through multiple generations of the EDW01 patient-derived xenograft model of breast cancer—insight into their role in epithelial mesenchymal transition in vivo gained from an in vitro model system. Breast Cancer Research, 2020, 22, 136.	2.2	16
10	Clinicopathologic significance of nuclear HER4 and phospho-YAP(S^{127}) in human breast cancers and matching brain metastases. Therapeutic Advances in Medical Oncology, 2020, 12, 175883592094625.	1.4	11
11	Phenotypic drift in metastatic progression of breast cancer: A case report with histologically heterogeneous lesions that are clonally related. Clinical Case Reports (discontinued), 2020, 8, 2725-2731.	0.2	1
12	Metaplastic breast cancers frequently express immune checkpoint markers FOXP3 and PD-L1. British Journal of Cancer, 2020, 123, 1665-1672.	2.9	26
13	Association of Sperm-Associated Antigen 5 and Treatment Response in Patients With Estrogen Receptor–Positive Breast Cancer. JAMA Network Open, 2020, 3, e209486.	2.8	2
14	Using whole-genome sequencing data to derive the homologous recombination deficiency scores. Npj Breast Cancer, 2020, 6, 33.	2.3	19
15	Digital spatial profiling application in breast cancer: a user's perspective. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2020, 477, 885-890.	1.4	16
16	Non-coding RNAs underlie genetic predisposition to breast cancer. Genome Biology, 2020, 21, 7.	3.8	21
17	Morphologic and Genomic Heterogeneity in the Evolution and Progression of Breast Cancer. Cancers, 2020, 12, 848.	1.7	14
18	Tradeoff between metabolic i-proteasome addiction and immune evasion in triple-negative breast cancer. Life Science Alliance, 2020, 3, e201900562.	1.3	11

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19	A Molecular and Morphological Deep-Dive Into Metaplastic Breast Cancers. Cancer Informatics, 2019, 18, 117693511985015.	0.9	1
20	LobSig is a multigene predictor of outcome in invasive lobular carcinoma. Npj Breast Cancer, 2019, 5, 18.	2.3	28
21	Breast Cancer Heterogeneity in Primary and Metastatic Disease. Advances in Experimental Medicine and Biology, 2019, 1152, 75-104.	0.8	27
22	Evaluation of novel immunohistochemical biomarkers for invasive lobular carcinoma. Pathology, 2019, 51, S77.	0.3	0
23	Whole-genome sequencing reveals clinically relevant insights into the aetiology of familial breast cancers. Annals of Oncology, 2019, 30, 1071-1079.	0.6	64
24	Diff-Quik Cytology Smears from Endobronchial Ultrasound Transbronchial Needle Aspiration Lymph Node Specimens as a Source of DNA for Next-Generation Sequencing Instead of Cell Blocks. Respiration, 2019, 97, 525-539.	1.2	25
25	Recent advances in breast cancer research impacting clinical diagnostic practice. Journal of Pathology, 2019, 247, 552-562.	2.1	24
26	Phenotypic and molecular dissection of metaplastic breast cancer and the prognostic implications. Journal of Pathology, 2019, 247, 214-227.	2.1	73
27	Breast cancer metastasis to gynaecological organs: a clinico–pathological and molecular profiling study. Journal of Pathology: Clinical Research, 2019, 5, 25-39.	1.3	31
28	Mixed ductal–lobular carcinomas: evidence for progression from ductal to lobular morphology. Journal of Pathology, 2018, 244, 460-468.	2.1	31
29	Multidimensional phenotyping of breast cancer cell lines to guide preclinical research. Breast Cancer Research and Treatment, 2018, 167, 289-301.	1.1	27
30	Expression of ARGLU and FBXL3 is highly prognostic for invasive lobular carcinoma. Pathology, 2018, 50, S68.	0.3	0
31	Breast Cancer—Pathology and Genetics. , 2018, , .		0
32	The Brisbane Breast Bank. Open Journal of Bioresources, 2018, 5, .	1.5	13
33	Immunohistochemical evaluation of FOXA1 expression and its correlation with breast cancer-specific survival: queensland follow up cohort. Pathology, 2017, 49, S70.	0.3	0
34	Association analysis identifies 65 new breast cancer risk loci. Nature, 2017, 551, 92-94.	13.7	1,099
35	Next-Generation Sequencing of Endobronchial Ultrasound Transbronchial Needle Aspiration Specimens in Lung Cancer. American Journal of Respiratory and Critical Care Medicine, 2017, 196, 388-391.	2.5	14
36	Molecular signatures in breast cancer. Methods, 2017, 131, 135-146.	1.9	47

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37	Breast Cancer Brain Metastases: Clonal Evolution in Clinical Context. International Journal of Molecular Sciences, 2017, 18, 152.	1.8	20
38	An epithelial to mesenchymal transition programme does not usually drive the phenotype of invasive lobular carcinomas. Journal of Pathology, 2016, 238, 489-494.	2.1	32
39	Long-range regulators of the lncRNA<i>HOTAIR</i>enhance its prognostic potential in breast cancer. Human Molecular Genetics, 2016, 25, 3269-3283.	1.4	58
40	â€˜Omics Approaches in Breast Cancer Research and Clinical Practice. Advances in Anatomic Pathology, 2016, 23, 356-367.	2.4	17
41	High content screening application for cell-type specific behaviour in heterogeneous primary breast epithelial subpopulations. Breast Cancer Research, 2016, 18, 18.	2.2	9
42	Forgotten Resources â€“ The Autopsy. , 2016, , 335-348.		0
43	Genomic Analysis. , 2016, , 83-106.		0
44	Integrated genomic and transcriptomic analysis of human brain metastases identifies alterations of potential clinical significance. Journal of Pathology, 2015, 237, 363-378.	2.1	98
45	Invasive lobular carcinoma of the breast: morphology, biomarkers and â€™omics. Breast Cancer Research, 2015, 17, 12.	2.2	256
46	Using the MCF10A/MCF10CA1a Breast Cancer Progression Cell Line Model to Investigate the Effect of Active, Mutant Forms of EGFR in Breast Cancer Development and Treatment Using Gefitinib. PLoS ONE, 2015, 10, e0125232.	1.1	27
47	Heregulin-HER3-HER2 signaling promotes matrix metalloproteinase-dependent blood-brain-barrier transendothelial migration of human breast cancer cell lines. Oncotarget, 2015, 6, 3932-3946.	0.8	60
48	Metastatic progression of breast cancer: insights from 50â€‰years of autopsies. Journal of Pathology, 2014, 232, 23-31.	2.1	161
49	Evaluating the repair of DNA derived from formalin-fixed paraffin-embedded tissues prior to genomic profiling by SNPâ€“CGH analysis. Laboratory Investigation, 2013, 93, 701-710.	1.7	26
50	Thrombospondin-4 expression is activated during the stromal response to invasive breast cancer. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2013, 463, 535-545.	1.4	54
51	Expression Profiling of Archival Tumors for Long-term Health Studies. Clinical Cancer Research, 2012, 18, 6136-6146.	3.2	32
52	Gene expression profiling of tumour epithelial and stromal compartments during breast cancer progression. Breast Cancer Research and Treatment, 2012, 135, 153-165.	1.1	111
53	Molecular pathology of pre-invasive breast disease in the screening setting: application in diagnosis and management. Diagnostic Histopathology, 2012, 18, 64-69.	0.2	0
54	The Stem Cell Marker CD133 Associates with Enhanced Colony Formation and Cell Motility in Colorectal Cancer. PLoS ONE, 2010, 5, e10714.	1.1	79

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55	The Apc1322T Mouse Develops Severe Polyposis Associated With Submaximal Nuclear Î²-Catenin Expression. Gastroenterology, 2009, 136, 2204-2213.e13.	0.6	55
56	Apc mice: Models, modifiers and mutants. Pathology Research and Practice, 2008, 204, 479-490.	1.0	143
57	Pregnancy does not influence colonic polyp multiplicity but may modulate upper gastrointestinal disease in patients with FAP. Journal of Medical Genetics, 2007, 44, 541-544.	1.5	1
58	Activation of AKT and nuclear accumulation of wild type TP53 and MDM2 in anal squamous cell carcinoma. International Journal of Cancer, 2007, 121, 2668-2673.	2.3	35
59	A Novel Exon Duplication Event Leading to a Truncating Germ-line Mutation of the APC Gene in a Familial Adenomatous Polyposis Family. Familial Cancer, 2006, 5, 205-208.	0.9	12
60	Genetic determinants modulate susceptibility to pregnancy-associated tumourigenesis in a recombinant line of Min mice. Human Molecular Genetics, 2006, 15, 3429-3435.	1.4	17
61	Alternatively Spliced Products of the Human Kinesin Light Chain 1 (*KNS2*) Gene. Traffic, 2003, 4, 576-580.	1.3	27