

Martha R Stampfer

List of Publications by Year in descending order

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85
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5,799
citations

61984

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86
all docs

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docs citations

86
times ranked

7436
citing authors

#	ARTICLE	IF	CITATIONS
1	Early growth response 2 (EGR2) is a novel regulator of the senescence programme. <i>Aging Cell</i> , 2021, 20, e13318.	6.7	16
2	Breast-Specific Molecular Clocks Comprised of <i>ELF5</i> Expression and Promoter Methylation Identify Individuals Susceptible to Cancer Initiation. <i>Cancer Prevention Research</i> , 2021, 14, 779-794.	1.5	11
3	Evidence for accelerated aging in mammary epithelia of women carrying germline BRCA1 or BRCA2 mutations. <i>Nature Aging</i> , 2021, 1, 838-849.	11.6	27
4	AXL Is a Driver of Stemness in Normal Mammary Gland and Breast Cancer. <i>IScience</i> , 2020, 23, 101649.	4.1	20
5	Superresolution microscopy reveals linkages between ribosomal DNA on heterologous chromosomes. <i>Journal of Cell Biology</i> , 2019, 218, 2492-2513.	5.2	40
6	Breast Tissue Biology Expands the Possibilities for Prevention of Age-Related Breast Cancers. <i>Frontiers in Cell and Developmental Biology</i> , 2019, 7, 174.	3.7	6
7	Experimental and pan-cancer genome analyses reveal widespread contribution of acrylamide exposure to carcinogenesis in humans. <i>Genome Research</i> , 2019, 29, 521-531.	5.5	57
8	Genetic variation and radiation quality impact cancer promoting cellular phenotypes in response to HZE exposure. <i>Life Sciences in Space Research</i> , 2019, 20, 101-112.	2.3	2
9	Delayed γ H2AX foci disappearance in mammary epithelial cells from aged women reveals an age-associated DNA repair defect. <i>Aging</i> , 2019, 11, 1510-1523.	3.1	13
10	Characterizing cellular mechanical phenotypes with mechano-node-pore sensing. <i>Microsystems and Nanoengineering</i> , 2018, 4, .	7.0	38
11	High-Dimensional Phenotyping Identifies Age-Emergent Cells in Human Mammary Epithelia. <i>Cell Reports</i> , 2018, 23, 1205-1219.	6.4	39
12	Different culture media modulate growth, heterogeneity, and senescence in human mammary epithelial cell cultures. <i>PLoS ONE</i> , 2018, 13, e0204645.	2.5	13
13	Quantitative phosphoproteomic analysis reveals reciprocal activation of receptor tyrosine kinases between cancer epithelial cells and stromal fibroblasts. <i>Clinical Proteomics</i> , 2018, 15, 21.	2.1	15
14	Microenvironment-Induced Non-sporadic Expression of the AXL and cKIT Receptors Are Related to Epithelial Plasticity and Drug Resistance. <i>Frontiers in Cell and Developmental Biology</i> , 2018, 6, 41.	3.7	22
15	Age-related gene expression in luminal epithelial cells is driven by a microenvironment made from myoepithelial cells. <i>Aging</i> , 2017, 9, 2026-2051.	3.1	21
16	184AA3: a xenograft model of ER+ breast adenocarcinoma. <i>Breast Cancer Research and Treatment</i> , 2016, 155, 37-52.	2.5	8
17	Delineating transcriptional networks of prognostic gene signatures refines treatment recommendations for lymph node-negative breast cancer patients. <i>FEBS Journal</i> , 2015, 282, 3455-3473.	4.7	12
18	The senescent methylome and its relationship with cancer, ageing and germline genetic variation in humans. <i>Genome Biology</i> , 2015, 16, 194.	8.8	40

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19	Age and the means of bypassing stasis influence the intrinsic subtype of immortalized human mammary epithelial cells. <i>Frontiers in Cell and Developmental Biology</i> , 2015, 3, 13.	3.7	25
20	A lincRNA connected to cell mortality and epigenetically-silenced in most common human cancers. <i>Epigenetics</i> , 2015, 10, 1074-1083.	2.7	28
21	Exome-wide mutation profile in benzo[a]pyrene-derived post-stasis and immortal human mammary epithelial cells. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2014, 775-776, 48-54.	1.7	29
22	Immortalization of normal human mammary epithelial cells in two steps by direct targeting of senescence barriers does not require gross genomic alterations. <i>Cell Cycle</i> , 2014, 13, 3423-3435.	2.6	60
23	Cellular senescence mediated by p16INK4A-coupled miRNA pathways. <i>Nucleic Acids Research</i> , 2014, 42, 1606-1618.	14.5	63
24	Age-Related Dysfunction in Mechanotransduction Impairs Differentiation of Human Mammary Epithelial Progenitors. <i>Cell Reports</i> , 2014, 7, 1926-1939.	6.4	74
25	Aging phenotypes in cultured normal human mammary epithelial cells are correlated with decreased telomerase activity independent of telomere length. <i>Genome Integrity</i> , 2013, 4, 4.	1.0	8
26	The Regulation of SOX7 and Its Tumor Suppressive Role in Breast Cancer. <i>American Journal of Pathology</i> , 2013, 183, 1645-1653.	3.8	52
27	DNA Repair Gene Patterns as Prognostic and Predictive Factors in Molecular Breast Cancer Subtypes. <i>Oncologist</i> , 2013, 18, 1063-1073.	3.7	75
28	Processing of Human Reduction Mammoplasty and Mastectomy Tissues for Cell Culture. <i>Journal of Visualized Experiments</i> , 2013, , .	0.3	52
29	Common chromosome fragile sites in human and murine epithelial cells and <i>FHIT/FRA3B</i> loss-induced global genome instability. <i>Genes Chromosomes and Cancer</i> , 2013, 52, 1017-1029.	2.8	54
30	Constitutive CCND1/CDK2 Activity Substitutes for p53 Loss, or MYC or Oncogenic RAS Expression in the Transformation of Human Mammary Epithelial Cells. <i>PLoS ONE</i> , 2013, 8, e53776.	2.5	22
31	miRNA Gene Promoters Are Frequent Targets of Aberrant DNA Methylation in Human Breast Cancer. <i>PLoS ONE</i> , 2013, 8, e54398.	2.5	110
32	Accumulation of Multipotent Progenitors with a Basal Differentiation Bias during Aging of Human Mammary Epithelia. <i>Cancer Research</i> , 2012, 72, 3687-3701.	0.9	94
33	Cell-Type Specific DNA Methylation Patterns Define Human Breast Cellular Identity. <i>PLoS ONE</i> , 2012, 7, e52299.	2.5	22
34	Oncogenes induce senescence with incomplete growth arrest and suppress the DNA damage response in immortalized cells. <i>Aging Cell</i> , 2011, 10, 949-961.	6.7	35
35	Epigenetic regulation of normal human mammary cell type-specific miRNAs. <i>Genome Research</i> , 2011, 21, 2026-2037.	5.5	68
36	TGF- β 2 signaling engages an ATM-CHK2-p53-independent RAS-induced senescence and prevents malignant transformation in human mammary epithelial cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 8668-8673.	7.1	86

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37	Self-organization is a dynamic and lineage-intrinsic property of mammary epithelial cells. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 3264-3269.	7.1	52
38	Role for DNA Methylation in the Regulation of miR-200c and miR-141 Expression in Normal and Cancer Cells. PLoS ONE, 2010, 5, e8697.	2.5	268
39	Exon-Level Microarray Analyses Identify Alternative Splicing Programs in Breast Cancer. Molecular Cancer Research, 2010, 8, 961-974.	3.4	121
40	Primary Cilium-Dependent and -Independent Hedgehog Signaling Inhibits p16INK4A. Molecular Cell, 2010, 40, 533-547.	9.7	52
41	Molecular Distinctions between Stasis and Telomere Attrition Senescence Barriers Shown by Long-term Culture of Normal Human Mammary Epithelial Cells. Cancer Research, 2009, 69, 7557-7568.	0.9	144
42	Stepwise DNA Methylation Changes Are Linked to Escape from Defined Proliferation Barriers and Mammary Epithelial Cell Immortalization. Cancer Research, 2009, 69, 5251-5258.	0.9	113
43	Human mammary progenitor cell fate decisions are products of interactions with combinatorial microenvironments. Integrative Biology (United Kingdom), 2009, 1, 70-79.	1.3	166
44	Cell type-specific DNA methylation patterns in the human breast. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 14076-14081.	7.1	210
45	Karyotypic Instability and Centrosome Aberrations in the Progeny of Finite Life-Span Human Mammary Epithelial Cells Exposed to Sparsely or Densely Ionizing Radiation. Radiation Research, 2008, 170, 23-32.	1.5	28
46	Inactivation of p53 Function in Cultured Human Mammary Epithelial Cells Turns the Telomere-Length Dependent Senescence Barrier from Agonescence into Crisis. Cell Cycle, 2007, 6, 1927-1936.	2.6	65
47	Transcriptional changes associated with breast cancer occur as normal human mammary epithelial cells overcome senescence barriers and become immortalized. Molecular Cancer, 2007, 6, 7.	19.2	44
48	ZNF652, A Novel Zinc Finger Protein, Interacts with the Putative Breast Tumor Suppressor CBFA2T3 to Repress Transcription. Molecular Cancer Research, 2006, 4, 655-665.	3.4	50
49	Caspase-independent cytochrome c release is a sensitive measure of low-level apoptosis in cell culture models. Aging Cell, 2005, 4, 217-222.	6.7	26
50	Accumulation and altered localization of telomere-associated protein TRF2 in immortally transformed and tumor-derived human breast cells. Oncogene, 2005, 24, 3369-3376.	5.9	37
51	Chromatin Inactivation Precedes De Novo DNA Methylation during the Progressive Epigenetic Silencing of the RASSF1A Promoter. Molecular and Cellular Biology, 2005, 25, 3923-3933.	2.3	123
52	In situ analyses of genome instability in breast cancer. Nature Genetics, 2004, 36, 984-988.	21.4	337
53	Loss of p53 function accelerates acquisition of telomerase activity in indefinite lifespan human mammary epithelial cell lines. Oncogene, 2003, 22, 5238-5251.	5.9	63
54	Human epithelial cell immortalization as a step in carcinogenesis. Cancer Letters, 2003, 194, 199-208.	7.2	70

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55	Raf-1-induced growth arrest in human mammary epithelial cells is p16-independent and is overcome in immortal cells during conversion. <i>Oncogene</i> , 2002, 21, 6328-6339.	5.9	82
56	Normal human mammary epithelial cells spontaneously escape senescence and acquire genomic changes. <i>Nature</i> , 2001, 409, 633-637.	27.8	604
57	Reduction of Cdc25A contributes to cyclin E1-Cdk2 inhibition at senescence in human mammary epithelial cells. <i>Oncogene</i> , 2000, 19, 5314-5323.	5.9	51
58	Culture models of human mammary epithelial cell transformation. <i>Journal of Mammary Gland Biology and Neoplasia</i> , 2000, 5, 365-378.	2.7	45
59	Establishment and Characterization of a Breast Cell Strain Containing a BRCA1 185delAG Mutation. <i>Gynecologic Oncology</i> , 2000, 77, 121-128.	1.4	11
60	Interindividual variation in CYP1A1 expression in breast tissue and the role of genetic polymorphism. <i>Carcinogenesis</i> , 2000, 21, 2119-2122.	2.8	31
61	Viral oncogenes accelerate conversion to immortality of cultured conditionally immortal human mammary epithelial cells. <i>Oncogene</i> , 1999, 18, 2169-2180.	5.9	44
62	Increased p16 expression with first senescence arrest in human mammary epithelial cells and extended growth capacity with p16 inactivation. <i>Oncogene</i> , 1998, 17, 199-205.	5.9	249
63	Gradual Phenotypic Conversion Associated with immortalization of Cultured Human Mammary Epithelial Cells. <i>Molecular Biology of the Cell</i> , 1997, 8, 2391-2405.	2.1	64
64	Insulin receptor overexpression in 184B5 human mammary epithelial cells induces a ligand-dependent transformed phenotype. <i>Journal of Cellular Biochemistry</i> , 1995, 57, 666-669.	2.6	59
65	Oncogene-induced basement membrane invasiveness in human mammary epithelial cells. <i>Clinical and Experimental Metastasis</i> , 1994, 12, 181-194.	3.3	78
66	Phospholipases A2 in ras-transformed and immortalized human mammary epithelial cells. <i>Cancer Letters</i> , 1994, 86, 11-21.	7.2	23
67	Growth, differentiation, and transformation of human mammary epithelial cells in culture. <i>Cancer Treatment and Research</i> , 1994, 71, 29-48.	0.5	11
68	TGF β induction of extracellular matrix associated proteins in normal and transformed human mammary epithelial cells in culture is independent of growth effects. <i>Journal of Cellular Physiology</i> , 1993, 155, 210-221.	4.1	65
69	Blockage of EGF Receptor Signal Transduction Causes Reversible Arrest of Normal and Immortal Human Mammary Epithelial Cells with Synchronous Reentry into the Cell Cycle. <i>Experimental Cell Research</i> , 1993, 208, 175-188.	2.6	77
70	p53 Mutations in human immortalized epithelial cell lines. <i>Carcinogenesis</i> , 1993, 14, 833-839.	2.8	406
71	Effects of Sequence of Thioated Oligonucleotides on Cultured Human Mammary Epithelial Cells. <i>Antisense Research and Development</i> , 1993, 3, 67-77.	3.1	133
72	Role of DNA repair in malignant neoplastic transformation of human mammary epithelial cells in culture. <i>Carcinogenesis</i> , 1992, 13, 1137-1141.	2.8	15

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73	Chromosomal changes in cultured human epithelial cells transformed by low- and high-let radiation. <i>Advances in Space Research</i> , 1992, 12, 127-136.	2.6	8
74	Stromal influences on transformation of human mammary epithelial cells overexpressingc-myc and SV40T. <i>Journal of Cellular Physiology</i> , 1990, 145, 207-216.	4.1	53
75	Radiation Studies on Sensitivity and Repair of Human Mammary Epithelial Cells. <i>International Journal of Radiation Biology</i> , 1989, 56, 605-609.	1.8	13
76	Chromosome analyses of human mammary epithelial cells at stages of chemical-induced transformation progression to immortality. <i>Cancer Genetics and Cytogenetics</i> , 1989, 37, 249-261.	1.0	69
77	Human mammary epithelial cells in culture: differentiation and transformation. <i>Cancer Treatment and Research</i> , 1988, 40, 1-24.	0.5	47
78	Response to Doxorubicin of Cultured Normal and Cancerous Human Mammary Epithelial Cells. <i>Journal of the National Cancer Institute</i> , 1985, , .	6.3	11
79	Factors influencing benzo[a]pyrene metabolism in human mammary epithelial cells in culture. <i>Carcinogenesis</i> , 1985, 6, 1017-1022.	2.8	21
80	Growth of diploid cells from breast cancers. <i>Cancer Genetics and Cytogenetics</i> , 1985, 16, 49-64.	1.0	71
81	Thioesterase II, a New Marker Enzyme for Human Cells of Breast Epithelial Origin. <i>Journal of the National Cancer Institute</i> , 1984, 73, 323-329.	6.3	16
82	Response of Cultured Normal Human Mammary Epithelial Cells to X Rays. <i>Radiation Research</i> , 1983, 96, 476.	1.5	17
83	Cholera toxin stimulation of human mammary epithelial cells in culture. <i>In Vitro</i> , 1982, 18, 531-537.	1.2	91
84	Culture of Human Mammary Epithelial Cells. , 0, , 95-135.		6
85	Viral oncogenes accelerate conversion to immortality of cultured conditionally immortal human mammary epithelial cells. , 0, .		2