

Laura Baglietto

List of Publications by Year in descending order

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Version: 2024-02-01

194
papers

15,994
citations

15495

65
h-index

19169

118
g-index

200
all docs

200
docs citations

200
times ranked

20707
citing authors

#	ARTICLE	IF	CITATIONS
1	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	9.4	960
2	Subtyping of Breast Cancer by Immunohistochemistry to Investigate a Relationship between Subtype and Short and Long Term Survival: A Collaborative Analysis of Data for 10,159 Cases from 12 Studies. <i>PLoS Medicine</i> , 2010, 7, e1000279.	3.9	764
3	Associations of Breast Cancer Risk Factors With Tumor Subtypes: A Pooled Analysis From the Breast Cancer Association Consortium Studies. <i>Journal of the National Cancer Institute</i> , 2011, 103, 250-263.	3.0	596
4	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 371-384.	9.4	493
5	The Clinical Phenotype of Lynch Syndrome Due to Germ-Line PMS2 Mutations. <i>Gastroenterology</i> , 2008, 135, 419-428.e1.	0.6	480
6	Newly discovered breast cancer susceptibility loci on 3p24 and 17q23.2. <i>Nature Genetics</i> , 2009, 41, 585-590.	9.4	434
7	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , 2013, 45, 392-398.	9.4	374
8	Risks of Lynch Syndrome Cancers for MSH6 Mutation Carriers. <i>Journal of the National Cancer Institute</i> , 2010, 102, 193-201.	3.0	328
9	GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 362-370.	9.4	326
10	Heterogeneity of Breast Cancer Associations with Five Susceptibility Loci by Clinical and Pathological Characteristics. <i>PLoS Genetics</i> , 2008, 4, e1000054.	1.5	315
11	A locus on 19p13 modifies risk of breast cancer in BRCA1 mutation carriers and is associated with hormone receptor-negative breast cancer in the general population. <i>Nature Genetics</i> , 2010, 42, 885-892.	9.4	309
12	Breast Cancer Risk From Modifiable and Nonmodifiable Risk Factors Among White Women in the United States. <i>JAMA Oncology</i> , 2016, 2, 1295.	3.4	285
13	A common variant at the TERT-CLPTM1L locus is associated with estrogen receptor-negative breast cancer. <i>Nature Genetics</i> , 2011, 43, 1210-1214.	9.4	279
14	A genome-wide association study identifies a new ovarian cancer susceptibility locus on 9p22.2. <i>Nature Genetics</i> , 2009, 41, 996-1000.	9.4	276
15	Mammographic Density Phenotypes and Risk of Breast Cancer: A Meta-analysis. <i>Journal of the National Cancer Institute</i> , 2014, 106, .	3.0	261
16	Genome-wide association analysis identifies three new breast cancer susceptibility loci. <i>Nature Genetics</i> , 2012, 44, 312-318.	9.4	256
17	Inositol polyphosphate 4-phosphatase II regulates PI3K/Akt signaling and is lost in human basal-like breast cancers. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 22231-22236.	3.3	249
18	Association of Body Mass Index and Age With Subsequent Breast Cancer Risk in Premenopausal Women. <i>JAMA Oncology</i> , 2018, 4, e181771.	3.4	210

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19	Functional Variants at the 11q13 Risk Locus for Breast Cancer Regulate Cyclin D1 Expression through Long-Range Enhancers. <i>American Journal of Human Genetics</i> , 2013, 92, 489-503.	2.6	201
20	Hypomethylation of smoking-related genes is associated with future lung cancer in four prospective cohorts. <i>Nature Communications</i> , 2015, 6, 10192.	5.8	197
21	Social adversity and epigenetic aging: a multi-cohort study on socioeconomic differences in peripheral blood DNA methylation. <i>Scientific Reports</i> , 2017, 7, 16266.	1.6	181
22	A meta-analysis of genome-wide association studies of breast cancer identifies two novel susceptibility loci at 6q14 and 20q11. <i>Human Molecular Genetics</i> , 2012, 21, 5373-5384.	1.4	168
23	Fruit and Vegetable Intake and Risk of Breast Cancer by Hormone Receptor Status. <i>Journal of the National Cancer Institute</i> , 2013, 105, 219-236.	3.0	164
24	<i>CHEK2</i> 1100delC Heterozygosity in Women With Breast Cancer Associated With Early Death, Breast Cancer-Specific Death, and Increased Risk of a Second Breast Cancer. <i>Journal of Clinical Oncology</i> , 2012, 30, 4308-4316.	0.8	162
25	A metabolomic study of biomarkers of meat and fish intake. <i>American Journal of Clinical Nutrition</i> , 2017, 105, 600-608.	2.2	156
26	DNA methylome analysis identifies accelerated epigenetic ageing associated with postmenopausal breast cancer susceptibility. <i>European Journal of Cancer</i> , 2017, 75, 299-307.	1.3	154
27	Low penetrance breast cancer susceptibility loci are associated with specific breast tumor subtypes: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2011, 20, 3289-3303.	1.4	152
28	Cancer Risks For Mismatch Repair Gene Mutation Carriers: A Population-Based Early Onset Case-Family Study. <i>Clinical Gastroenterology and Hepatology</i> , 2006, 4, 489-498.	2.4	151
29	Genome-wide association study identifies 25 known breast cancer susceptibility loci as risk factors for triple-negative breast cancer. <i>Carcinogenesis</i> , 2014, 35, 1012-1019.	1.3	145
30	Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. <i>Nature Communications</i> , 2013, 4, 1628.	5.8	144
31	Evidence of Gene-Environment Interactions between Common Breast Cancer Susceptibility Loci and Established Environmental Risk Factors. <i>PLoS Genetics</i> , 2013, 9, e1003284.	1.5	136
32	Breast Cancer Risk After Recent Childbirth. <i>Annals of Internal Medicine</i> , 2019, 170, 22.	2.0	120
33	A review of the reporting and handling of missing data in cohort studies with repeated assessment of exposure measures. <i>BMC Medical Research Methodology</i> , 2012, 12, 96.	1.4	119
34	DNA methylation changes measured in pre-diagnostic peripheral blood samples are associated with smoking and lung cancer risk. <i>International Journal of Cancer</i> , 2017, 140, 50-61.	2.3	115
35	Circulating steroid hormone concentrations in postmenopausal women in relation to body size and composition. <i>Breast Cancer Research and Treatment</i> , 2009, 115, 171-179.	1.1	113
36	Common Breast Cancer Susceptibility Loci Are Associated with Triple-Negative Breast Cancer. <i>Cancer Research</i> , 2011, 71, 6240-6249.	0.4	109

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37	Genome-wide association study identifies multiple loci associated with both mammographic density and breast cancer risk. <i>Nature Communications</i> , 2014, 5, 5303.	5.8	109
38	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. <i>Nature Communications</i> , 2017, 8, 15724.	5.8	106
39	Association of DNA Methylation-Based Biological Age With Health Risk Factors and Overall and Cause-Specific Mortality. <i>American Journal of Epidemiology</i> , 2018, 187, 529-538.	1.6	106
40	Long-Term Exposure to Ambient Air Pollution and Incidence of Postmenopausal Breast Cancer in 15 European Cohorts within the ESCAPE Project. <i>Environmental Health Perspectives</i> , 2017, 125, 107005.	2.8	104
41	A PALB2 mutation associated with high risk of breast cancer. <i>Breast Cancer Research</i> , 2010, 12, R109.	2.2	102
42	Common Breast Cancer Susceptibility Variants in <i>LSP1</i> and <i>RAD51L1</i> Are Associated with Mammographic Density Measures that Predict Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 1156-1166.	1.1	101
43	Alcohol consumption and breast cancer risk by estrogen receptor status: in a pooled analysis of 20 studies. <i>International Journal of Epidemiology</i> , 2016, 45, 916-928.	0.9	101
44	19p13.1 Is a Triple-Negative-Specific Breast Cancer Susceptibility Locus. <i>Cancer Research</i> , 2012, 72, 1795-1803.	0.4	100
45	Cross-Cancer Genome-Wide Analysis of Lung, Ovary, Breast, Prostate, and Colorectal Cancer Reveals Novel Pleiotropic Associations. <i>Cancer Research</i> , 2016, 76, 5103-5114.	0.4	100
46	Risk of Estrogen Receptor-Positive and -Negative Breast Cancer and Single-Nucleotide Polymorphism 2q35-rs13387042. <i>Journal of the National Cancer Institute</i> , 2009, 101, 1012-1018.	3.0	99
47	Fine-Scale Mapping of the FGFR2 Breast Cancer Risk Locus: Putative Functional Variants Differentially Bind FOXA1 and E2F1. <i>American Journal of Human Genetics</i> , 2013, 93, 1046-1060.	2.6	98
48	Identification and molecular characterization of a new ovarian cancer susceptibility locus at 17q21.31. <i>Nature Communications</i> , 2013, 4, 1627.	5.8	98
49	Refined histopathological predictors of BRCA1 and BRCA2 mutation status: a large-scale analysis of breast cancer characteristics from the BCAC, CIMBA, and ENIGMA consortia. <i>Breast Cancer Research</i> , 2014, 16, 3419.	2.2	97
50	Epigenome-wide association study reveals decreased average methylation levels years before breast cancer diagnosis. <i>Clinical Epigenetics</i> , 2015, 7, 67.	1.8	95
51	Healthy lifestyle and risk of breast cancer among postmenopausal women in the European Prospective Investigation into Cancer and Nutrition cohort study. <i>International Journal of Cancer</i> , 2015, 136, 2640-2648.	2.3	95
52	Circulating Steroid Hormone Levels and Risk of Breast Cancer for Postmenopausal Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 492-502.	1.1	94
53	Circulating Insulin-Like Growth Factor-I and Binding Protein-3 and the Risk of Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 763-768.	1.1	93
54	Epigenome-wide methylation in DNA from peripheral blood as a marker of risk for breast cancer. <i>Breast Cancer Research and Treatment</i> , 2014, 148, 665-673.	1.1	93

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55	Carotenoid intakes and risk of breast cancer defined by estrogen receptor and progesterone receptor status: a pooled analysis of 18 prospective cohort studies. <i>American Journal of Clinical Nutrition</i> , 2012, 95, 713-725.	2.2	92
56	Measures of familial aggregation depend on definition of family history: meta-analysis for colorectal cancer. <i>Journal of Clinical Epidemiology</i> , 2006, 59, 114-124.	2.4	89
57	Association of ESR1 gene tagging SNPs with breast cancer risk. <i>Human Molecular Genetics</i> , 2009, 18, 1131-1139.	1.4	84
58	Assessing interactions between the associations of common genetic susceptibility variants, reproductive history and body mass index with breast cancer risk in the breast cancer association consortium: a combined case-control study. <i>Breast Cancer Research</i> , 2010, 12, R110.	2.2	82
59	The role of genetic breast cancer susceptibility variants as prognostic factors. <i>Human Molecular Genetics</i> , 2012, 21, 3926-3939.	1.4	80
60	Does dietary folate intake modify effect of alcohol consumption on breast cancer risk? Prospective cohort study. <i>BMJ: British Medical Journal</i> , 2005, 331, 807.	2.4	77
61	Factors associated with breast cancer recurrences or mortality and dynamic prediction of death using history of cancer recurrences: the French E3N cohort. <i>BMC Cancer</i> , 2018, 18, 171.	1.1	75
62	Associations of common variants at 1p11.2 and 14q24.1 (RAD51L1) with breast cancer risk and heterogeneity by tumor subtype: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2011, 20, 4693-4706.	1.4	71
63	Pre-menopausal serum sex hormone levels in relation to breast cancer risk, overall and by hormone receptor status-Results from the EPIC cohort. <i>International Journal of Cancer</i> , 2014, 134, 1947-1957.	2.3	71
64	Weight change and prostate cancer incidence and mortality. <i>International Journal of Cancer</i> , 2012, 131, 1711-1719.	2.3	70
65	Body Size, Weight Change, and Risk of Colon Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 2978-2986.	1.1	67
66	Alcohol intake and breast cancer in the European prospective investigation into cancer and nutrition. <i>International Journal of Cancer</i> , 2015, 137, 1921-1930.	2.3	65
67	The use of DNA from archival dried blood spots with the Infinium HumanMethylation450 array. <i>BMC Biotechnology</i> , 2013, 13, 23.	1.7	62
68	Relationships between circulating hormone levels, mammographic percent density and breast cancer risk factors in postmenopausal women. <i>Breast Cancer Research and Treatment</i> , 2008, 108, 57-67.	1.1	59
69	Five Polymorphisms and Breast Cancer Risk: Results from the Breast Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 1610-1616.	1.1	57
70	Smoking and blood DNA methylation: an epigenome-wide association study and assessment of reversibility. <i>Epigenetics</i> , 2020, 15, 358-368.	1.3	56
71	Novel Associations between Common Breast Cancer Susceptibility Variants and Risk-Predicting Mammographic Density Measures. <i>Cancer Research</i> , 2015, 75, 2457-2467.	0.4	55
72	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2014, 23, 6096-6111.	1.4	53

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73	Reproductive and hormone-related risk factors for epithelial ovarian cancer by histologic pathways, invasiveness and histologic subtypes: Results from the EPIC cohort. <i>International Journal of Cancer</i> , 2015, 137, 1196-1208.	2.3	53
74	Appraising the causal relevance of DNA methylation for risk of lung cancer. <i>International Journal of Epidemiology</i> , 2019, 48, 1493-1504.	0.9	53
75	Inflammatory Markers and Risk of Epithelial Ovarian Cancer by Tumor Subtypes: The EPIC Cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 951-961.	1.1	51
76	Comparison of 6q25 Breast Cancer Hits from Asian and European Genome Wide Association Studies in the Breast Cancer Association Consortium (BCAC). <i>PLoS ONE</i> , 2012, 7, e42380.	1.1	51
77	Alcohol consumption is associated with widespread changes in blood DNA methylation: Analysis of cross-sectional and longitudinal data. <i>Addiction Biology</i> , 2021, 26, e12855.	1.4	49
78	Dietary Patterns and Prostate Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 3126-3129.	1.1	48
79	Association Between a Germline OCA2 Polymorphism at Chromosome 15q13.1 and Estrogen Receptor-Negative Breast Cancer Survival. <i>Journal of the National Cancer Institute</i> , 2010, 102, 650-662.	3.0	48
80	Risk of Ovarian Cancer and the NF- κ B Pathway: Genetic Association with <i>IL1A</i> and <i>TNFSF10</i> . <i>Cancer Research</i> , 2014, 74, 852-861.	0.4	48
81	Associations of Mammographic Dense and Nondense Areas and Body Mass Index With Risk of Breast Cancer. <i>American Journal of Epidemiology</i> , 2014, 179, 475-483.	1.6	48
82	Vegetable and fruit consumption and the risk of hormone receptor-defined breast cancer in the EPIC cohort. <i>American Journal of Clinical Nutrition</i> , 2016, 103, 168-177.	2.2	48
83	Autosomal dominant inheritance of prostate cancer: a confirmatory study. <i>Urology</i> , 2001, 57, 97-101.	0.5	47
84	AVERAGE VOLUME OF ALCOHOL CONSUMED, TYPE OF BEVERAGE, DRINKING PATTERN AND THE RISK OF DEATH FROM ALL CAUSES. <i>Alcohol and Alcoholism</i> , 2006, 41, 664-671.	0.9	45
85	Mitochondrial DNA copy number variation, leukocyte telomere length, and breast cancer risk in the European Prospective Investigation into Cancer and Nutrition (EPIC) study. <i>Breast Cancer Research</i> , 2018, 20, 29.	2.2	44
86	After hMSH2 and hMLH1? what next? Analysis of three-generational, population-based, early-onset colorectal cancer families. <i>International Journal of Cancer</i> , 2002, 102, 166-171.	2.3	43
87	An epidemiological model for prediction of endometrial cancer risk in Europe. <i>European Journal of Epidemiology</i> , 2016, 31, 51-60.	2.5	43
88	Demographic, lifestyle, and other factors in relation to antimüllerian hormone levels in mostly late premenopausal women. <i>Fertility and Sterility</i> , 2017, 107, 1012-1022.e2.	0.5	43
89	Dietary intake of B vitamins and methionine and breast cancer risk. <i>Cancer Causes and Control</i> , 2013, 24, 1555-1563.	0.8	41
90	Dietary Intake of B Vitamins and Methionine and Colorectal Cancer Risk. <i>Nutrition and Cancer</i> , 2013, 65, 659-667.	0.9	41

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91	Circulating Osteopontin and Prediction of Hepatocellular Carcinoma Development in a Large European Population. <i>Cancer Prevention Research</i> , 2016, 9, 758-765.	0.7	41
92	Serum oncostatin M at baseline predicts mucosal healing in Crohn's disease patients treated with infliximab. <i>Alimentary Pharmacology and Therapeutics</i> , 2020, 52, 284-291.	1.9	41
93	Alcohol consumption and risk of glioblastoma; evidence from the Melbourne collaborative cohort study. <i>International Journal of Cancer</i> , 2011, 128, 1929-1934.	2.3	40
94	Associations between Weight in Early Adulthood, Change in Weight, and Breast Cancer Risk in Postmenopausal Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 1409-1416.	1.1	40
95	Genetic Predisposition to In Situ and Invasive Lobular Carcinoma of the Breast. <i>PLoS Genetics</i> , 2014, 10, e1004285.	1.5	39
96	The effect of socioeconomic status on survival from colorectal cancer in the Melbourne Collaborative Cohort Study. <i>Social Science and Medicine</i> , 2009, 68, 290-297.	1.8	38
97	Dietary intake of B vitamins and methionine and prostate cancer incidence and mortality. <i>Cancer Causes and Control</i> , 2012, 23, 855-863.	0.8	37
98	Post-GWAS gene-environment interplay in breast cancer: results from the Breast and Prostate Cancer Cohort Consortium and a meta-analysis on 79 000 women. <i>Human Molecular Genetics</i> , 2014, 23, 5260-5270.	1.4	37
99	Proportion of premenopausal and postmenopausal breast cancers attributable to known risk factors: Estimates from the EPIC cohort. <i>International Journal of Cancer</i> , 2016, 138, 2415-2427.	2.3	37
100	Physical activity, body size and composition, and risk of ovarian cancer. <i>Cancer Causes and Control</i> , 2010, 21, 2183-2194.	0.8	36
101	Additive Interactions Between Susceptibility Single-Nucleotide Polymorphisms Identified in Genome-Wide Association Studies and Breast Cancer Risk Factors in the Breast and Prostate Cancer Cohort Consortium. <i>American Journal of Epidemiology</i> , 2014, 180, 1018-1027.	1.6	36
102	Endogenous androgens and risk of epithelial invasive ovarian cancer by tumor characteristics in the European Prospective Investigation into Cancer and Nutrition. <i>International Journal of Cancer</i> , 2015, 136, 399-410.	2.3	36
103	Longitudinal Study of Mammographic Density Measures That Predict Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 651-660.	1.1	36
104	Tyrosine Kinase Inhibitors Play an Antiviral Action in Patients Affected by Chronic Myeloid Leukemia: A Possible Model Supporting Their Use in the Fight Against SARS-CoV-2. <i>Frontiers in Oncology</i> , 2020, 10, 1428.	1.3	36
105	The CoV-2 outbreak: how hematologists could help to fight Covid-19. <i>Pharmacological Research</i> , 2020, 157, 104866.	3.1	36
106	11q13 is a susceptibility locus for hormone receptor positive breast cancer. <i>Human Mutation</i> , 2012, 33, 1123-1132.	1.1	35
107	Global measures of peripheral blood-derived DNA methylation as a risk factor in the development of mature B-cell neoplasms. <i>Epigenomics</i> , 2016, 8, 55-66.	1.0	35
108	Investigation of gene-environment interactions between 47 newly identified breast cancer susceptibility loci and environmental risk factors. <i>International Journal of Cancer</i> , 2015, 136, E685-96.	2.3	34

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109	Blood DNA methylation and breast cancer risk: a meta-analysis of four prospective cohort studies. <i>Breast Cancer Research</i> , 2019, 21, 62.	2.2	34
110	Alcohol consumption and prostate cancer risk: Results from the Melbourne collaborative cohort study. <i>International Journal of Cancer</i> , 2006, 119, 1501-1504.	2.3	33
111	Missense Variants in <i>ATM</i> in 26,101 Breast Cancer Cases and 29,842 Controls. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 2143-2151.	1.1	33
112	A large-scale assessment of two-way SNP interactions in breast cancer susceptibility using 46 450 cases and 42 461 controls from the breast cancer association consortium. <i>Human Molecular Genetics</i> , 2014, 23, 1934-1946.	1.4	32
113	A Phase III Preoperative Biomarker Trial of Fenretinide in Ascitic Ovarian Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 1914-1919.	1.1	31
114	Pre-diagnostic polyphenol intake and breast cancer survival: the European Prospective Investigation into Cancer and Nutrition (EPIC) cohort. <i>Breast Cancer Research and Treatment</i> , 2015, 154, 389-401.	1.1	31
115	Genome-Wide Measures of Peripheral Blood Dna Methylation and Prostate Cancer Risk in a Prospective Nested Case-Control Study. <i>Prostate</i> , 2017, 77, 471-478.	1.2	31
116	Joint association of mammographic density adjusted for age and body mass index and polygenic risk score with breast cancer risk. <i>Breast Cancer Research</i> , 2019, 21, 68.	2.2	31
117	Population-Based Estimate of Prostate Cancer Risk for Carriers of the HOXB13 Missense Mutation G84E. <i>PLoS ONE</i> , 2013, 8, e54727.	1.1	31
118	Segregation analysis of urothelial cell carcinoma. <i>European Journal of Cancer</i> , 2006, 42, 1428-1433.	1.3	30
119	Circulating prolactin and in situ breast cancer risk in the European EPIC cohort: a case-control study. <i>Breast Cancer Research</i> , 2015, 17, 49.	2.2	30
120	Mammographic density and risk of breast cancer by mode of detection and tumor size: a case-control study. <i>Breast Cancer Research</i> , 2016, 18, 63.	2.2	30
121	Assessment of serum cytokines predicts clinical and endoscopic outcomes to vedolizumab in ulcerative colitis patients. <i>British Journal of Clinical Pharmacology</i> , 2020, 86, 1296-1305.	1.1	30
122	Reproductive factors and epithelial ovarian cancer survival in the EPIC cohort study. <i>British Journal of Cancer</i> , 2015, 113, 1622-1631.	2.9	29
123	Identifying and correcting epigenetics measurements for systematic sources of variation. <i>Clinical Epigenetics</i> , 2018, 10, 38.	1.8	29
124	Identification of New Genetic Susceptibility Loci for Breast Cancer Through Consideration of Gene-Environment Interactions. <i>Genetic Epidemiology</i> , 2014, 38, 84-93.	0.6	28
125	Asthma, Asthma Medications, and Prostate Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 2318-2324.	1.1	27
126	Confirmation of 5p12 As a Susceptibility Locus for Progesterone-Receptor-Positive, Lower Grade Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 2222-2231.	1.1	27

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127	Acrylamide and Glycidamide Hemoglobin Adducts and Epithelial Ovarian Cancer: A Nested Caseâ€“Control Study in Nonsmoking Postmenopausal Women from the EPIC Cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 127-134.	1.1	27
128	Investigation of circulating metabolites associated with breast cancer risk by untargeted metabolomics: a caseâ€“control study nested within the French E3N cohort. <i>British Journal of Cancer</i> , 2021, 124, 1734-1743.	2.9	27
129	A Two-by-Two Factorial Trial Comparing Oral with Transdermal Estrogen Therapy and Fenretinide with Placebo on Breast Cancer Biomarkers. <i>Clinical Cancer Research</i> , 2004, 10, 4389-4397.	3.2	26
130	Genetic modifiers of menopausal hormone replacement therapy and breast cancer risk: a genomeâ€“wide interaction study. <i>Endocrine-Related Cancer</i> , 2013, 20, 875-887.	1.6	26
131	A treelet transform analysis to relate nutrient patterns to the risk of hormonal receptor-defined breast cancer in the European Prospective Investigation into Cancer and Nutrition (EPIC). <i>Public Health Nutrition</i> , 2016, 19, 242-254.	1.1	26
132	Psychological distress in the academic population and its association with socio-demographic and lifestyle characteristics during COVID-19 pandemic lockdown: Results from a large multicenter Italian study. <i>PLoS ONE</i> , 2021, 16, e0248370.	1.1	26
133	Factors associated with circulating levels of insulin-like growth factor-I and insulin-like growth factor binding protein-3 in 740 women at risk for breast cancer. <i>Breast Cancer Research and Treatment</i> , 2004, 88, 63-73.	1.1	25
134	Change in Body Size and Mortality: Results from the Melbourne Collaborative Cohort Study. <i>PLoS ONE</i> , 2014, 9, e99672.	1.1	25
135	Causal mediation analysis in presence of multiple mediators uncausally related. <i>International Journal of Biostatistics</i> , 2021, 17, 191-221.	0.4	25
136	Change in weight and waist circumference and risk of colorectal cancer: results from the Melbourne Collaborative Cohort Study. <i>BMC Cancer</i> , 2016, 16, 157.	1.1	24
137	Cirrus: An Automated Mammography-Based Measure of Breast Cancer Risk Based on Textural Features. <i>JNCI Cancer Spectrum</i> , 2018, 2, pky057.	1.4	24
138	Predicting interval and screen-detected breast cancers from mammographic density defined by different brightness thresholds. <i>Breast Cancer Research</i> , 2018, 20, 152.	2.2	24
139	Genome-wide association study of subtype-specific epithelial ovarian cancer risk alleles using pooled DNA. <i>Human Genetics</i> , 2014, 133, 481-497.	1.8	23
140	The Premenopausal Breast Cancer Collaboration: A Pooling Project of Studies Participating in the National Cancer Institute Cohort Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 1360-1369.	1.1	23
141	Interval breast cancer risk associations with breast density, family history and breast tissue aging. <i>International Journal of Cancer</i> , 2020, 147, 375-382.	2.3	22
142	Acrylamide and glycidamide hemoglobin adduct levels and endometrial cancer risk: A nested caseâ€“control study in nonsmoking postmenopausal women from the <sc>EPIC</sc> cohort. <i>International Journal of Cancer</i> , 2016, 138, 1129-1138.	2.3	21
143	Epigenome-wide association study for lifetime estrogen exposure identifies an epigenetic signature associated with breast cancer risk. <i>Clinical Epigenetics</i> , 2019, 11, 66.	1.8	21
144	Baseline and lifetime alcohol consumption and risk of differentiated thyroid carcinoma in the EPIC study. <i>British Journal of Cancer</i> , 2015, 113, 840-847.	2.9	20

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145	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , 2016, 141, 386-401.	0.6	18
146	Stochastic Epigenetic Mutations Are Associated with Risk of Breast Cancer, Lung Cancer, and Mature B-cell Neoplasms. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 2026-2037.	1.1	18
147	HCV microRNA elimination in two prisons in Milan, Italy: A model of care. <i>Journal of Viral Hepatitis</i> , 2020, 27, 1444-1454.	1.0	18
148	Novel mammogram-based measures improve breast cancer risk prediction beyond an established mammographic density measure. <i>International Journal of Cancer</i> , 2021, 148, 2193-2202.	2.3	18
149	Telomere Length and Male Fertility. <i>International Journal of Molecular Sciences</i> , 2021, 22, 3959.	1.8	18
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