

# Argyrios Ziogas

## List of Publications by Year in descending order

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Version: 2024-02-01

250  
papers

17,692  
citations

13865

67  
h-index

16650

123  
g-index

256  
all docs

256  
docs citations

256  
times ranked

21280  
citing authors

#	ARTICLE	IF	CITATIONS
1	High Prediagnosis Inflammation-Related Risk Score Associated with Decreased Ovarian Cancer Survival. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 443-452.	2.5	2
2	Better late than never: Brachytherapy is more important than timing in treatment of locally advanced cervical cancer. <i>Gynecologic Oncology</i> , 2022, 164, 348-356.	1.4	11
3	Rare germline copy number variants (CNVs) and breast cancer risk. <i>Communications Biology</i> , 2022, 5, 65.	4.4	6
4	Polygenic risk modeling for prediction of epithelial ovarian cancer risk. <i>European Journal of Human Genetics</i> , 2022, 30, 349-362.	2.8	23
5	Association of Glyphosate Exposure with Blood DNA Methylation in a Cross-Sectional Study of Postmenopausal Women. <i>Environmental Health Perspectives</i> , 2022, 130, 47001.	6.0	9
6	Genome-wide interaction analysis of menopausal hormone therapy use and breast cancer risk among 62,370 women. <i>Scientific Reports</i> , 2022, 12, 6199.	3.3	2
7	Combined Associations of a Polygenic Risk Score and Classical Risk Factors With Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2021, 113, 329-337.	6.3	45
8	Implementation of human papillomavirus video education for women participating in mass cervical cancer screening in Tanzania. <i>American Journal of Obstetrics and Gynecology</i> , 2021, 224, 105.e1-105.e9.	1.3	8
9	Expanding Our Understanding of Ovarian Cancer Risk: The Role of Incomplete Pregnancies. <i>Journal of the National Cancer Institute</i> , 2021, 113, 301-308.	6.3	8
10	Factors Associated with Women's Unwillingness to Decrease Alcohol Intake to Decrease Breast Cancer Risk. <i>Journal of Primary Care and Community Health</i> , 2021, 12, 215013272110002.	2.1	1
11	CYP3A7*1C allele: linking premenopausal oestrogen and progesterone levels with risk of hormone receptor-positive breast cancers. <i>British Journal of Cancer</i> , 2021, 124, 842-854.	6.4	5
12	A case-only study to identify genetic modifiers of breast cancer risk for BRCA1/BRCA2 mutation carriers. <i>Nature Communications</i> , 2021, 12, 1078.	12.8	19
13	A Population-Based Study of Genes Previously Implicated in Breast Cancer. <i>New England Journal of Medicine</i> , 2021, 384, 440-451.	27.0	414
14	Complicated placenta accreta spectrum: identifying a high-risk cohort. <i>Journal of Maternal-Fetal and Neonatal Medicine</i> , 2021, , 1-9.	1.5	0
15	Comparison of Perioperative Outcomes for Radical Nephrectomy Based on Surgical Approach for Masses Greater than 10cm. <i>Journal of Endourology</i> , 2021, 35, 1785-1792.	2.1	2
16	Association of mammographic density with blood DNA methylation. <i>Epigenetics</i> , 2021, , 1-16.	2.7	3
17	Surgery as women's work: gender in presentations at gynecologic conferences. <i>American Journal of Obstetrics and Gynecology</i> , 2021, 225, 454-455.	1.3	2
18	Functional annotation of the 2q35 breast cancer risk locus implicates a structural variant in influencing activity of a long-range enhancer element. <i>American Journal of Human Genetics</i> , 2021, 108, 1190-1203.	6.2	6

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19	Ambient air pollution and ovarian cancer survival in California. <i>Gynecologic Oncology</i> , 2021, 163, 155-161.	1.4	11
20	Association of germline genetic variants with breast cancer-specific survival in patient subgroups defined by clinic-pathological variables related to tumor biology and type of systemic treatment. <i>Breast Cancer Research</i> , 2021, 23, 86.	5.0	7
21	Mendelian randomisation study of smoking exposure in relation to breast cancer risk. <i>British Journal of Cancer</i> , 2021, 125, 1135-1145.	6.4	9
22	Breast Cancer Risk Factors and Survival by Tumor Subtype: Pooled Analyses from the Breast Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 623-642.	2.5	19
23	Germline variants and breast cancer survival in patients with distant metastases at primary breast cancer diagnosis. <i>Scientific Reports</i> , 2021, 11, 19787.	3.3	2
24	Visual inspection with acetic acid screening for cervical cancer among women receiving anti-retroviral therapy for human immunodeficiency virus infection in northern Tanzania. <i>Journal of Obstetrics and Gynaecology Research</i> , 2021, 47, 4365-4370.	1.3	1
25	Endometriosis and menopausal hormone therapy impact the hysterectomy-ovarian cancer association. <i>Gynecologic Oncology</i> , 2021, , .	1.4	5
26	Mammography screening and mortality by risk status in the California teachers study. <i>BMC Cancer</i> , 2021, 21, 1341.	2.6	4
27	Age-dependent interaction between sex and geographic ultraviolet index in melanoma risk. <i>Journal of the American Academy of Dermatology</i> , 2020, 82, 1102-1108.e3.	1.2	22
28	Impact of the affordable care act (ACA) Medicaid expansion on early stage diagnosis and guideline-adherent care for ovarian cancer patients in California. <i>Gynecologic Oncology</i> , 2020, 159, e18-e19.	1.4	0
29	A California Cancer Registry Analysis of Urothelial and Non-urothelial Bladder Cancer Subtypes: Epidemiology, Treatment, and Survival. <i>Clinical Genitourinary Cancer</i> , 2020, 18, e330-e336.	1.9	12
30	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 837-848.	6.2	39
31	Ovarian cancer in California: Guideline adherence, survival, and the impact of geographic location, 1996-2014. <i>Cancer Epidemiology</i> , 2020, 69, 101825.	1.9	7
32	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. <i>Scientific Reports</i> , 2020, 10, 9688.	3.3	2
33	Rationale, Study Design, and Cohort Characteristics for the Markers for Environmental Exposures (MEE) Study. <i>International Journal of Environmental Research and Public Health</i> , 2020, 17, 1774.	2.6	3
34	Multi-Systemic Biological Risk and Cancer Mortality: The NHANES III Study. <i>Scientific Reports</i> , 2020, 10, 5047.	3.3	12
35	Menopausal hormone therapy prior to the diagnosis of ovarian cancer is associated with improved survival. <i>Gynecologic Oncology</i> , 2020, 158, 702-709.	1.4	15
36	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	1.3	32

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37	A Risk-Adjusted Model for Ovarian Cancer Care and Disparities in Access to High-Performing Hospitals. <i>Obstetrics and Gynecology</i> , 2020, 135, 328-339.	2.4	11
38	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , 2020, 11, 312.	12.8	30
39	Disparities by race, socioeconomic status, and insurance type in the receipt of NCCN guideline concordant care for select cancer types in California.. <i>Journal of Clinical Oncology</i> , 2020, 38, 7031-7031.	1.6	8
40	Population-based analysis of guideline adherence for patients with rectal cancer in California.. <i>Journal of Clinical Oncology</i> , 2020, 38, e19045-e19045.	1.6	0
41	Relationship of the breast ductal carcinoma in situ (DCIS) immune microenvironment with clinicopathological features: An institutional experience.. <i>Journal of Clinical Oncology</i> , 2020, 38, e12565-e12565.	1.6	0
42	Disparities in the receipt of National Comprehensive Cancer Network (NCCN) guideline adherent care in triple-negative breast cancer (TNBC) by race/ethnicity, socioeconomic status, and insurance type.. <i>Journal of Clinical Oncology</i> , 2020, 38, 1080-1080.	1.6	0
43	The FANCM:p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. <i>Npj Breast Cancer</i> , 2019, 5, 38.	5.2	28
44	Evaluation of clear cell subtypes of ovarian and uterine malignancies with anti-PD-L1 and anti-PD1 immunohistochemical expression and their association with stage and survival. <i>Gynecologic Oncology</i> , 2019, 155, 483-488.	1.4	10
45	Two truncating variants in FANCC and breast cancer risk. <i>Scientific Reports</i> , 2019, 9, 12524.	3.3	5
46	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	12.8	88
47	Secondhand smoke, obesity, and risk of type II diabetes among California teachers. <i>Annals of Epidemiology</i> , 2019, 32, 35-42.	1.9	9
48	Association between genetically predicted polycystic ovary syndrome and ovarian cancer: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2019, 48, 822-830.	1.9	22
49	Racial and Socioeconomic Disparities in Bladder Cancer Survival: Analysis of the California Cancer Registry. <i>Clinical Genitourinary Cancer</i> , 2019, 17, e995-e1002.	1.9	34
50	Evaluation of vitamin D biosynthesis and pathway target genes reveals UGT2A1/2 and EGFR polymorphisms associated with epithelial ovarian cancer in African American Women. <i>Cancer Medicine</i> , 2019, 8, 2503-2513.	2.8	6
51	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , 2019, 10, 1741.	12.8	90
52	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. <i>American Journal of Human Genetics</i> , 2019, 104, 21-34.	6.2	711
53	A comprehensive gene-environment interaction analysis in Ovarian Cancer using genome-wide significant common variants. <i>International Journal of Cancer</i> , 2019, 144, 2192-2205.	5.1	12
54	Associations of obesity and circulating insulin and glucose with breast cancer risk: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2019, 48, 795-806.	1.9	81

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55	Feasibility of visual inspection with acetic acid (VIA) screening for cervical cancer in Tanzania with emphasis on special populations.. Journal of Clinical Oncology, 2019, 37, 5527-5527.	1.6	2
56	Contribution of Geographic Location to Disparities in Ovarian Cancer Treatment. Journal of the National Comprehensive Cancer Network: JNCCN, 2019, 17, 1318-1329.	4.9	15
57	Identification of a gene expression signature predicting survival in oral cavity squamous cell carcinoma using Monte Carlo cross validation. Oral Oncology, 2018, 78, 72-79.	1.5	2
58	Adult height is associated with increased risk of ovarian cancer: a Mendelian randomisation study. British Journal of Cancer, 2018, 118, 1123-1129.	6.4	15
59	Disparities in Adherence to National Comprehensive Cancer Network Treatment Guidelines and Survival for Stage IB-IIA Cervical Cancer in California. Obstetrics and Gynecology, 2018, 131, 899-908.	2.4	43
60	Assessment of moderate coffee consumption and risk of epithelial ovarian cancer: a Mendelian randomization study. International Journal of Epidemiology, 2018, 47, 450-459.	1.9	15
61	Perception matters: Stressful life events increase breast cancer risk. Journal of Psychosomatic Research, 2018, 110, 46-53.	2.6	14
62	Racial/ethnic differences in the epidemiology of ovarian cancer: a pooled analysis of 12 case-control studies. International Journal of Epidemiology, 2018, 47, 460-472.	1.9	33
63	Nonparametric Adjustment for Measurement Error in Time-to-Event Data: Application to Risk Prediction Models. Journal of the American Statistical Association, 2018, 113, 14-25.	3.1	4
64	Robust Tests for Additive Gene-Environment Interaction in Case-Control Studies Using Gene-Environment Independence. American Journal of Epidemiology, 2018, 187, 366-377.	3.4	8
65	Negative Valence Life Events Promote Breast Cancer Development. Clinical Breast Cancer, 2018, 18, e521-e528.	2.4	5
66	Polycystic Ovary Syndrome, Oligomenorrhea, and Risk of Ovarian Cancer Histotypes: Evidence from the Ovarian Cancer Association Consortium. Cancer Epidemiology Biomarkers and Prevention, 2018, 27, 174-182.	2.5	20
67	Trends in Treatment Patterns and Clinical Outcomes in Young Women Diagnosed With Ductal Carcinoma In Situ. Clinical Breast Cancer, 2018, 18, e179-e185.	2.4	17
68	Breast Cancer Characteristics in Middle Eastern Women Immigrants Compared With Non-Hispanic White Women in California. JNCI Cancer Spectrum, 2018, 2, pky014.	2.9	5
69	Diet Quality Scores Inversely Associated with Postmenopausal Breast Cancer Risk Are Not Associated with Premenopausal Breast Cancer Risk in the California Teachers Study. Journal of Nutrition, 2018, 148, 1830-1837.	2.9	24
70	Variants in genes encoding small GTPases and association with epithelial ovarian cancer susceptibility. PLoS ONE, 2018, 13, e0197561.	2.5	9
71	A Case-Control Study of the Genetic Variability in Reactive Oxygen Species-Metabolizing Enzymes in Melanoma Risk. International Journal of Molecular Sciences, 2018, 19, 242.	4.1	10
72	rs495139 in the TYMS-ENOSF1 Region and Risk of Ovarian Carcinoma of Mucinous Histology. International Journal of Molecular Sciences, 2018, 19, 2473.	4.1	3

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73	Laminin 332 expression and prognosis in breast cancer. <i>Human Pathology</i> , 2018, 82, 289-296.	2.0	18
74	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. <i>Nature Genetics</i> , 2018, 50, 968-978.	21.4	184
75	Enrichment of putative PAX8 target genes at serous epithelial ovarian cancer susceptibility loci. <i>British Journal of Cancer</i> , 2017, 116, 524-535.	6.4	23
76	Cigarette smoking is associated with adverse survival among women with ovarian cancer: Results from a pooled analysis of 19 studies. <i>International Journal of Cancer</i> , 2017, 140, 2422-2435.	5.1	25
77	Impact of community disadvantage and air pollution burden on geographic disparities of ovarian cancer survival in California. <i>Environmental Research</i> , 2017, 156, 388-393.	7.5	34
78	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017, 49, 680-691.	21.4	356
79	Use of common analgesic medications and ovarian cancer survival: results from a pooled analysis in the Ovarian Cancer Association Consortium. <i>British Journal of Cancer</i> , 2017, 116, 1223-1228.	6.4	13
80	Pelvic Inflammatory Disease and the Risk of Ovarian Cancer and Borderline Ovarian Tumors: A Pooled Analysis of 13 Case-Control Studies. <i>American Journal of Epidemiology</i> , 2017, 185, 8-20.	3.4	61
81	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	27.8	1,099
82	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	21.4	289
83	Genome-Wide Testing of Exonic Variants and Breast Cancer Risk in the California Teachers Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 1462-1465.	2.5	0
84	History of Comorbidities and Survival of Ovarian Cancer Patients, Results from the Ovarian Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 1470-1473.	2.5	10
85	Cancer risk in different generations of <i>Middle Eastern</i> immigrants to <i>California</i> , 1988-2013. <i>International Journal of Cancer</i> , 2017, 141, 2260-2269.	5.1	7
86	Sex differences in the association of cutaneous melanoma incidence rates and geographic ultraviolet light exposure. <i>Journal of the American Academy of Dermatology</i> , 2017, 76, 499-505.e3.	1.2	66
87	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. <i>Genetics in Medicine</i> , 2017, 19, 599-603.	2.4	67
88	No Evidence That Genetic Variation in the Myeloid-Derived Suppressor Cell Pathway Influences Ovarian Cancer Survival. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 420-424.	2.5	3
89	Cost-effectiveness of primary debulking surgery when compared to neoadjuvant chemotherapy in the management of stage III C and IV epithelial ovarian cancer. <i>ClinicoEconomics and Outcomes Research</i> , 2016, Volume 8, 397-406.	1.9	10
90	Adult body mass index and risk of ovarian cancer by subtype: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2016, 45, 884-895.	1.9	71

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91	Exome genotyping arrays to identify rare and low frequency variants associated with epithelial ovarian cancer risk. <i>Human Molecular Genetics</i> , 2016, 25, 3600-3612.	2.9	17
92	<i>PALB2</i> , <i>CHEK2</i> and <i>ATM</i> rare variants and cancer risk: data from COGS. <i>Journal of Medical Genetics</i> , 2016, 53, 800-811.	3.2	174
93	Assessing the genetic architecture of epithelial ovarian cancer histological subtypes. <i>Human Genetics</i> , 2016, 135, 741-756.	3.8	19
94	Cancer burden in four countries of the Middle East Cancer Consortium (Cyprus; Jordan; Israel; Izmir) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 <i>Cancer Epidemiology</i> , 2016, 44, 195-202.	1.9	16
95	Association of vitamin D levels and risk of ovarian cancer: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2016, 45, 1619-1630.	1.9	111
96	Risk Prediction for Epithelial Ovarian Cancer in 11 United Statesâ€“Based Case-Control Studies: Incorporation of Epidemiologic Risk Factors and 17 Confirmed Genetic Loci. <i>American Journal of Epidemiology</i> , 2016, 184, 555-569.	3.4	32
97	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. <i>Cancer Discovery</i> , 2016, 6, 1052-1067.	9.4	157
98	Novel polymorphisms in caspase-8 are associated with breast cancer risk in the California Teachers Study. <i>BMC Cancer</i> , 2016, 16, 14.	2.6	18
99	Age- and Tumor Subtypeâ€“Specific Breast Cancer Risk Estimates for <i>CH</i> <i>EK</i> <i>2</i> *110delC Carriers. <i>Journal of Clinical Oncology</i> , 2016, 34, 2750-2760.	1.6	152
100	Treatment for T1a Renal Cancer Substratified by Size: â€œLess is Moreâ€• <i>Journal of Urology</i> , 2016, 196, 1000-1007.	0.4	26
101	The association between socioeconomic status and tumour stage at diagnosis of ovarian cancer: A pooled analysis of 18 case-control studies. <i>Cancer Epidemiology</i> , 2016, 41, 71-79.	1.9	20
102	Investigation of Exomic Variants Associated with Overall Survival in Ovarian Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 446-454.	2.5	9
103	Evidence of a genetic link between endometriosis and ovarian cancer. <i>Fertility and Sterility</i> , 2016, 105, 35-43.e10.	1.0	37
104	Assessment of variation in immunosuppressive pathway genes reveals <i>TGFBR2</i> to be associated with risk of clear cell ovarian cancer. <i>Oncotarget</i> , 2016, 7, 69097-69110.	1.8	5
105	Inherited variants affecting RNA editing may contribute to ovarian cancer susceptibility: results from a large-scale collaboration. <i>Oncotarget</i> , 2016, 7, 72381-72394.	1.8	13
106	A targeted genetic association study of epithelial ovarian cancer susceptibility. <i>Oncotarget</i> , 2016, 7, 7381-7389.	1.8	7
107	Measuring the quality of surgical decisions for Latina breast cancer patients. <i>Health Expectations</i> , 2015, 18, 2389-2400.	2.6	15
108	Epithelialâ€“Mesenchymal Transition (EMT) Gene Variants and Epithelial Ovarian Cancer (EOC) Risk. <i>Genetic Epidemiology</i> , 2015, 39, 689-697.	1.3	22

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109	Clinical Implementation of a Breast Cancer Risk Assessment Program in a Multiethnic Patient Population: Which Risk Model to Use?. <i>Breast Journal</i> , 2015, 21, 562-564.	1.0	6
110	Common Genetic Variation In Cellular Transport Genes and Epithelial Ovarian Cancer (EOC) Risk. <i>PLoS ONE</i> , 2015, 10, e0128106.	2.5	44
111	Sociodemographic Disparities in Advanced Ovarian Cancer Survival and Adherence to Treatment Guidelines. <i>Obstetrics and Gynecology</i> , 2015, 125, 833-842.	2.4	97
112	Prediction of Breast Cancer Risk Based on Profiling With Common Genetic Variants. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	6.3	428
113	Cell-type-specific enrichment of risk-associated regulatory elements at ovarian cancer susceptibility loci. <i>Human Molecular Genetics</i> , 2015, 24, 3595-3607.	2.9	40
114	Observed-to-expected ratio for adherence to treatment guidelines as a quality of care indicator for ovarian cancer. <i>Gynecologic Oncology</i> , 2015, 139, 495-499.	1.4	23
115	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015, 47, 164-171.	21.4	221
116	The Association Between Circulating Total Folate and Folate Vitamers With Overall Survival After Postmenopausal Breast Cancer Diagnosis. <i>Nutrition and Cancer</i> , 2015, 67, 442-448.	2.0	13
117	Network-Based Integration of GWAS and Gene Expression Identifies a <i>HOX</i> -Centric Network Associated with Serous Ovarian Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1574-1584.	2.5	28
118	Socioeconomic status as a predictor of adherence to treatment guidelines for early-stage ovarian cancer. <i>Gynecologic Oncology</i> , 2015, 138, 121-127.	1.4	49
119	Genome-wide Analysis Identifies Novel Loci Associated with Ovarian Cancer Outcomes: Findings from the Ovarian Cancer Association Consortium. <i>Clinical Cancer Research</i> , 2015, 21, 5264-5276.	7.0	33
120	Spatial analysis of advanced-stage ovarian cancer mortality in California. <i>American Journal of Obstetrics and Gynecology</i> , 2015, 213, 43.e1-43.e8.	1.3	25
121	Polymorphisms in a Putative Enhancer at the 10q21.2 Breast Cancer Risk Locus Regulate NRBF2 Expression. <i>American Journal of Human Genetics</i> , 2015, 97, 22-34.	6.2	37
122	Impact of race, socioeconomic status, and the health care system on the treatment of advanced-stage ovarian cancer in California. <i>American Journal of Obstetrics and Gynecology</i> , 2015, 212, 468.e1-468.e9.	1.3	73
123	Costs of treatment for elderly women with advanced ovarian cancer in a Medicare population. <i>Gynecologic Oncology</i> , 2015, 137, 479-484.	1.4	14
124	Evaluating the ovarian cancer gonadotropin hypothesis: A candidate gene study. <i>Gynecologic Oncology</i> , 2015, 136, 542-548.	1.4	15
125	Impact of National Cancer Institute Comprehensive Cancer Centers on Ovarian Cancer Treatment and Survival. <i>Journal of the American College of Surgeons</i> , 2015, 220, 940-950.	0.5	94
126	Cis-eQTL analysis and functional validation of candidate susceptibility genes for high-grade serous ovarian cancer. <i>Nature Communications</i> , 2015, 6, 8234.	12.8	63



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127	Common variants at the <i>CHEK2</i> gene locus and risk of epithelial ovarian cancer. <i>Carcinogenesis</i> , 2015, 36, 1341-1353.	2.8	24
128	Shared genetics underlying epidemiological association between endometriosis and ovarian cancer. <i>Human Molecular Genetics</i> , 2015, 24, 5955-5964.	2.9	68
129	Common Genetic Variation in Circadian Rhythm Genes and Risk of Epithelial Ovarian Cancer (EOC). <i>Journal of Genetics and Genome Research</i> , 2015, 2, .	0.3	25
130	Human Nail Clippings as a Source of DNA for Genetic Studies. <i>Open Journal of Epidemiology</i> , 2015, 05, 41-50.	0.4	8
131	Evaluation of unanticipated 30-day readmission in patients with advanced stage epithelial ovarian cancer.. <i>Journal of Clinical Oncology</i> , 2015, 33, e17684-e17684.	1.6	0
132	Gender difference in the association of melanoma etiology to solar UV exposure.. <i>Journal of Clinical Oncology</i> , 2015, 33, e20012-e20012.	1.6	0
133	Variation in NF- $\kappa$ B Signaling Pathways and Survival in Invasive Epithelial Ovarian Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1421-1427.	2.5	13
134	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.	5.0	97
135	Risk of Ovarian Cancer and the NF- $\kappa$ B Pathway: Genetic Association with <i>IL1A</i> and <i>TNFSF10</i> . <i>Cancer Research</i> , 2014, 74, 852-861.	0.9	48
136	Large-Scale Evaluation of Common Variation in Regulatory T Cell-Related Genes and Ovarian Cancer Outcome. <i>Cancer Immunology Research</i> , 2014, 2, 332-340.	3.4	21
137	Spatial analysis of adherence to treatment guidelines for advanced-stage ovarian cancer and the impact of race and socioeconomic status. <i>Gynecologic Oncology</i> , 2014, 134, 60-67.	1.4	99
138	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.	2.9	32
139	Role of Primary Tumor Resection Among Chemotherapy-Treated Patients with Synchronous Stage IV Colorectal Cancer: A Survival Analysis. <i>Journal of Gastrointestinal Surgery</i> , 2014, 18, 592-598.	1.7	22
140	Genome-wide association study of subtype-specific epithelial ovarian cancer risk alleles using pooled DNA. <i>Human Genetics</i> , 2014, 133, 481-497.	3.8	23
141	High-volume ovarian cancer care: Survival impact and disparities in access for advanced-stage disease. <i>Gynecologic Oncology</i> , 2014, 132, 403-410.	1.4	141
142	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.	2.8	145
143	Attitudes Toward Cancer Clinical Trial Participation in Young Adults with a History of Cancer and a Healthy College Student Sample: A Preliminary Investigation. <i>Journal of Adolescent and Young Adult Oncology</i> , 2014, 3, 20-27.	1.3	25
144	Aspirin, Nonaspirin Nonsteroidal Anti-inflammatory Drug, and Acetaminophen Use and Risk of Invasive Epithelial Ovarian Cancer: A Pooled Analysis in the Ovarian Cancer Association Consortium. <i>Journal of the National Cancer Institute</i> , 2014, 106, djt431-djt431.	6.3	186

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145	Genetic variation in mitotic regulatory pathway genes is associated with breast tumor grade. <i>Human Molecular Genetics</i> , 2014, 23, 6034-6046.	2.9	12
146	Genetic variation at CYP3A is associated with age at menarche and breast cancer risk: a case-control study. <i>Breast Cancer Research</i> , 2014, 16, R51.	5.0	14
147	Dietary Risk Factors for Sporadic Creutzfeldt-Jakob Disease: A Confirmatory Case-Control Study. <i>British Journal of Medicine and Medical Research</i> , 2014, 4, 2388-2417.	0.2	3
148	Ocular Tonometry and Sporadic Creutzfeldt - Jakob disease (sCJD): A Confirmatory Case-Control Study. <i>British Journal of Medicine and Medical Research</i> , 2014, 4, 2322-2333.	0.2	4
149	Multilevel socioeconomic effects on quality of life in adolescent and young adult survivors of leukemia and lymphoma. <i>Quality of Life Research</i> , 2013, 22, 1339-1351.	3.1	19
150	GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 362-370.	21.4	326
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