

Regina M Santella

List of Publications by Year in descending order

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291
papers

16,833
citations

14615

66
h-index

24106

111
g-index

315
all docs

315
docs citations

315
times ranked

21748
citing authors

#	ARTICLE	IF	CITATIONS
1	Case-control study in ALS using the National ALS Registry: lead and agricultural chemicals are potential risk factors. <i>Amyotrophic Lateral Sclerosis and Frontotemporal Degeneration</i> , 2022, 23, 190-202.	2.2	11
2	Environmental exposure and clinical correlates of hepatocellular carcinoma in New York City: a case only study. <i>Cancer Causes and Control</i> , 2022, 33, 153-159.	1.8	1
3	Phthalates and Phenols, Leukocyte Telomere Length, and Breast Cancer Risk and Mortality in the Long Island Breast Cancer Study Project. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 117-123.	1.9	5
4	Predictors of urinary polycyclic aromatic hydrocarbon metabolites in girls from the San Francisco Bay Area. <i>Environmental Research</i> , 2022, 205, 112534.	7.7	4
5	Improvement on recovery and reproducibility for quantifying urinary mono-hydroxylated polycyclic aromatic hydrocarbons (OH-PAHs). <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2022, 1192, 123113.	2.4	5
6	Dietary Acid Load, Serum Polychlorinated Biphenyl Levels, and Mortality Following Breast Cancer in the Long Island Breast Cancer Study Project. <i>International Journal of Environmental Research and Public Health</i> , 2022, 19, 374.	2.7	5
7	The associations of healthy lifestyle index with breast cancer incidence and mortality in a population-based study. <i>Breast Cancer</i> , 2022, 29, 957-966.	3.0	10
8	Common Childhood Viruses and Pubertal Timing: The LEGACY Girls Study. <i>American Journal of Epidemiology</i> , 2021, 190, 766-778.	3.7	3
9	The Steroid Metabolome and Breast Cancer Risk in Women with a Family History of Breast Cancer: The Novel Role of Adrenal Androgens and Glucocorticoids. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 89-96.	1.9	10
10	MicroRNA-Based Cancer Mortality Risk Scoring System and hTERT Expression in Early-Stage Oral Squamous Cell Carcinoma. <i>Journal of Oncology</i> , 2021, 2021, 1-11.	1.4	1
11	PAM50- and immunohistochemistry-based subtypes of breast cancer and their relationship with breast cancer mortality in a population-based study. <i>Breast Cancer</i> , 2021, 28, 1235-1242.	3.0	7
12	Reproductive and environmental exposures and the breast cancer risk in Taiwanese women. <i>Scientific Reports</i> , 2021, 11, 15656.	3.4	6
13	Associations Between Polymorphisms in Genes Related to Oxidative Stress and DNA Repair, Interactions With Serum Antioxidants, and Prostate Cancer Risk: Results From the Prostate Cancer Prevention Trial. <i>Frontiers in Oncology</i> , 2021, 11, 808715.	2.9	5
14	DDT exposure during pregnancy and DNA methylation alterations in female offspring in the Child Health and Development Study. <i>Reproductive Toxicology</i> , 2020, 92, 138-147.	3.1	14
15	Identifying Novel Genetic Markers Through a Transcription-Wide Association Study: Can This Be a Path to Reducing the Burden of Pancreatic Cancer?. <i>Journal of the National Cancer Institute</i> , 2020, 112, 977-978.	6.4	1
16	Influence of pubertal development on urinary oxidative stress biomarkers in adolescent girls in the New York LEGACY cohort. <i>Free Radical Research</i> , 2020, 54, 431-441.	3.3	6
17	Circulating growth factor concentrations and breast cancer risk: a nested case-control study of IGF-1, IGFBP-3, and breast cancer in a family-based cohort. <i>Breast Cancer Research</i> , 2020, 22, 109.	5.1	9
18	Urinary Estrogen Metabolites and Long-Term Mortality Following Breast Cancer. <i>JNCI Cancer Spectrum</i> , 2020, 4, pkaa014.	2.8	0

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19	MicroRNA-based risk scoring system to identify early-stage oral squamous cell carcinoma patients at high-risk for cancer-specific mortality. <i>Head and Neck</i> , 2020, 42, 1699-1712.	2.0	36
20	Gene expression profiles for low-dose exposure to diethyl phthalate in rodents and humans: a translational study with implications for breast carcinogenesis. <i>Scientific Reports</i> , 2020, 10, 7067.	3.4	20
21	Plasma creatinine and oxidative stress biomarkers in amyotrophic lateral sclerosis. <i>Amyotrophic Lateral Sclerosis and Frontotemporal Degeneration</i> , 2020, 21, 263-272.	2.2	20
22	High levels of global genome methylation in patients with retinoblastoma. <i>Oncology Letters</i> , 2020, 20, 715-723.	1.8	9
23	Genetic polymorphisms of diabetes-related genes, their interaction with diabetes status, and breast cancer incidence and mortality: The Long Island Breast Cancer Study Project. <i>Molecular Carcinogenesis</i> , 2019, 58, 436-446.	2.9	14
24	Prediagnosis aspirin use, DNA methylation, and mortality after breast cancer: A population-based study. <i>Cancer</i> , 2019, 125, 3836-3844.	4.1	13
25	Changes in human peripheral blood mononuclear cell (HPBMC) populations and T-cell subsets associated with arsenic and polycyclic aromatic hydrocarbon exposures in a Bangladesh cohort. <i>PLoS ONE</i> , 2019, 14, e0220451.	2.5	20
26	An increase in circulating B cells and B cell activation markers in peripheral blood is associated with cigarette smoking in a male cohort in Bangladesh. <i>Toxicology and Applied Pharmacology</i> , 2019, 384, 114783.	2.9	7
27	Reproductive characteristics are associated with gene-specific promoter methylation status in breast cancer. <i>BMC Cancer</i> , 2019, 19, 926.	2.6	4
28	Urinary concentrations of environmental phenols and their associations with breast cancer incidence and mortality following breast cancer. <i>Environment International</i> , 2019, 130, 104890.	10.1	71
29	Assessment of arsenic and polycyclic aromatic hydrocarbon (PAH) exposures on immune function among males in Bangladesh. <i>PLoS ONE</i> , 2019, 14, e0216662.	2.5	24
30	Association of Prepubertal and Adolescent Androgen Concentrations With Timing of Breast Development and Family History of Breast Cancer. <i>JAMA Network Open</i> , 2019, 2, e190083.	6.0	9
31	Reproductive characteristics modify the association between global DNA methylation and breast cancer risk in a population-based sample of women. <i>PLoS ONE</i> , 2019, 14, e0210884.	2.5	5
32	Rasagiline for amyotrophic lateral sclerosis: A randomized, controlled trial. <i>Muscle and Nerve</i> , 2019, 59, 201-207.	2.3	36
33	Breast cancer family history and allele-specific DNA methylation in the legacy girls study. <i>Epigenetics</i> , 2018, 13, 240-250.	2.9	10
34	Smoking, Sex, and Non-Small Cell Lung Cancer: Steroid Hormone Receptors in Tumor Tissue (S0424). <i>Journal of the National Cancer Institute</i> , 2018, 110, 734-742.	6.4	32
35	Associations between polymorphisms in genes related to estrogen metabolism and function and prostate cancer risk: results from the Prostate Cancer Prevention Trial. <i>Carcinogenesis</i> , 2018, 39, 125-133.	2.8	14
36	Aflatoxin B1 exposure increases the risk of hepatocellular carcinoma associated with hepatitis C virus infection or alcohol consumption. <i>European Journal of Cancer</i> , 2018, 94, 37-46.	2.9	65

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37	Maternal cigarette smoking during pregnancy and offspring DNA methylation in midlife. <i>Epigenetics</i> , 2018, 13, 129-134.	2.9	63
38	Novel Predictors of Breast Cancer Survival Derived from miRNA Activity Analysis. <i>Clinical Cancer Research</i> , 2018, 24, 581-591.	7.2	24
39	Urinary Phthalate Metabolite Concentrations and Breast Cancer Incidence and Survival following Breast Cancer: The Long Island Breast Cancer Study Project. <i>Environmental Health Perspectives</i> , 2018, 126, 047013.	8.2	43
40	Response to H. Nabi et al.. <i>Journal of the National Cancer Institute</i> , 2018, 110, 1424-1425.	6.4	0
41	Dietary lipids differentially modulate the initiation of experimental breast carcinogenesis through their influence on hepatic xenobiotic metabolism and DNA damage in the mammary gland. <i>Journal of Nutritional Biochemistry</i> , 2017, 43, 68-77.	4.3	5
42	Modification of the association between recreational physical activity and survival after breast cancer by promoter methylation in breast cancer-related genes. <i>Breast Cancer Research</i> , 2017, 19, 19.	5.1	18
43	Aflatoxin B ₁ exposure increases the risk of cirrhosis and hepatocellular carcinoma in chronic hepatitis B virus carriers. <i>International Journal of Cancer</i> , 2017, 141, 711-720.	5.4	90
44	Arsenic exposures alter clinical indicators of anemia in a male population of smokers and non-smokers in Bangladesh. <i>Toxicology and Applied Pharmacology</i> , 2017, 331, 62-68.	2.9	22
45	Global Level of Plasma DNA Methylation is Associated with Overall Survival in Patients with Hepatocellular Carcinoma. <i>Annals of Surgical Oncology</i> , 2017, 24, 3788-3795.	2.0	19
46	Dependence of cancer risk from environmental exposures on underlying genetic susceptibility: an illustration with polycyclic aromatic hydrocarbons and breast cancer. <i>British Journal of Cancer</i> , 2017, 116, 1229-1233.	6.6	55
47	Association between variants in genes involved in the immune response and prostate cancer risk in men randomized to the finasteride arm in the Prostate Cancer Prevention Trial. <i>Prostate</i> , 2017, 77, 908-919.	2.3	22
48	Genetic polymorphisms of phase I metabolizing enzyme genes, their interaction with lifetime grilled and smoked meat intake, and breast cancer incidence. <i>Annals of Epidemiology</i> , 2017, 27, 208-214.e1.	2.1	16
49	Plasma DNA methylation marker and hepatocellular carcinoma risk prediction model for the general population. <i>Carcinogenesis</i> , 2017, 38, 1021-1028.	2.8	40
50	Polycyclic aromatic hydrocarbons and postmenopausal breast cancer: An evaluation of effect measure modification by body mass index and weight change. <i>Environmental Research</i> , 2017, 152, 17-25.	7.7	25
51	Environmental Tobacco Smoke Exposure and Survival Following Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 278-280.	1.9	2
52	Telomere Length and Risk of Hepatocellular Carcinoma: A Nested Case-Control Study in Taiwan Cancer Screening Program Cohort. <i>Anticancer Research</i> , 2017, 37, 637-644.	1.1	6
53	DNA Methylation in Breast Tumor from High-risk Women in the Breast Cancer Family Registry. <i>Anticancer Research</i> , 2017, 37, 659-664.	1.1	7
54	Vehicular Traffic-Related Polycyclic Aromatic Hydrocarbon Exposure and Breast Cancer Incidence: The Long Island Breast Cancer Study Project (LIBCSP). <i>Environmental Health Perspectives</i> , 2016, 124, 30-38.	8.2	78

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55	Key genes involved in the immune response are generally not associated with intraprostatic inflammation in men without a prostate cancer diagnosis: Results from the prostate cancer prevention trial. <i>Prostate</i> , 2016, 76, 565-574.	2.3	5
56	Polymorphisms in DNA repair genes, traffic-related polycyclic aromatic hydrocarbon exposure and breast cancer incidence. <i>International Journal of Cancer</i> , 2016, 139, 310-321.	5.4	28
57	Long-term Diet and Biomarker Changes after a Short-term Intervention among Hispanic Breast Cancer Survivors: The <i>Cocinar Para Su Salud</i> Randomized Controlled Trial. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 1491-1502.	1.9	37
58	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016, 7, 11375.	13.2	96
59	Urinary 8-oxo-7,8-dihydro-2-deoxyguanosine analysis by an improved ELISA: An inter-laboratory comparison study. <i>Free Radical Biology and Medicine</i> , 2016, 95, 169-179.	4.5	25
60	NEpiC: a network-assisted algorithm for epigenetic studies using mean and variance combined signals. <i>Nucleic Acids Research</i> , 2016, 44, e134-e134.	14.0	13
61	Sources of polycyclic aromatic hydrocarbons are associated with gene-specific promoter methylation in women with breast cancer. <i>Environmental Research</i> , 2016, 145, 93-100.	7.7	53
62	Exposure to multiple sources of polycyclic aromatic hydrocarbons and breast cancer incidence. <i>Environment International</i> , 2016, 89-90, 185-192.	10.1	126
63	DNA methylation modifies the association between obesity and survival after breast cancer diagnosis. <i>Breast Cancer Research and Treatment</i> , 2016, 156, 183-194.	2.5	18
64	Environmental Exposures and Hepatocellular Carcinoma. <i>Journal of Clinical and Translational Hepatology</i> , 2016, 1, 138-43.	1.4	10
65	Identifying microRNA panels specifically associated with hepatocellular carcinoma and its different etiologies. <i>Hepatoma Research</i> , 2016, 2, 151.	1.5	12
66	Evaluating normalization approaches for the better identification of aberrant microRNAs associated with hepatocellular carcinoma. <i>Hepatoma Research</i> , 2016, 2, 305-315.	1.5	13
67	Mismatch Repair Polymorphisms as Markers of Breast Cancer Prevalence in the Breast Cancer Family Registry. <i>Anticancer Research</i> , 2016, 36, 4437-4442.	1.1	25
68	Plasma Adiponectin and Hepatocellular Carcinoma Survival Among Patients Without Liver Transplantation. <i>Anticancer Research</i> , 2016, 36, 5307-5314.	1.1	12
69	Association between polymorphisms of <i>APE1</i> and <i>OGG1</i> and risk of colorectal cancer in Taiwan. <i>World Journal of Gastroenterology</i> , 2016, 22, 3372-3380.	3.4	17
70	Blood DNA methylation markers in prospectively identified hepatocellular carcinoma cases and controls from Taiwan. <i>World Journal of Hepatology</i> , 2016, 8, 301.	2.0	6
71	Variation in genes involved in the immune response and prostate cancer risk in the placebo arm of the Prostate Cancer Prevention Trial. <i>Prostate</i> , 2015, 75, 1403-1418.	2.3	25
72	Promoter Hypermethylation in White Blood Cell DNA and Breast Cancer Risk. <i>Journal of Cancer</i> , 2015, 6, 819-824.	2.6	29

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73	Genome-Wide Expression of MicroRNAs Is Regulated by DNA Methylation in Hepatocarcinogenesis. <i>Gastroenterology Research and Practice</i> , 2015, 2015, 1-12.	1.5	20
74	Global DNA Methylation, Measured by the Luminometric Methylation Assay (LUMA), Associates with Postmenopausal Breast Cancer in Non-Obese and Physically Active Women. <i>Journal of Cancer</i> , 2015, 6, 548-554.	2.6	7
75	Polycyclic aromatic hydrocarbon (PAH)â€“DNA adducts and breast cancer: modification by gene promoter methylation in a population-based study. <i>Cancer Causes and Control</i> , 2015, 26, 1791-1802.	1.8	22
76	Dietary Modifications, Weight Loss, and Changes in Metabolic Markers Affect Global DNA Methylation in Hispanic, African American, and Afro-Caribbean Breast Cancer Survivors,. <i>Journal of Nutrition</i> , 2015, 145, 783-790.	2.7	61
77	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , 2015, 47, 373-380.	20.4	528
78	Integrative epigenomic and genomic filtering for methylation markers in hepatocellular carcinomas. <i>BMC Medical Genomics</i> , 2015, 8, 28.	1.5	27
79	Effect of Finasteride on Serum Androstenedione and Risk of Prostate Cancer Within the Prostate Cancer Prevention Trial: Differential Effect on High- and Low-grade Disease. <i>Urology</i> , 2015, 85, 616-620.	1.4	8
80	Genetic polymorphisms in DNA repair and oxidative stress pathways may modify the association between body size and postmenopausal breast cancer. <i>Annals of Epidemiology</i> , 2015, 25, 263-269.	2.1	9
81	Dietary intake of fish, polyunsaturated fatty acids, and survival after breast cancer: A populationâ€“based followâ€“up study on Long Island, New York. <i>Cancer</i> , 2015, 121, 2244-2252.	4.1	30
82	Polyunsaturated fatty acid interactions and breast cancer incidence: a population-based case-control study on Long Island, New York. <i>Annals of Epidemiology</i> , 2015, 25, 929-935.	2.1	29
83	Vitamin D-related gene polymorphisms, plasma 25-hydroxyvitamin D, and breast cancer risk. <i>Cancer Causes and Control</i> , 2015, 26, 187-203.	1.8	61
84	Gene-Specific Promoter Methylation Status in Hormone-Receptor-Positive Breast Cancer Associates with Postmenopausal Body Size and Recreational Physical Activity. <i>International Journal of Cancer and Clinical Research</i> , 2015, 2, .	0.1	12
85	Exploration of Deregulated Long Non-Coding RNAs in Association with Hepatocarcinogenesis and Survival. <i>Cancers</i> , 2015, 7, 1847-1862.	3.8	16
86	Genome-Wide Methylation Analyses in Glioblastoma Multiforme. <i>PLoS ONE</i> , 2014, 9, e89376.	2.5	45
87	Differences in DNA methylation by extent of breast cancer family history in unaffected women. <i>Epigenetics</i> , 2014, 9, 243-248.	2.9	23
88	Correlations in global DNA methylation measures in peripheral blood mononuclear cells and granulocytes. <i>Epigenetics</i> , 2014, 9, 1504-1510.	2.9	15
89	ALS Multicenter Cohort Study of Oxidative Stress (ALS COSMOS): Study methodology, recruitment, and baseline demographic and disease characteristics. <i>Amyotrophic Lateral Sclerosis and Frontotemporal Degeneration</i> , 2014, 15, 192-203.	2.2	35
90	Correlation of DNA methylation levels in blood and saliva DNA in young girls of the LEGACY Girls study. <i>Epigenetics</i> , 2014, 9, 929-933.	2.9	33

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91	Polymorphisms in DNA repair genes, recreational physical activity and breast cancer risk. <i>International Journal of Cancer</i> , 2014, 134, 654-663.	5.4	25
92	Genetic variation in multiple biologic pathways, flavonoid intake, and breast cancer. <i>Cancer Causes and Control</i> , 2014, 25, 215-226.	1.8	10
93	A Genome-wide Association Study of Early-Onset Breast Cancer Identifies <i>PFKM</i> as a Novel Breast Cancer Gene and Supports a Common Genetic Spectrum for Breast Cancer at Any Age. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 658-669.	1.9	79
94	Clinical perspective on oxidative stress in sporadic amyotrophic lateral sclerosis. <i>Free Radical Biology and Medicine</i> , 2013, 65, 509-527.	4.5	278
95	DNA double-strand break repair genotype and phenotype and breast cancer risk within sisters from the New York site of the Breast Cancer Family Registry (BCFR). <i>Cancer Causes and Control</i> , 2013, 24, 2157-2168.	1.8	14
96	Human and Methodological Sources of Variability in the Measurement of Urinary 8-Oxo-7,8-dihydro-2-deoxyguanosine. <i>Antioxidants and Redox Signaling</i> , 2013, 18, 2377-2391.	5.5	135
97	Exploring genome-wide DNA methylation profiles altered in hepatocellular carcinoma using Infinium HumanMethylation 450 BeadChips. <i>Epigenetics</i> , 2013, 8, 34-43.	2.9	145
98	Exploration of Genome-Wide Circulating MicroRNA in Hepatocellular Carcinoma: MiR-483-5p as a Potential Biomarker. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 2364-2373.	1.9	100
99	Early life socioeconomic factors and genomic DNA methylation in mid-life. <i>Epigenetics</i> , 2013, 8, 23-27.	2.9	78
100	Global DNA methylation in a population with aflatoxin B ₁ exposure. <i>Epigenetics</i> , 2013, 8, 962-969.	2.9	30
101	Genome-Wide and Differential Proteomic Analysis of Hepatitis B Virus and Aflatoxin B1 Related Hepatocellular Carcinoma in Guangxi, China. <i>PLoS ONE</i> , 2013, 8, e83465.	2.5	28
102	Phase IB Randomized, Double-Blinded, Placebo-Controlled, Dose Escalation Study of Polyphenon E in Women with Hormone Receptor-Negative Breast Cancer. <i>Cancer Prevention Research</i> , 2012, 5, 1144-1154.	1.6	86
103	Global DNA methylation levels in white blood cells as a biomarker for hepatocellular carcinoma risk: a nested case-control study. <i>Carcinogenesis</i> , 2012, 33, 1340-1345.	2.8	39
104	Genomic Methylation Changes Over Time in Peripheral Blood Mononuclear Cell DNA: Differences by Assay Type and Baseline Values. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 1314-1318.	1.9	24
105	Global DNA methylation levels in white blood cell DNA from sisters discordant for breast cancer from the New York site of the Breast Cancer Family Registry. <i>Epigenetics</i> , 2012, 7, 868-874.	2.9	40
106	Genome-wide aberrant DNA methylation of microRNA host genes in hepatocellular carcinoma. <i>Epigenetics</i> , 2012, 7, 1230-1237.	2.9	60
107	Genetic Variation in GPX1 Is Associated with GPX1 Activity in a Comprehensive Analysis of Genetic Variations in Selenoenzyme Genes and Their Activity and Oxidative Stress in Humans. <i>Journal of Nutrition</i> , 2012, 142, 419-426.	2.7	24
108	Adult global DNA methylation in relation to pre-natal nutrition. <i>International Journal of Epidemiology</i> , 2012, 41, 116-123.	2.0	65

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109	Repetitive element DNA methylation levels in white blood cell DNA from sisters discordant for breast cancer from the New York site of the Breast Cancer Family Registry. <i>Carcinogenesis</i> , 2012, 33, 1946-1952.	2.8	67
110	Polymorphisms in oxidative stress genes, physical activity, and breast cancer risk. <i>Cancer Causes and Control</i> , 2012, 23, 1949-1958.	1.8	20
111	White blood cell global methylation and IL-6 promoter methylation in association with diet and lifestyle risk factors in a cancer-free population. <i>Epigenetics</i> , 2012, 7, 606-614.	2.9	80
112	Selenium, Selenoenzymes, Oxidative Stress and Risk of Neoplastic Progression from Barrett's Esophagus: Results from Biomarkers and Genetic Variants. <i>PLoS ONE</i> , 2012, 7, e38612.	2.5	28
113	Dysregulation of circulating microRNAs and prediction of aggressive prostate cancer. <i>Prostate</i> , 2012, 72, 1469-1477.	2.3	168
114	Genome-wide DNA methylation profiles in hepatocellular carcinoma. <i>Hepatology</i> , 2012, 55, 1799-1808.	8.1	181
115	DNA methylation in peripheral blood measured by LUMA is associated with breast cancer in a population-based study. <i>FASEB Journal</i> , 2012, 26, 2657-2666.	0.5	77
116	Genetic polymorphisms in telomere pathway genes, telomere length, and breast cancer survival. <i>Breast Cancer Research and Treatment</i> , 2012, 134, 393-400.	2.5	38
117	Biologic and epigenetic impact of commuting to work by car or using public transportation: A case-control study. <i>Preventive Medicine</i> , 2012, 54, 229-233.	3.5	23
118	Prognostic significance of gene-specific promoter hypermethylation in breast cancer patients. <i>Breast Cancer Research and Treatment</i> , 2012, 131, 197-205.	2.5	79
119	Topical application of green tea polyphenol (EGCG) epigallocatechin-3-gallate for prevention of recurrent oral neoplastic lesions. <i>Journal of Orofacial Sciences</i> , 2012, 4, 43.	0.2	15
120	Global hypomethylation in hepatocellular carcinoma and its relationship to aflatoxin B ₁ exposure. <i>World Journal of Hepatology</i> , 2012, 4, 169.	2.0	28
121	DNA methylation in white blood cells. <i>Epigenetics</i> , 2011, 6, 828-837.	2.9	306
122	Significant differences in global genomic DNA methylation by gender and race/ethnicity in peripheral blood. <i>Epigenetics</i> , 2011, 6, 623-629.	2.9	342
123	The influence of one-carbon metabolism on gene promoter methylation in a population-based breast cancer study. <i>Epigenetics</i> , 2011, 6, 1276-1283.	2.9	20
124	Serum estrogen levels and prostate cancer risk in the prostate cancer prevention trial: a nested case-control study. <i>Cancer Causes and Control</i> , 2011, 22, 1121-1131.	1.8	43
125	The Definition of Life: A Survey of Obstetricians and Neonatologists in New York City Hospitals Regarding Extremely Premature Births. <i>Maternal and Child Health Journal</i> , 2011, 15, 446-452.	1.5	17
126	Global DNA methylation levels in girls with and without a family history of breast cancer. <i>Epigenetics</i> , 2011, 6, 29-33.	2.9	31

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127	Physical activity and global genomic DNA methylation in a cancer-free population. <i>Epigenetics</i> , 2011, 6, 293-299.	2.9	161
128	Global methylation profiles in DNA from different blood cell types. <i>Epigenetics</i> , 2011, 6, 76-85.	2.9	129
129	Prenatal Smoke Exposure and Genomic DNA Methylation in a Multiethnic Birth Cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 2518-2523.	1.9	94
130	Dietary Patterns Are Associated with Levels of Global Genomic DNA Methylation in a Cancer-Free Population. <i>Journal of Nutrition</i> , 2011, 141, 1165-1171.	2.7	104
131	Replication and Functional Genomic Analyses of the Breast Cancer Susceptibility Locus at 6q25.1 Generalize Its Importance in Women of Chinese, Japanese, and European Ancestry. <i>Cancer Research</i> , 2011, 71, 1344-1355.	0.9	72
132	Repeat polymorphisms in estrogen metabolism genes and prostate cancer risk: results from the Prostate Cancer Prevention Trial. <i>Carcinogenesis</i> , 2011, 32, 1500-1506.	2.8	23
133	Immunologic Detection of Benzo(a)pyreneâ€DNA Adducts. <i>Methods in Molecular Biology</i> , 2011, 682, 271-278.	0.0	5
134	Common genetic variations in the LEP and LEPR genes, obesity and breast cancer incidence and survival. <i>Breast Cancer Research and Treatment</i> , 2010, 120, 745-752.	2.5	47
135	Gene promoter methylation is associated with increased mortality among women with breast cancer. <i>Breast Cancer Research and Treatment</i> , 2010, 121, 685-692.	2.5	41
136	Serum Oxidized Protein and Prostate Cancer Risk within the Prostate Cancer Prevention Trial. <i>Cancer Prevention Research</i> , 2010, 3, 478-483.	1.6	12
137	Transition of a Clinical Trial into Translational Research: The Prostate Cancer Prevention Trial Experience. <i>Cancer Prevention Research</i> , 2010, 3, 1523-1533.	1.6	19
138	Airborne particulate metals in the New York City subway: A pilot study to assess the potential for health impacts. <i>Environmental Research</i> , 2010, 110, 1-11.	7.7	76
139	Multiple Genetic Variants in Telomere Pathway Genes and Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 219-228.	1.9	47
140	High intakes of choline and betaine reduce breast cancer mortality in a populationâ€based study. <i>FASEB Journal</i> , 2009, 23, 4022-4028.	0.5	86
141	Effect of Selenium and Vitamin E on Risk of Prostate Cancer and Other Cancers. <i>JAMA - Journal of the American Medical Association</i> , 2009, 301, 39.	7.0	1,856
142	PAHâ€DNA Adducts, Cigarette Smoking, <i>GST</i> Polymorphisms, and Breast Cancer Risk. <i>Environmental Health Perspectives</i> , 2009, 117, 552-558.	8.2	55
143	Aberrant Methylation of RASSF1A in Plasma DNA Before Breast Cancer Diagnosis in the Breast Cancer Family Registry. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 2723-2725.	1.9	27
144	Association between Plasma 25-Hydroxyvitamin D and Breast Cancer Risk. <i>Cancer Prevention Research</i> , 2009, 2, 598-604.	1.6	115

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145	Mutations in <i>p53</i> , p53 protein overexpression and breast cancer survival. <i>Journal of Cellular and Molecular Medicine</i> , 2009, 13, 3847-3857.	3.6	39
146	HINT1 inhibits β -catenin/TCF4, USF2 and NF κ B activity in human hepatoma cells. <i>International Journal of Cancer</i> , 2009, 124, 1526-1534.	5.4	42
147	Telomere length, oxidative damage, antioxidants and breast cancer risk. <i>International Journal of Cancer</i> , 2009, 124, 1637-1643.	5.4	142
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