

Manuela Gariboldi

List of Publications by Year in descending order

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97
papers

9,167
citations

101543

36
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101
all docs

101
docs citations

101
times ranked

13771
citing authors

#	ARTICLE	IF	CITATIONS
1	Force Sensing on Cells and Tissues by Atomic Force Microscopy. <i>Sensors</i> , 2022, 22, 2197.	3.8	12
2	Exploring the association with disease recurrence of miRNAs predictive of colorectal cancer. <i>International Journal of Biological Markers</i> , 2022, 37, 102-109.	1.8	0
3	Preventive Anti-inflammatory Diet to Reduce Gastrointestinal Inflammation in Familial Adenomatous Polyposis Patients: A Prospective Pilot Study. <i>Cancer Prevention Research</i> , 2021, 14, 963-972.	1.5	8
4	Management of Dietary Habits and Diarrhea in Fap Individuals: A Mediterranean Low-Inflammatory Dietary Intervention. <i>Nutrients</i> , 2021, 13, 3988.	4.1	2
5	Plasma miRNA-based signatures in CRC screening programs. <i>International Journal of Cancer</i> , 2020, 146, 1164-1173.	5.1	35
6	Trial watch : the gut microbiota as a tool to boost the clinical efficacy of anticancer immunotherapy. <i>Oncolimmunology</i> , 2020, 9, 1774298.	4.6	22
7	A Pilot Low-Inflammatory Dietary Intervention to Reduce Inflammation and Improve Quality of Life in Patients With Familial Adenomatous Polyposis: Protocol Description and Preliminary Results. <i>Integrative Cancer Therapies</i> , 2019, 18, 153473541984640.	2.0	10
8	Workflow for Circulating miRNA Identification and Development in Cancer Research: Methodological Considerations. , 2018, , 103-117.		1
9	A methodological procedure for evaluating the impact of hemolysis on circulating microRNAs. <i>Oncology Letters</i> , 2017, 13, 315-320.	1.8	52
10	MIF/CD74 axis is a target for novel therapies in colon carcinomatosis. <i>Journal of Experimental and Clinical Cancer Research</i> , 2017, 36, 16.	8.6	43
11	Metformin transiently inhibits colorectal cancer cell proliferation as a result of either AMPK activation or increased ROS production. <i>Scientific Reports</i> , 2017, 7, 15992.	3.3	102
12	Abstract B19: Disruption of energy homeostasis as an approach to block the proliferation of colon carcinomatosis. , 2017, , .		0
13	Comment on "Circulating cell-free miRNAs as biomarker for triple-negative breast cancer" Methodological challenges in combining miRNAs as circulating biomarkers. <i>British Journal of Cancer</i> , 2016, 114, e5-e5.	6.4	2
14	miR-342 overexpression results in a synthetic lethal phenotype in <i>BRCA1</i> -mutant HCC1937 breast cancer cells. <i>Oncotarget</i> , 2016, 7, 18594-18604.	1.8	20
15	Integration of genome scale data for identifying new players in colorectal cancer. <i>World Journal of Gastroenterology</i> , 2016, 22, 534.	3.3	3
16	Moving from Discovery to Validation in Circulating microRNA Research. <i>International Journal of Biological Markers</i> , 2015, 30, 258-261.	1.8	2
17	The Effects of miR-20a on p21: Two Mechanisms Blocking Growth Arrest in TGF- β -Responsive Colon Carcinoma. <i>Journal of Cellular Physiology</i> , 2015, 230, 3105-3114.	4.1	46
18	DPD and UGT1A1 deficiency in colorectal cancer patients receiving triplet chemotherapy with fluoropyrimidines, oxaliplatin and irinotecan. <i>British Journal of Clinical Pharmacology</i> , 2015, 80, 581-588.	2.4	52

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19	Abstract 1161: Metformin has an inhibitory effect on cell proliferation but does not induce death in colorectal cancer. , 2015, , .		3
20	Abstract CT121: Metronomic capecitabine and bevacizumab is an active combination in patients with relapsed peritoneal pseudomyxoma. , 2015, , .		0
21	A normalization strategy for the analysis of plasma microRNA qPCR data in colorectal cancer. International Journal of Cancer, 2014, 134, 2016-2018.	5.1	8
22	Combined analysis of chromosomal instabilities and gene expression for colon cancer progression inference. Journal of Clinical Bioinformatics, 2014, 4, 2.	1.2	15
23	Circulating miR-378 in plasma: a reliable, haemolysis-independent biomarker for colorectal cancer. British Journal of Cancer, 2014, 110, 1001-1007.	6.4	118
24	Activity of temozolomide in patients with advanced chemorefractory colorectal cancer and MGMT promoter methylation. Annals of Oncology, 2014, 25, 404-408.	1.2	67
25	NqA: An R-based algorithm for the normalization and analysis of microRNA quantitative real-time polymerase chain reaction data. Analytical Biochemistry, 2014, 461, 7-9.	2.4	15
26	Circulating Free DNA in a Screening Program for Early Colorectal Cancer Detection. Tumori, 2014, 100, 115-121.	1.1	39
27	miR-342 Regulates BRCA1 Expression through Modulation of ID4 in Breast Cancer. PLoS ONE, 2014, 9, e87039.	2.5	59
28	Circulating free DNA in a screening program for early colorectal cancer detection. Tumori, 2014, 100, 115-21.	1.1	27
29	Targeting metabolism for cancer treatment and prevention: metformin, an old drug with multi-faceted effects. Oncogene, 2013, 32, 1475-1487.	5.9	204
30	Gold-Nanoparticle-Based Colorimetric Discrimination of Cancer-Related Point Mutations with Picomolar Sensitivity. ACS Nano, 2013, 7, 5530-5538.	14.6	101
31	Role of cMET in the Development and Progression of Colorectal Cancer. International Journal of Molecular Sciences, 2013, 14, 18056-18077.	4.1	47
32	Effects of Warm Ischemic Time on Gene Expression Profiling in Colorectal Cancer Tissues and Normal Mucosa. PLoS ONE, 2013, 8, e53406.	2.5	44
33	Copyâ€œNumber Alterations for Tumor Progression Inference. Lecture Notes in Computer Science, 2013, , 104-109.	1.3	11
34	miRNA Profiling in Colorectal Cancer Highlights miR-1 Involvement in MET-Dependent Proliferation. Molecular Cancer Research, 2012, 10, 504-515.	3.4	123
35	Methylation status in patients with early stage colon cancer: A new prognostic marker?. International Journal of Cancer, 2012, 130, 488-489.	5.1	10
36	Abstract 2295: miR-342 modulates ID4 expression in breast cancer. , 2012, , .		0

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37	Gene expression analysis reveals a different transcriptomic landscape in female and male breast cancer. <i>Breast Cancer Research and Treatment</i> , 2011, 127, 601-610.	2.5	88
38	Transcriptional characteristics of familial non- <i>BRCA1/BRCA2</i> breast tumors. <i>International Journal of Cancer</i> , 2011, 128, 2635-2644.	5.1	11
39	ERG Deregulation Induces PIM1 Over-Expression and Aneuploidy in Prostate Epithelial Cells. <i>PLoS ONE</i> , 2011, 6, e28162.	2.5	25
40	Abstract 3959: Growth suppression by TGF- β 2 in responsive colon carcinoma is opposed by miR-20a through mechanisms altering p21WAF1 up-regulation. , 2011, , .		0
41	Chromosome band 17q21 in breast cancer: Significant association between <i>beclin 1</i> loss and <i>HER2/NEU</i> amplification. <i>Genes Chromosomes and Cancer</i> , 2010, 49, 901-909.	2.8	41
42	Molecular markers for prediction of risk of radiation-related injury to normal tissue. <i>Journal of Nucleic Acids Investigation</i> , 2010, 1, 11.	0.8	2
43	Abstract 4038: Identification of microRNAs involved in colorectal cancer progression. , 2010, , .		0
44	Molecular markers for prediction of risk of radiation-related injury to normal tissue. <i>Journal of Nucleic Acids Investigation</i> , 2010, 1, 11.	0.8	4
45	Misbehaviour of XIST RNA in Breast Cancer Cells. <i>PLoS ONE</i> , 2009, 4, e5559.	2.5	75
46	Integrative approach for prioritizing cancer genes in sporadic colon cancer. <i>Genes Chromosomes and Cancer</i> , 2009, 48, 953-962.	2.8	47
47	Gene expression profiling integrated into network modelling reveals heterogeneity in the mechanisms of <i>BRCA1</i> tumorigenesis. <i>British Journal of Cancer</i> , 2009, 101, 1469-1480.	6.4	13
48	To Bleed or Not to Bleed. A Prediction Based on Individual Gene Profiling Combined With Dose-Volume Histogram Shapes in Prostate Cancer Patients Undergoing Three-Dimensional Conformal Radiation Therapy. <i>International Journal of Radiation Oncology Biology Physics</i> , 2009, 74, 1431-1440.	0.8	55
49	Invasiveness gene signature predicts a favorable outcome also in estrogen receptor-positive primary breast cancers treated with adjuvant tamoxifen. <i>Breast Cancer Research and Treatment</i> , 2008, 111, 389-390.	2.5	3
50	Metallothionein 1G acts as an oncosuppressor in papillary thyroid carcinoma. <i>Laboratory Investigation</i> , 2008, 88, 474-481.	3.7	60
51	Re: Molecular Basis for Estrogen Receptor β Deficiency in <i>BRCA1</i> -Linked Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2008, 100, 752-753.	6.3	2
52	Patterns and changes in gene expression following neo-adjuvant anti-estrogen treatment in estrogen receptor-positive breast cancer. <i>Endocrine-Related Cancer</i> , 2008, 15, 439-449.	3.1	16
53	Challenges in Projecting Clustering Results Across Gene Expression Profiling Datasets. <i>Journal of the National Cancer Institute</i> , 2007, 99, 1715-1723.	6.3	88
54	Analysis of gene expression identifies PLAB as a mediator of the apoptotic activity of fenretinide in human ovarian cancer cells. <i>Oncogene</i> , 2007, 26, 3952-3962.	5.9	14

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55	Transcriptional network dynamics in macrophage activation. <i>Genomics</i> , 2006, 88, 133-142.	2.9	125
56	Specific gene expression profiles distinguish among functional allelic variants of the mouse <i>Pthlh</i> gene in transfected human cancer cells. <i>Oncogene</i> , 2006, 25, 4501-4504.	5.9	8
57	Molecular predictors of response and outcome in ovarian cancer. <i>Critical Reviews in Oncology/Hematology</i> , 2006, 60, 19-37.	4.4	36
58	Regulation of lipocalin-2 gene by the cancer chemopreventive retinoid 4-HPR. <i>International Journal of Cancer</i> , 2006, 119, 1599-1606.	5.1	15
59	Mâ€CAM expression as marker of poor prognosis in epithelial ovarian cancer. <i>International Journal of Cancer</i> , 2006, 119, 1920-1926.	5.1	78
60	Gene expression profile identifies a rare epithelioid variant case of pleomorphic liposarcoma carrying FUS-CHOP transcript. <i>Histopathology</i> , 2005, 46, 334-341.	2.9	21
61	RESPONSE: Re: Limits of Predictive Models Using Microarray Data for Breast Cancer Clinical Treatment Outcome. <i>Journal of the National Cancer Institute</i> , 2005, 97, 1852-1853.	6.3	2
62	Re: Limits of Predictive Models Using Microarray Data for Breast Cancer Clinical Treatment Outcome. <i>Journal of the National Cancer Institute</i> , 2005, 97, 1851-1852.	6.3	10
63	Induction of a proinflammatory program in normal human thyrocytes by the RET/PTC1 oncogene. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 14825-14830.	7.1	311
64	Limits of Predictive Models Using Microarray Data for Breast Cancer Clinical Treatment Outcome. <i>Journal of the National Cancer Institute</i> , 2005, 97, 927-930.	6.3	110
65	The Transcriptional Landscape of the Mammalian Genome. <i>Science</i> , 2005, 309, 1559-1563.	12.6	3,227
66	Gene discovery in genetically labeled single dopaminergic neurons of the retina. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 5069-5074.	7.1	70
67	CTAB-Urea Method Purifies RNA from Melanin for cDNA Microarray Analysis. <i>Pigment Cell & Melanoma Research</i> , 2004, 17, 312-315.	3.6	26
68	Gene expression profiling of advanced ovarian cancer: characterization of a molecular signature involving fibroblast growth factor 2. <i>Oncogene</i> , 2004, 23, 8171-8183.	5.9	75
69	Alternative mutations of BRAF, RET and NTRK1 are associated with similar but distinct gene expression patterns in papillary thyroid cancer. <i>Oncogene</i> , 2004, 23, 7436-7440.	5.9	239
70	VE-Cadherin Expression and Clustering Maintain Low Levels of Survivin in Endothelial Cells. <i>American Journal of Pathology</i> , 2004, 165, 181-189.	3.8	34
71	Allele-specific patterns of the mouse parathyroid hormone-related protein: influences on cell adhesion and migration. <i>Oncogene</i> , 2003, 22, 7711-7715.	5.9	8
72	Gene expression profile of normal lungs predicts genetic predisposition to lung cancer in mice. <i>Carcinogenesis</i> , 2003, 24, 1819-1826.	2.8	10

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73	Stearoyl-CoA desaturase 1 (Scd1) gene overexpression is associated with genetic predisposition to hepatocarcinogenesis in mice and rats. <i>Carcinogenesis</i> , 2002, 23, 1933-1936.	2.8	81
74	Analysis of the mouse transcriptome based on functional annotation of 60,770 full-length cDNAs. <i>Nature</i> , 2002, 420, 563-573.	27.8	1,548
75	Functional annotation of a full-length mouse cDNA collection. <i>Nature</i> , 2001, 409, 685-690.	27.8	653
76	A cancer modifier role for parathyroid hormone-related protein. <i>Oncogene</i> , 2000, 19, 5324-5328.	5.9	28
77	Genetic mapping and analysis of mouse p27 Kip1 gene as Pas1 candidate gene. <i>Mammalian Genome</i> , 2000, 11, 338-339.	2.2	0
78	Predisposition to lung tumorigenesis. <i>Toxicology Letters</i> , 2000, 112-113, 257-263.	0.8	13
79	Analysis of loss of heterozygosity in neoplastic nodules induced by diethylnitrosamine in the resistant BFF1 rat strain. <i>Carcinogenesis</i> , 1999, 20, 1363-1368.	2.8	9
80	Linkage Disequilibrium and Physical Mapping of <i>Pas1</i> in Mice. <i>Genome Research</i> , 1999, 9, 639-646.	5.5	35
81	Analysis of the retinoic acid receptor β gene as a candidate for the pulmonary adenoma resistance 1 gene. <i>Molecular Carcinogenesis</i> , 1998, 21, 13-16.	2.7	2
82	<i>Pas1</i> is a common lung cancer susceptibility locus in three mouse strains. <i>Mammalian Genome</i> , 1997, 8, 801-804.	2.2	25
83	Genetic mapping of a pulmonary adenoma resistance locus (<i>Par1</i>) in mouse. <i>Nature Genetics</i> , 1996, 12, 455-457.	21.4	64
84	Analysis of loss of heterozygosity in murine hepatocellular tumors. <i>Molecular Carcinogenesis</i> , 1995, 13, 191-200.	2.7	23
85	Genetic mapping and expression analysis of the murine DNA ligase I gene. <i>Molecular Carcinogenesis</i> , 1995, 14, 71-74.	2.7	9
86	Mapping of the <i>Hmg1</i> gene and of seven related sequences in the mouse. <i>Mammalian Genome</i> , 1995, 6, 581-585.	2.2	21
87	Chromosome mapping of nine tropomyosin-related sequences in mice. <i>Mammalian Genome</i> , 1995, 6, 273-277.	2.2	2
88	Mapping of body weight loci on mouse Chromosome X. <i>Mammalian Genome</i> , 1995, 6, 778-781.	2.2	70
89	Different Susceptibility to Lung Tumorigenesis in Mice with an Identical <i>Kras2</i> Intron 2. <i>Genomics</i> , 1995, 29, 438-444.	2.9	33
90	Genetics of liver tumor susceptibility in mice. <i>Toxicology Letters</i> , 1995, 82-83, 613-619.	0.8	28

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91	Expression in lung tumors and genetic mapping of the novel murine protein kinase C δ . Molecular Carcinogenesis, 1994, 9, 111-113.	2.7	4
92	Comparative Mapping of the Actin-Binding Protein 280 Genes in Human and Mouse. Genomics, 1994, 21, 428-430.	2.9	21
93	Genetic Mapping of the Mouse CDC25Mm Gene, a Ras-Specific Guanine Nucleotide-Releasing Factor, to Chromosome 9. Genomics, 1994, 21, 451-453.	2.9	12
94	Multiple Loci Affect Genetic Predisposition to Hepatocarcinogenesis in Mice. Genomics, 1994, 23, 118-124.	2.9	93
95	A major susceptibility locus to murine lung carcinogenesis maps on chromosome 6. Nature Genetics, 1993, 3, 132-136.	21.4	127
96	Esophageal Carcinoma. Acta Radiologica, 1987, 28, 177-180.	1.1	2
97	Esophageal Carcinoma. Acta Radiologica, 1987, 28, 177-180.	1.1	1