

Cathy B Moelans

List of Publications by Year in descending order

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Version: 2024-02-01

70
papers

1,890
citations

257450

24
h-index

276875

41
g-index

72
all docs

72
docs citations

72
times ranked

3292
citing authors

#	ARTICLE	IF	CITATIONS
1	Receptor Conversion in Distant Breast Cancer Metastases: A Systematic Review and Meta-analysis. <i>Journal of the National Cancer Institute</i> , 2018, 110, 568-580.	6.3	198
2	Frequent promoter hypermethylation of <i>BRCA2</i> , <i>CDH13</i> , <i>MSH6</i> , <i>PAX5</i> , <i>PAX6</i> and <i>WT1</i> in ductal carcinoma <i>in situ</i> and invasive breast cancer. <i>Journal of Pathology</i> , 2011, 225, 222-231.	4.5	118
3	Absence of chromosome 17 polysomy in breast cancer: analysis by CEP17 chromogenic <i>in situ</i> hybridization and multiplex ligation-dependent probe amplification. <i>Breast Cancer Research and Treatment</i> , 2010, 120, 1-7.	2.5	104
4	Molecular profiling of invasive breast cancer by multiplex ligation-dependent probe amplification-based copy number analysis of tumor suppressor and oncogenes. <i>Modern Pathology</i> , 2010, 23, 1029-1039.	5.5	90
5	Prognostic value of estrogen receptor $\hat{\pm}$ and progesterone receptor conversion in distant breast cancer metastases. <i>Cancer</i> , 2012, 118, 4929-4935.	4.1	81
6	Influence of decalcification procedures on immunohistochemistry and molecular pathology in breast cancer. <i>Modern Pathology</i> , 2016, 29, 1460-1470.	5.5	62
7	Mutation Profiling of Key Cancer Genes in Primary Breast Cancers and Their Distant Metastases. <i>Cancer Research</i> , 2018, 78, 3112-3121.	0.9	57
8	Oncogene amplification in male breast cancer: analysis by multiplex ligation-dependent probe amplification. <i>Breast Cancer Research and Treatment</i> , 2012, 135, 49-58.	2.5	53
9	Promoter hypermethylation in male breast cancer: analysis by multiplex ligation-dependent probe amplification. <i>Breast Cancer Research</i> , 2012, 14, R101.	5.0	49
10	Formaldehyde substitute fixatives: effects on nucleic acid preservation. <i>Journal of Clinical Pathology</i> , 2011, 64, 960-967.	2.0	48
11	Formaldehyde Substitute Fixatives. <i>American Journal of Clinical Pathology</i> , 2011, 136, 548-556.	0.7	48
12	Frequent discordance in PD-1 and PD-L1 expression between primary breast tumors and their matched distant metastases. <i>Clinical and Experimental Metastasis</i> , 2019, 36, 29-37.	3.3	47
13	Low frequency of HER2 amplification and overexpression in early onset gastric cancer. <i>Cellular Oncology (Dordrecht)</i> , 2011, 34, 89-95.	4.4	41
14	Simultaneous detection of TOP2A and HER2 gene amplification by multiplex ligation-dependent probe amplification in breast cancer. <i>Modern Pathology</i> , 2010, 23, 62-70.	5.5	38
15	Molecular differences between ductal carcinoma <i>in situ</i> and adjacent invasive breast carcinoma: a multiplex ligation-dependent probe amplification study. <i>Cellular Oncology (Dordrecht)</i> , 2011, 34, 475-482.	4.4	38
16	Validation of DNA promoter hypermethylation biomarkers in breast cancer – a short report. <i>Cellular Oncology (Dordrecht)</i> , 2014, 37, 297-303.	4.4	37
17	Clinical relevance of copy number profiling in oral and oropharyngeal squamous cell carcinoma. <i>Cancer Medicine</i> , 2015, 4, 1525-1535.	2.8	37
18	HER-2/ <i>neu</i> Testing and Therapy in Gastroesophageal Adenocarcinoma. <i>Pathology Research International</i> , 2011, 2011, 1-10.	1.4	35

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19	Genomic evolution from primary breast carcinoma to distant metastasis: Few copy number changes of breast cancer related genes. <i>Cancer Letters</i> , 2014, 344, 138-146.	7.2	34
20	HER-2/neu amplification testing in breast cancer by Multiplex Ligation-dependent Probe Amplification: influence of manual- and laser microdissection. <i>BMC Cancer</i> , 2009, 9, 4.	2.6	32
21	Upregulation of Claudin-4, CAIX and GLUT-1 in distant breast cancer metastases. <i>BMC Cancer</i> , 2014, 14, 864.	2.6	32
22	Global transcriptional analysis identifies a novel role for SOX4 in tumor-induced angiogenesis. <i>ELife</i> , 2018, 7, .	6.0	32
23	Progressive APOBEC3B mRNA expression in distant breast cancer metastases. <i>PLoS ONE</i> , 2017, 12, e0171343.	2.5	31
24	A Novel Diagnostic Tool for Selecting Patients With Mesenchymal-Type Colon Cancer Reveals Intratumor Subtype Heterogeneity. <i>Journal of the National Cancer Institute</i> , 2017, 109, .	6.3	30
25	HER-2/neu amplification testing in breast cancer by multiplex ligation-dependent probe amplification in comparison with immunohistochemistry and in situ hybridization. <i>Cellular Oncology</i> , 2009, 31, 1-10.	1.9	28
26	The molecular genetic make-up of male breast cancer. <i>Endocrine-Related Cancer</i> , 2019, 26, 779-794.	3.1	27
27	Promoter hypermethylation using 24-gene array in early head and neck cancer. <i>Epigenetics</i> , 2014, 9, 1220-1227.	2.7	24
28	Unravelling site-specific breast cancer metastasis: a microRNA expression profiling study. <i>Oncotarget</i> , 2017, 8, 3111-3123.	1.8	24
29	DNA promoter hypermethylation in nipple fluid: a potential tool for early breast cancer detection. <i>Oncotarget</i> , 2016, 7, 24778-24791.	1.8	24
30	Optimal Fixation Conditions and DNA Extraction Methods for MLPA Analysis on FFPE Tissue-Derived DNA. <i>American Journal of Clinical Pathology</i> , 2017, 147, aqw205.	0.7	22
31	Methylation-Specific Multiplex Ligation-Dependent Probe Amplification (MS-MLPA). <i>Methods in Molecular Biology</i> , 2018, 1708, 537-549.	0.9	22
32	Molecular Differences between Ductal Carcinoma <i>In Situ</i> and Adjacent Invasive Breast Carcinoma: A Multiplex Ligation-Dependent Probe Amplification Study. <i>Analytical Cellular Pathology</i> , 2010, 33, 165-173.	1.4	21
33	Clonal intratumor heterogeneity of promoter hypermethylation in breast cancer by MS-MLPA. <i>Modern Pathology</i> , 2014, 27, 869-874.	5.5	20
34	Analysis of copy number changes on chromosome 16q in male breast cancer by multiplex ligation-dependent probe amplification. <i>Modern Pathology</i> , 2013, 26, 1461-1467.	5.5	19
35	<sc>FOXA</sc> 1 levels are decreased in pleural breast cancer metastases after adjuvant endocrine therapy, and this is associated with poor outcome. <i>Molecular Oncology</i> , 2018, 12, 1884-1894.	4.6	19
36	ESR1 amplification is rare in breast cancer and is associated with high grade and high proliferation: a multiplex ligation-dependent probe amplification study. <i>Cellular Oncology (Dordrecht)</i> , 2011, 34, 489-494.	4.4	18

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37	ESR1 Amplification is Rare in Breast Cancer and is Associated with High Grade and High Proliferation: A Multiplex Ligation-Dependent Probe Amplification Study. <i>Analytical Cellular Pathology</i> , 2010, 33, 13-18.	1.4	16
38	Implications of rarity of chromosome 17 polysomy in breast cancer. <i>Lancet Oncology</i> , The, 2011, 12, 1087-1089.	10.7	16
39	Molecular differences between ductal carcinoma in situ and adjacent invasive breast carcinoma: a multiplex ligation-dependent probe amplification study. <i>Analytical Cellular Pathology</i> , 2010, 33, 165-73.	1.4	16
40	ESR1 Amplification in Breast Cancer by Optimized RNase FISH: Frequent but Low-Level and Heterogeneous. <i>PLoS ONE</i> , 2013, 8, e84189.	2.5	14
41	Loss of steroid hormone receptors is common in malignant pleural and peritoneal effusions of breast cancer patients treated with endocrine therapy. <i>Oncotarget</i> , 2017, 8, 55550-55561.	1.8	14
42	Copy number changes at 8p11-12 predict adverse clinical outcome and chemo- and radiotherapy response in breast cancer. <i>Oncotarget</i> , 2018, 9, 17078-17092.	1.8	14
43	Chromosome 17 copy number changes in male breast cancer. <i>Cellular Oncology (Dordrecht)</i> , 2015, 38, 237-245.	4.4	12
44	Promoter hypermethylation profiling of distant breast cancer metastases. <i>Breast Cancer Research and Treatment</i> , 2015, 151, 41-55.	2.5	11
45	Methylation biomarkers for pleomorphic lobular breast cancer - a short report. <i>Cellular Oncology (Dordrecht)</i> , 2015, 38, 397-405.	4.4	10
46	PD-1 and PD-L1 Expression in Male Breast Cancer in Comparison with Female Breast Cancer. <i>Targeted Oncology</i> , 2018, 13, 769-777.	3.6	10
47	Role of columnar cell lesions in breast carcinogenesis: analysis of chromosome 16 copy number changes by multiplex ligation-dependent probe amplification. <i>Modern Pathology</i> , 2018, 31, 1816-1833.	5.5	10
48	Lessons Learned from Setting Up a Prospective, Longitudinal, Multicenter Study with Women at High Risk for Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 441-449.	2.5	10
49	ESR1 amplification is rare in breast cancer and is associated with high grade and high proliferation: a multiplex ligation-dependent probe amplification study. <i>Analytical Cellular Pathology</i> , 2010, 33, 13-8.	1.4	10
50	Molecular profile of nasopharyngeal carcinoma: analysing tumour suppressor gene promoter hypermethylation by multiplex ligation-dependent probe amplification. <i>Journal of Clinical Pathology</i> , 2018, 71, 351-359.	2.0	8
51	Methylation Profile of X-Chromosome-Related Genes in Male Breast Cancer. <i>Frontiers in Oncology</i> , 2020, 10, 784.	2.8	8
52	Promoter hypermethylation in ductal carcinoma in situ of the male breast. <i>Endocrine-Related Cancer</i> , 2019, 26, 575-584.	3.1	8
53	CEP17 copy number increase does not indicate polysomy 17. <i>Journal of Clinical Pathology</i> , 2014, 67, 454.1-455.	2.0	7
54	Validation of a fully automated HER2 staining kit in breast cancer. <i>Cellular Oncology</i> , 2010, 32, 149-55.	1.9	7

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55	Nipple Aspirate Fluid at a Glance. <i>Cancers</i> , 2022, 14, 159.	3.7	7
56	Copy number profiling of oncogenes in ductal carcinoma in situ of the male breast. <i>Endocrine-Related Cancer</i> , 2018, 25, 173-184.	3.1	6
57	Application of Nipple Aspirate Fluid miRNA Profiles for Early Breast Cancer Detection and Management. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5814.	4.1	6
58	OUP accepted manuscript. <i>Clinical Chemistry</i> , 2022, , .	3.2	5
59	The Physiological MicroRNA Landscape in Nipple Aspirate Fluid: Differences and Similarities with Breast Tissue, Breast Milk, Plasma and Serum. <i>International Journal of Molecular Sciences</i> , 2020, 21, 8466.	4.1	4
60	The changing microRNA landscape by color and cloudiness: a cautionary tale for nipple aspirate fluid biomarker analysis. <i>Cellular Oncology (Dordrecht)</i> , 2021, 44, 1339-1349.	4.4	4
61	Differences in cancer gene copy number alterations between Epstein-Barr virus-positive and Epstein-Barr virus-negative nasopharyngeal carcinoma. <i>Head and Neck</i> , 2018, 40, 1986-1998.	2.0	3
62	Amplification Testing in Breast Cancer by Multiplex Ligation-Dependent Probe Amplification of Microdissected Tissue. <i>Methods in Molecular Biology</i> , 2011, 755, 107-118.	0.9	3
63	Genomic copy number alterations as biomarkers for triple negative pregnancy-associated breast cancer. <i>Cellular Oncology (Dordrecht)</i> , 2022, 45, 591-600.	4.4	3
64	Added Value of HER-2 Amplification Testing by Multiplex Ligation-Dependent Probe Amplification in Invasive Breast Cancer. <i>PLoS ONE</i> , 2013, 8, e82018.	2.5	2
65	Heterogeneity in Signaling Pathway Activity within Primary and between Primary and Metastatic Breast Cancer. <i>Cancers</i> , 2021, 13, 1345.	3.7	2
66	HER-2/neu Amplification Testing in Breast Cancer by Multiplex Ligation-Dependent Probe Amplification in Comparison with Immunohistochemistry and In Situ Hybridization. <i>Analytical Cellular Pathology</i> , 2009, 31, 1-10.	1.4	2
67	Validation of a Fully Automated HER2 Staining Kit in Breast Cancer. <i>Analytical Cellular Pathology</i> , 2010, 32, 149-155.	1.4	1
68	Multiplex Ligation-Dependent Probe Amplification to Detect HER2 Amplification in Breast Cancer: New Insights in Optimal Cut-Off Value. <i>Analytical Cellular Pathology</i> , 2010, 32, 311-312.	1.4	1
69	Response to A. Matikas et al.. <i>Journal of the National Cancer Institute</i> , 2018, 110, 1282-1283.	6.3	0
70	Patient-centered research: how do women tolerate nipple fluid aspiration as a potential screening tool for breast cancer?. <i>BMC Cancer</i> , 2022, 22, .	2.6	0