

Regina M Santella

List of Publications by Year in descending order

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Version: 2024-02-01

327
papers

17,066
citations

15880

67
h-index

26792

111
g-index

328
all docs

328
docs citations

328
times ranked

21986
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.	1.1	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.	0.8	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.	1.1	5
4	Predictors of urinary polycyclic aromatic hydrocarbon metabolites in girls from the San Francisco Bay Area. <i>Environmental Research</i> , 2022, 205, 112534.	3.7	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.	1.2	4
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.	1.2	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.	1.3	8
8	Common Childhood Viruses and Pubertal Timing: The LEGACY Girls Study. <i>American Journal of Epidemiology</i> , 2021, 190, 766-778.	1.6	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.	1.1	8
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.	0.6	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.	1.3	7
12	Reproductive and environmental exposures and the breast cancer risk in Taiwanese women. <i>Scientific Reports</i> , 2021, 11, 15656.	1.6	6
13	Urinary parabens and breast cancer risk: Modification and interaction by LINE-1/LUMA methylation in the Long Island Breast Cancer Study Project. <i>ISEE Conference Abstracts</i> , 2021, 2021, .	0.0	0
14	Associations of Phthalates and Phenols, Telomere Length, and Breast Cancer in the Long Island Breast Cancer Study Project. <i>ISEE Conference Abstracts</i> , 2021, 2021, .	0.0	0
15	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.	1.3	4
16	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.	1.3	13
17	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.	3.0	1
18	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.	1.5	5

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19	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.	2.2	8
20	Urinary Estrogen Metabolites and Long-Term Mortality Following Breast Cancer. <i>JNCI Cancer Spectrum</i> , 2020, 4, pkaa014.	1.4	0
21	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.	0.9	27
22	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.	1.6	19
23	Plasma creatinine and oxidative stress biomarkers in amyotrophic lateral sclerosis. <i>Amyotrophic Lateral Sclerosis and Frontotemporal Degeneration</i> , 2020, 21, 263-272.	1.1	20
24	High levels of global genome methylation in patients with retinoblastoma. <i>Oncology Letters</i> , 2020, 20, 715-723.	0.8	8
25	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.	1.3	13
26	Prediagnosis aspirin use, DNA methylation, and mortality after breast cancer: A population-based study. <i>Cancer</i> , 2019, 125, 3836-3844.	2.0	13
27	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.	1.1	16
28	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.	1.3	6
29	Reproductive characteristics are associated with gene-specific promoter methylation status in breast cancer. <i>BMC Cancer</i> , 2019, 19, 926.	1.1	4
30	Urinary concentrations of environmental phenols and their associations with breast cancer incidence and mortality following breast cancer. <i>Environment International</i> , 2019, 130, 104890.	4.8	66
31	Assessment of arsenic and polycyclic aromatic hydrocarbon (PAH) exposures on immune function among males in Bangladesh. <i>PLoS ONE</i> , 2019, 14, e0216662.	1.1	24
32	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.	2.8	7
33	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.	1.1	5
34	Rasagiline for amyotrophic lateral sclerosis: A randomized, controlled trial. <i>Muscle and Nerve</i> , 2019, 59, 201-207.	1.0	35
35	Tumor expression of environmental chemical-responsive genes and breast cancer mortality. <i>Endocrine-Related Cancer</i> , 2019, 26, 843-851.	1.6	18
36	Title is missing!. , 2019, 14, e0220451.		0

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37	Title is missing!. , 2019, 14, e0220451.		0
38	Title is missing!. , 2019, 14, e0220451.		0
39	Title is missing!. , 2019, 14, e0220451.		0
40	Breast cancer family history and allele-specific DNA methylation in the legacy girls study. <i>Epigenetics</i> , 2018, 13, 240-250.	1.3	10
41	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.	3.0	32
42	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.	1.3	14
43	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.	1.3	56
44	Maternal cigarette smoking during pregnancy and offspring DNA methylation in midlife. <i>Epigenetics</i> , 2018, 13, 129-134.	1.3	61
45	Novel Predictors of Breast Cancer Survival Derived from miRNA Activity Analysis. <i>Clinical Cancer Research</i> , 2018, 24, 581-591.	3.2	21
46	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.	2.8	36
47	Response to H. Nabi et al.. <i>Journal of the National Cancer Institute</i> , 2018, 110, 1424-1425.	3.0	0
48	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.	1.9	4
49	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.	2.2	18
50	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.	2.3	86
51	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.	1.3	21
52	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.	0.7	19
53	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.	2.9	54
54	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.	1.2	21

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55	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.	0.9	15
56	Plasma DNA methylation marker and hepatocellular carcinoma risk prediction model for the general population. <i>Carcinogenesis</i> , 2017, 38, 1021-1028.	1.3	37
57	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.	3.7	24
58	Environmental Tobacco Smoke Exposure and Survival Following Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 278-280.	1.1	2
59	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.	0.5	6
60	DNA Methylation in Breast Tumor from High-risk Women in the Breast Cancer Family Registry. <i>Anticancer Research</i> , 2017, 37, 659-664.	0.5	7
61	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.	2.8	73
62	Genetically Predicted Body Mass Index and Breast Cancer Risk: Mendelian Randomization Analyses of Data from 145,000 Women of European Descent. <i>PLoS Medicine</i> , 2016, 13, e1002105.	3.9	118
63	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.	1.2	5
64	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.	2.3	28
65	Association of genetic susceptibility variants for type 2 diabetes with breast cancer risk in women of European ancestry. <i>Cancer Causes and Control</i> , 2016, 27, 679-693.	0.8	21
66	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.	1.1	33
67	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016, 7, 11375.	5.8	93
68	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.	1.3	24
69	NEpiC: a network-assisted algorithm for epigenetic studies using mean and variance combined signals. <i>Nucleic Acids Research</i> , 2016, 44, e134-e134.	6.5	13
70	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.	3.7	52
71	Exposure to multiple sources of polycyclic aromatic hydrocarbons and breast cancer incidence. <i>Environment International</i> , 2016, 89-90, 185-192.	4.8	122
72	DNA methylation modifies the association between obesity and survival after breast cancer diagnosis. <i>Breast Cancer Research and Treatment</i> , 2016, 156, 183-194.	1.1	17

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73	Environmental Exposures and Hepatocellular Carcinoma. <i>Journal of Clinical and Translational Hepatology</i> , 2016, 1, 138-43.	0.7	10
74	Identifying microRNA panels specifically associated with hepatocellular carcinoma and its different etiologies. <i>Hepatoma Research</i> , 2016, 2, 151.	0.6	12
75	Evaluating normalization approaches for the better identification of aberrant microRNAs associated with hepatocellular carcinoma. <i>Hepatoma Research</i> , 2016, 2, 305-315.	0.6	13
76	Mismatch Repair Polymorphisms as Markers of Breast Cancer Prevalence in the Breast Cancer Family Registry. <i>Anticancer Research</i> , 2016, 36, 4437-4442.	0.5	24
77	Plasma Adiponectin and Hepatocellular Carcinoma Survival Among Patients Without Liver Transplantation. <i>Anticancer Research</i> , 2016, 36, 5307-5314.	0.5	12
78	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.	1.4	17
79	Blood DNA methylation markers in prospectively identified hepatocellular carcinoma cases and controls from Taiwan. <i>World Journal of Hepatology</i> , 2016, 8, 301.	0.8	6
80	DNA Repair Gene Expression Levels as Indicators of Breast Cancer in the Breast Cancer Family Registry. <i>Anticancer Research</i> , 2016, 36, 4039-44.	0.5	8
81	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.	1.2	25
82	Promoter Hypermethylation in White Blood Cell DNA and Breast Cancer Risk. <i>Journal of Cancer</i> , 2015, 6, 819-824.	1.2	28
83	Genome-Wide Expression of MicroRNAs Is Regulated by DNA Methylation in Hepatocarcinogenesis. <i>Gastroenterology Research and Practice</i> , 2015, 2015, 1-12.	0.7	20
84	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.	1.2	7
85	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.	0.8	22
86	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.	1.3	59
87	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.	9.4	513
88	Integrative epigenomic and genomic filtering for methylation markers in hepatocellular carcinomas. <i>BMC Medical Genomics</i> , 2015, 8, 28.	0.7	24
89	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.	0.5	8
90	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.	0.9	8

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91	Dietary intake of fish, polyunsaturated fatty acids, and survival after breast cancer: A population-based follow-up study on Long Island, New York. <i>Cancer</i> , 2015, 121, 2244-2252.	2.0	28
92	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.	0.9	26
93	Vitamin D-related gene polymorphisms, plasma 25-hydroxyvitamin D, and breast cancer risk. <i>Cancer Causes and Control</i> , 2015, 26, 187-203.	0.8	60
94	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
95	Exploration of Deregulated Long Non-Coding RNAs in Association with Hepatocarcinogenesis and Survival. <i>Cancers</i> , 2015, 7, 1847-1862.	1.7	16
96	microRNA Expression in Prospectively Collected Blood as a Potential Biomarker of Breast Cancer Risk in the BCFR. <i>Anticancer Research</i> , 2015, 35, 3969-77.	0.5	26
97	Genome-Wide Methylation Analyses in Glioblastoma Multiforme. <i>PLoS ONE</i> , 2014, 9, e89376.	1.1	45
98	Differences in DNA methylation by extent of breast cancer family history in unaffected women. <i>Epigenetics</i> , 2014, 9, 243-248.	1.3	23
99	Correlations in global DNA methylation measures in peripheral blood mononuclear cells and granulocytes. <i>Epigenetics</i> , 2014, 9, 1504-1510.	1.3	15
100	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.	1.1	35
101	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.	1.3	32
102	Polymorphisms in DNA repair genes, recreational physical activity and breast cancer risk. <i>International Journal of Cancer</i> , 2014, 134, 654-663.	2.3	24
103	Genetic variation in multiple biologic pathways, flavonoid intake, and breast cancer. <i>Cancer Causes and Control</i> , 2014, 25, 215-226.	0.8	10
104	A Genome-wide Association Study of Early-Onset Breast Cancer Identifies <i>PCSK9</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.	1.1	77
105	Prognostic value of miR-375 and miR-214-3p in early stage oral squamous cell carcinoma. <i>American Journal of Translational Research (discontinued)</i> , 2014, 6, 580-92.	0.0	24
106	Clinical perspective on oxidative stress in sporadic amyotrophic lateral sclerosis. <i>Free Radical Biology and Medicine</i> , 2013, 65, 509-527.	1.3	269
107	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.	0.8	14
108	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.	2.5	130

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109	Exploring genome-wide DNA methylation profiles altered in hepatocellular carcinoma using Infinium HumanMethylation 450 BeadChips. <i>Epigenetics</i> , 2013, 8, 34-43.	1.3	144
110	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.	1.1	97
111	Early life socioeconomic factors and genomic DNA methylation in mid-life. <i>Epigenetics</i> , 2013, 8, 23-27.	1.3	76
112	Global DNA methylation in a population with aflatoxin B ₁ exposure. <i>Epigenetics</i> , 2013, 8, 962-969.	1.3	30
113	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.	1.1	27
114	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.	0.7	86
115	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.	1.3	39
116	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.	1.1	24
117	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.	1.3	40
118	Genome-wide aberrant DNA methylation of microRNA host genes in hepatocellular carcinoma. <i>Epigenetics</i> , 2012, 7, 1230-1237.	1.3	59
119	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.	1.3	23
120	Adult global DNA methylation in relation to pre-natal nutrition. <i>International Journal of Epidemiology</i> , 2012, 41, 116-123.	0.9	64
121	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.	1.3	66
122	Polymorphisms in oxidative stress genes, physical activity, and breast cancer risk. <i>Cancer Causes and Control</i> , 2012, 23, 1949-1958.	0.8	18
123	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.	1.3	80
124	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.	1.1	28
125	Dysregulation of circulating microRNAs and prediction of aggressive prostate cancer. <i>Prostate</i> , 2012, 72, 1469-1477.	1.2	167
126	Genome-wide DNA methylation profiles in hepatocellular carcinoma. <i>Hepatology</i> , 2012, 55, 1799-1808.	3.6	178

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127	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.	0.2	76
128	Genetic polymorphisms in telomere pathway genes, telomere length, and breast cancer survival. <i>Breast Cancer Research and Treatment</i> , 2012, 134, 393-400.	1.1	38
129	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.	1.6	22
130	Prognostic significance of gene-specific promoter hypermethylation in breast cancer patients. <i>Breast Cancer Research and Treatment</i> , 2012, 131, 197-205.	1.1	78
131	Topical application of green tea polyphenol (âˆ™) epigallocatechin-3-gallate for prevention of recurrent oral neoplastic lesions. <i>Journal of Orofacial Sciences</i> , 2012, 4, 43.	0.1	15
132	Global hypomethylation in hepatocellular carcinoma and its relationship to aflatoxin B ₁ exposure. <i>World Journal of Hepatology</i> , 2012, 4, 169.	0.8	28
133	Serum adiponectin and overall survival in a prospective cohort of patients with hepatocellular carcinoma.. <i>Journal of Clinical Oncology</i> , 2012, 30, 205-205.	0.8	0
134	DNA methylation in white blood cells. <i>Epigenetics</i> , 2011, 6, 828-837.	1.3	304
135	Significant differences in global genomic DNA methylation by gender and race/ethnicity in peripheral blood. <i>Epigenetics</i> , 2011, 6, 623-629.	1.3	331
136	The influence of one-carbon metabolism on gene promoter methylation in a population-based breast cancer study. <i>Epigenetics</i> , 2011, 6, 1276-1283.	1.3	20
137	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.	0.8	42
138	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.	0.7	16
139	Global DNA methylation levels in girls with and without a family history of breast cancer. <i>Epigenetics</i> , 2011, 6, 29-33.	1.3	31
140	Physical activity and global genomic DNA methylation in a cancer-free population. <i>Epigenetics</i> , 2011, 6, 293-299.	1.3	154
141	Global methylation profiles in DNA from different blood cell types. <i>Epigenetics</i> , 2011, 6, 76-85.	1.3	128
142	Prenatal Smoke Exposure and Genomic DNA Methylation in a Multiethnic Birth Cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 2518-2523.	1.1	94
143	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.	1.3	101
144	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.	0.4	71

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145	Repeat polymorphisms in estrogen metabolism genes and prostate cancer risk: results from the Prostate Cancer Prevention Trial. <i>Carcinogenesis</i> , 2011, 32, 1500-1506.	1.3	23
146	Immunologic Detection of Benzo(a)pyreneâ€“DNA Adducts. <i>Methods in Molecular Biology</i> , 2011, 682, 271-278.	0.4	5
147	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.	1.1	47
148	Gene promoter methylation is associated with increased mortality among women with breast cancer. <i>Breast Cancer Research and Treatment</i> , 2010, 121, 685-692.	1.1	41
149	Serum Oxidized Protein and Prostate Cancer Risk within the Prostate Cancer Prevention Trial. <i>Cancer Prevention Research</i> , 2010, 3, 478-483.	0.7	12
150	Transition of a Clinical Trial into Translational Research: The Prostate Cancer Prevention Trial Experience. <i>Cancer Prevention Research</i> , 2010, 3, 1523-1533.	0.7	19
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