

# Robert A Vierkant

## List of Publications by Year in descending order

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Version: 2024-02-01

314  
papers

17,150  
citations

15495

65  
h-index

23514

111  
g-index

326  
all docs

326  
docs citations

326  
times ranked

19689  
citing authors

#	ARTICLE	IF	CITATIONS
1	MCM3 is a novel proliferation marker associated with longer survival for patients with tubo-ovarian high-grade serous carcinoma. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2022, 480, 855-871.	1.4	8
2	High Prediagnosis Inflammation-Related Risk Score Associated with Decreased Ovarian Cancer Survival. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 443-452.	1.1	2
3	Polygenic risk modeling for prediction of epithelial ovarian cancer risk. <i>European Journal of Human Genetics</i> , 2022, 30, 349-362.	1.4	23
4	Automated quantification of levels of breast terminal duct lobular (TDLU) involution using deep learning. <i>Npj Breast Cancer</i> , 2022, 8, 13.	2.3	6
5	Serum hormone levels and normal breast histology among premenopausal women. <i>Breast Cancer Research and Treatment</i> , 2022, , .	1.1	0
6	Towards defining morphologic parameters of normal parous and nulliparous breast tissues by artificial intelligence. <i>Breast Cancer Research</i> , 2022, 24, .	2.2	1
7	Cross-Cancer Genome-Wide Association Study of Endometrial Cancer and Epithelial Ovarian Cancer Identifies Genetic Risk Regions Associated with Risk of Both Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 217-228.	1.1	12
8	Population-based targeted sequencing of 54 candidate genes identifies <i>PALB2</i> as a susceptibility gene for high-grade serous ovarian cancer. <i>Journal of Medical Genetics</i> , 2021, 58, 305-313.	1.5	26
9	Associations between tissue-based CD3+ T-cell lymphocyte count and colorectal cancer survival in a prospective cohort of older women. <i>Molecular Carcinogenesis</i> , 2021, 60, 15-24.	1.3	1
10	Pleiotropy-guided transcriptome imputation from normal and tumor tissues identifies candidate susceptibility genes for breast and ovarian cancer. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100042.	1.0	6
11	Clinically Actionable Findings Derived From Predictive Genomic Testing Offered in a Medical Practice Setting. <i>Mayo Clinic Proceedings</i> , 2021, 96, 1407-1417.	1.4	6
12	N-Terminal Pro Brain Natriuretic Peptide, sST2, and Galectin-3 Levels in Breast Cancer Survivors. <i>Journal of Clinical Medicine</i> , 2021, 10, 3313.	1.0	5
13	Somatic mutations in benign breast disease tissues and association with breast cancer risk. <i>BMC Medical Genomics</i> , 2021, 14, 185.	0.7	2
14	Breast Cancer Risk and Use of Nonsteroidal Anti-inflammatory Agents After a Benign Breast Biopsy. <i>Cancer Prevention Research</i> , 2020, 13, 967-976.	0.7	9
15	Clinical and pathological associations of PTEN expression in ovarian cancer: a multicentre study from the Ovarian Tumour Tissue Analysis Consortium. <i>British Journal of Cancer</i> , 2020, 123, 793-802.	2.9	35
16	Bioinformatics and DNA-extraction strategies to reliably detect genetic variants from FFPE breast tissue samples. <i>BMC Genomics</i> , 2019, 20, 689.	1.2	37
17	A combination of the immunohistochemical markers CK7 and SATB2 is highly sensitive and specific for distinguishing primary ovarian mucinous tumors from colorectal and appendiceal metastases. <i>Modern Pathology</i> , 2019, 32, 1834-1846.	2.9	54
18	Episodic and Chronic Migraine in Primary Care. <i>Headache</i> , 2019, 59, 1042-1051.	1.8	21

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19	Genome-wide Analysis of Common Copy Number Variation and Epithelial Ovarian Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 1117-1126.	1.1	21
20	MyD88 and TLR4 Expression in Epithelial Ovarian Cancer. <i>Mayo Clinic Proceedings</i> , 2018, 93, 307-320.	1.4	22
21	Oncologic Safety of Prophylactic Nipple-Sparing Mastectomy in a Population With <i>BRCA</i> Mutations. <i>JAMA Surgery</i> , 2018, 153, 123.	2.2	140
22	Model for Predicting Breast Cancer Risk in Women With Atypical Hyperplasia. <i>Journal of Clinical Oncology</i> , 2018, 36, 1840-1846.	0.8	22
23	Evaluation of 2 breast cancer risk models in a benign breast disease cohort. <i>Cancer</i> , 2018, 124, 3319-3328.	2.0	7
24	Association of p16 expression with prognosis varies across ovarian carcinoma histotypes: an Ovarian Tumor Tissue Analysis consortium study. <i>Journal of Pathology: Clinical Research</i> , 2018, 4, 250-261.	1.3	70
25	Variants in genes encoding small GTPases and association with epithelial ovarian cancer susceptibility. <i>PLoS ONE</i> , 2018, 13, e0197561.	1.1	9
26	rs495139 in the TYMS-ENOSF1 Region and Risk of Ovarian Carcinoma of Mucinous Histology. <i>International Journal of Molecular Sciences</i> , 2018, 19, 2473.	1.8	3
27	Mammographic breast density and risk of breast cancer in women with atypical hyperplasia: an observational cohort study from the Mayo Clinic Benign Breast Disease (BBD) cohort. <i>BMC Cancer</i> , 2017, 17, 84.	1.1	23
28	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017, 49, 680-691.	9.4	356
29	Cytotoxic T Cells and Granzyme B Associated with Improved Colorectal Cancer Survival in a Prospective Cohort of Older Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 622-631.	1.1	68
30	Breast Cancer Risk and Progressive Histology in Serial Benign Biopsies. <i>Journal of the National Cancer Institute</i> , 2017, 109, .	3.0	10
31	Dose-Response Association of CD8 <sup>+</sup> Tumor-Infiltrating Lymphocytes and Survival Time in High-Grade Serous Ovarian Cancer. <i>JAMA Oncology</i> , 2017, 3, e173290.	3.4	260
32	Postlactational involution biomarkers plasminogen and phospho-STAT3 are linked with active age-related lobular involution. <i>Breast Cancer Research and Treatment</i> , 2017, 166, 133-143.	1.1	0
33	NanoString-based breast cancer risk prediction for women with sclerosing adenosis. <i>Breast Cancer Research and Treatment</i> , 2017, 166, 641-650.	1.1	10
34	Association between mammographic breast density and histologic features of benign breast disease. <i>Breast Cancer Research</i> , 2017, 19, 134.	2.2	24
35	<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.	1.5	174
36	Chronic Recreational Physical Inactivity and Epithelial Ovarian Cancer Risk: Evidence from the Ovarian Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 1114-1124.	1.1	32

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37	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.	1.6	32
38	Breast cancer risk by the extent and type of atypical hyperplasia. <i>Cancer</i> , 2016, 122, 3087-3088.	2.0	10
39	Extent of atypical hyperplasia stratifies breast cancer risk in 2 independent cohorts of women. <i>Cancer</i> , 2016, 122, 2971-2978.	2.0	48
40	The inflammatory microenvironment in epithelial ovarian cancer: a role for TLR4 and MyD88 and related proteins. <i>Tumor Biology</i> , 2016, 37, 13279-13286.	0.8	41
41	Benign Breast Disease and the Risk of Breast Cancer. <i>Obstetrical and Gynecological Survey</i> , 2016, 71, 472-473.	0.2	6
42	Chronic Recreational Physical Inactivity and Epithelial Ovarian Cancer Risk. <i>Obstetrical and Gynecological Survey</i> , 2016, 71, 528-530.	0.2	0
43	Clinicopathologic features of breast cancers that develop in women with previous benign breast disease. <i>Cancer</i> , 2016, 122, 378-385.	2.0	31
44	Impaired innate, humoral, and cellular immunity despite a take in smallpox vaccine recipients. <i>Vaccine</i> , 2016, 34, 3283-3290.	1.7	16
45	Recreational physical inactivity and mortality in women with invasive epithelial ovarian cancer: evidence from the Ovarian Cancer Association Consortium. <i>British Journal of Cancer</i> , 2016, 115, 95-101.	2.9	39
46	Hormonal and Reproductive Factors and Risk of Myeloproliferative Neoplasms in Postmenopausal Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 151-157.	1.1	3
47	Personalizing Aspirin Use for Targeted Breast Cancer Chemoprevention in Postmenopausal Women. <i>Mayo Clinic Proceedings</i> , 2016, 91, 71-80.	1.4	20
48	Assessment of Multifactor Geneâ€‘Environment Interactions and Ovarian Cancer Risk: Candidate Genes, Obesity, and Hormone-Related Risk Factors. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 780-790.	1.1	10
49	Clinical and Emergent Biomarkers and Their Relationship to the Prognosis of Ovarian Cancer. <i>Oncology</i> , 2016, 90, 59-68.	0.9	9
50	Tumor eosinophil infiltration and improved survival of colorectal cancer patients: Iowa Women's Health Study. <i>Modern Pathology</i> , 2016, 29, 516-527.	2.9	65
51	Natural history of age-related lobular involution and impact on breast cancer risk. <i>Breast Cancer Research and Treatment</i> , 2016, 155, 423-430.	1.1	29
52	Investigation of Exomic Variants Associated with Overall Survival in Ovarian Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 446-454.	1.1	9
53	Mucocele-like lesions of the breast: a clinical outcome and histologic analysis of 102 cases. <i>Human Pathology</i> , 2016, 49, 33-38.	1.1	29
54	Evidence of a genetic link between endometriosis and ovarian cancer. <i>Fertility and Sterility</i> , 2016, 105, 35-43.e10.	0.5	37

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55	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , 2016, 141, 386-401.	0.6	18
56	Assessment of variation in immunosuppressive pathway genes reveals TGFBR2 to be associated with risk of clear cell ovarian cancer. <i>Oncotarget</i> , 2016, 7, 69097-69110.	0.8	5
57	A targeted genetic association study of epithelial ovarian cancer susceptibility. <i>Oncotarget</i> , 2016, 7, 7381-7389.	0.8	7
58	Prior oral contraceptive use in ovarian cancer patients: assessing associations with overall and progression-free survival. <i>BMC Cancer</i> , 2015, 15, 711.	1.1	9
59	Epithelialâ€Mesenchymal Transition (EMT) Gene Variants and Epithelial Ovarian Cancer (EOC) Risk. <i>Genetic Epidemiology</i> , 2015, 39, 689-697.	0.6	22
60	Tumor Budding in Colorectal Carcinoma. <i>American Journal of Surgical Pathology</i> , 2015, 39, 1340-1346.	2.1	95
61	Common Genetic Variation In Cellular Transport Genes and Epithelial Ovarian Cancer (EOC) Risk. <i>PLoS ONE</i> , 2015, 10, e0128106.	1.1	44
62	Associations between Environmental Exposures and Incident Colorectal Cancer by ESR2 Protein Expression Level in a Population-Based Cohort of Older Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 713-719.	1.1	10
63	Cell-type-specific enrichment of risk-associated regulatory elements at ovarian cancer susceptibility loci. <i>Human Molecular Genetics</i> , 2015, 24, 3595-3607.	1.4	40
64	Flat epithelial atypia and risk of breast cancer: A Mayo cohort study. <i>Cancer</i> , 2015, 121, 1548-1555.	2.0	85
65	Model for Individualized Prediction of Breast Cancer Risk After a Benign Breast Biopsy. <i>Journal of Clinical Oncology</i> , 2015, 33, 923-929.	0.8	51
66	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015, 47, 164-171.	9.4	221
67	Genome-wide significant risk associations for mucinous ovarian carcinoma. <i>Nature Genetics</i> , 2015, 47, 888-897.	9.4	78
68	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.	1.1	28
69	Obesity and survival among women with ovarian cancer: results from the Ovarian Cancer Association Consortium. <i>British Journal of Cancer</i> , 2015, 113, 817-826.	2.9	111
70	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.	3.2	33
71	Evaluating the ovarian cancer gonadotropin hypothesis: A candidate gene study. <i>Gynecologic Oncology</i> , 2015, 136, 542-548.	0.6	15
72	Ki-67 expression in sclerosing adenosis and adjacent normal breast terminal ductal lobular units: a nested caseâ€control study from the Mayo Benign Breast Disease Cohort. <i>Breast Cancer Research and Treatment</i> , 2015, 151, 89-97.	1.1	13

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73	Cis-eQTL analysis and functional validation of candidate susceptibility genes for high-grade serous ovarian cancer. <i>Nature Communications</i> , 2015, 6, 8234.	5.8	63
74	Common variants at the <i>CHEK2</i> gene locus and risk of epithelial ovarian cancer. <i>Carcinogenesis</i> , 2015, 36, 1341-1353.	1.3	24
75	Reply to M.H. Gail et al. <i>Journal of Clinical Oncology</i> , 2015, 33, 2830-2831.	0.8	0
76	Gene signature model for breast cancer risk prediction for women with sclerosing adenosis. <i>Breast Cancer Research and Treatment</i> , 2015, 152, 687-694.	1.1	11
77	Complex fibroadenoma and breast cancer risk: a Mayo Clinic Benign Breast Disease Cohort Study. <i>Breast Cancer Research and Treatment</i> , 2015, 153, 397-405.	1.1	61
78	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
79	Understanding the Premalignant Potential of Atypical Hyperplasia through Its Natural History: A Longitudinal Cohort Study. <i>Cancer Prevention Research</i> , 2014, 7, 211-217.	0.7	192
80	Evidence for a time-dependent association between FOLR1 expression and survival from ovarian carcinoma: implications for clinical testing. An Ovarian Tumour Tissue Analysis consortium study. <i>British Journal of Cancer</i> , 2014, 111, 2297-2307.	2.9	76
81	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.	1.1	13
82	Anthropometric, medical history and lifestyle risk factors for myeloproliferative neoplasms in The Iowa Women's Health Study cohort. <i>International Journal of Cancer</i> , 2014, 134, 1741-1750.	2.3	42
83	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.	0.4	48
84	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.	1.6	21
85	Regional Differences in Breast Cancer Biomarkers in American Indian and Alaska Native Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 409-415.	1.1	6
86	Sclerosing adenosis and risk of breast cancer. <i>Breast Cancer Research and Treatment</i> , 2014, 144, 205-212.	1.1	72
87	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
88	Associations between Cigarette Smoking, Hormone Therapy, and Folate Intake with Incident Colorectal Cancer by TP53 Protein Expression Level in a Population-Based Cohort of Older Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 350-355.	1.1	11
89	Kernel canonical correlation analysis for assessing gene-gene interactions and application to ovarian cancer. <i>European Journal of Human Genetics</i> , 2014, 22, 126-131.	1.4	33
90	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

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91	Consortium analysis of gene and gene-gene interactions in purine and pyrimidine metabolism pathways with ovarian carcinoma risk. <i>Molecular Nutrition and Food Research</i> , 2014, 58, 2023-2035.	1.5	16
92	Development and Pilot Evaluation of Native CREST—a Cancer Research Experience and Student Training Program for Navajo Undergraduate Students. <i>Journal of Cancer Education</i> , 2013, 28, 92-99.	0.6	4
93	Hormone-receptor expression and ovarian cancer survival: an Ovarian Tumor Tissue Analysis consortium study. <i>Lancet Oncology</i> , The, 2013, 14, 853-862.	5.1	335
94	ABCB1 (MDR1) polymorphisms and ovarian cancer progression and survival: A comprehensive analysis from the Ovarian Cancer Association Consortium and The Cancer Genome Atlas. <i>Gynecologic Oncology</i> , 2013, 131, 8-14.	0.6	55
95	GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 362-370.	9.4	326
96	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
97	Cigarette smoking and risk of ovarian cancer: a pooled analysis of 21 case-control studies. <i>Cancer Causes and Control</i> , 2013, 24, 989-1004.	0.8	84
98	Associations Between Colorectal Cancer Molecular Markers and Pathways With Clinicopathologic Features in Older Women. <i>Gastroenterology</i> , 2013, 145, 348-356.e2.	0.6	49
99	Associations between polymorphisms in the antiviral TRIM genes and measles vaccine immunity. <i>Human Immunology</i> , 2013, 74, 768-774.	1.2	24
100	Intake of coffee, caffeine and other methylxanthines and risk of Type I vs Type II endometrial cancer. <i>British Journal of Cancer</i> , 2013, 109, 1908-1913.	2.9	33
101	Epigenome-wide ovarian cancer analysis identifies a methylation profile differentiating clear-cell histology with epigenetic silencing of the HERG K+ channel. <i>Human Molecular Genetics</i> , 2013, 22, 3038-3047.	1.4	60
102	Human Leukocyte Antigens and Cellular Immune Responses to Anthrax Vaccine Adsorbed. <i>Infection and Immunity</i> , 2013, 81, 2584-2591.	1.0	22
103	Biomarker-Based Ovarian Carcinoma Typing: A Histologic Investigation in the Ovarian Tumor Tissue Analysis Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 1677-1686.	1.1	70
104	Combined and Interactive Effects of Environmental and GWAS-Identified Risk Factors in Ovarian Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 880-890.	1.1	54
105	Obesity and risk of ovarian cancer subtypes: evidence from the Ovarian Cancer Association Consortium. <i>Endocrine-Related Cancer</i> , 2013, 20, 251-262.	1.6	169
106	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
107	Analysis of Over 10,000 Cases Finds No Association between Previously Reported Candidate Polymorphisms and Ovarian Cancer Outcome. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 987-992.	1.1	20
108	Identification and molecular characterization of a new ovarian cancer susceptibility locus at 17q21.31. <i>Nature Communications</i> , 2013, 4, 1627.	5.8	98

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109	Inherited Variants in Regulatory T Cell Genes and Outcome of Ovarian Cancer. <i>PLoS ONE</i> , 2013, 8, e53903.	1.1	20
110	Associations Between Intake of Folate and Related Micronutrients with Molecularly Defined Colorectal Cancer Risks in the Iowa Women's Health Study. <i>Nutrition and Cancer</i> , 2012, 64, 899-910.	0.9	33
111	Gene Set Analysis of Survival Following Ovarian Cancer Implicates Macrolide Binding and Intracellular Signaling Genes. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 529-536.	1.1	7
112	Identification of a novel percent mammographic density locus at 12q24. <i>Human Molecular Genetics</i> , 2012, 21, 3299-3305.	1.4	31
113	Common Variation in Nemo-Like Kinase Is Associated with Risk of Ovarian Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 523-528.	1.1	8
114	Ovarian Cancer Risk Associated with Inherited Inflammation-Related Variants. <i>Cancer Research</i> , 2012, 72, 1064-1069.	0.4	45
115	Atypical Apocrine Adenosis of the Breast: Long-term Follow-up in 37 Patients. <i>Archives of Pathology and Laboratory Medicine</i> , 2012, 136, 179-182.	1.2	30
116	Effects of vitamin A and D receptor gene polymorphisms/haplotypes on immune responses to measles vaccine. <i>Pharmacogenetics and Genomics</i> , 2012, 22, 20-31.	0.7	38
117	Genome-wide analysis of polymorphisms associated with cytokine responses in smallpox vaccine recipients. <i>Human Genetics</i> , 2012, 131, 1403-1421.	1.8	75
118	Genome-wide genetic associations with IFN $\gamma$ response to smallpox vaccine. <i>Human Genetics</i> , 2012, 131, 1433-1451.	1.8	47
119	ABO blood group and risk of epithelial ovarian cancer within the Ovarian Cancer Association Consortium. <i>Cancer Causes and Control</i> , 2012, 23, 1805-1810.	0.8	35
120	Consistency of HLA associations between two independent measles vaccine cohorts: A replication study. <i>Vaccine</i> , 2012, 30, 2146-2152.	1.7	44
121	Independence of measles-specific humoral and cellular immune responses to vaccination. <i>Human Immunology</i> , 2012, 73, 474-479.	1.2	14
122	Replication of associations between cytokine and cytokine receptor single nucleotide polymorphisms and measles-specific adaptive immunophenotypic extremes. <i>Human Immunology</i> , 2012, 73, 636-640.	1.2	12
123	Postmenopausal Hormone Therapy and Colorectal Cancer Risk in Relation to Somatic <i>KRAS</i> Mutation Status among Older Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 681-684.	1.1	25
124	Cigarette Smoking and Colorectal Cancer Risk by <i>KRAS</i> Mutation Status Among Older Women. <i>American Journal of Gastroenterology</i> , 2012, 107, 782-789.	0.2	32
125	Postmenopausal hormone therapy and colorectal cancer risk by molecularly defined subtypes among older women. <i>Gut</i> , 2012, 61, 1299-1305.	6.1	36
126	Impact of cytokine and cytokine receptor gene polymorphisms on cellular immunity after smallpox vaccination. <i>Gene</i> , 2012, 510, 59-65.	1.0	34

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127	Regular Multivitamin Supplement Use, Single Nucleotide Polymorphisms in ATIC, SHMT2, and SLC46A1, and Risk of Ovarian Carcinoma. <i>Frontiers in Genetics</i> , 2012, 3, 33.	1.1	4
128	Germline Copy Number Variation and Ovarian Cancer Survival. <i>Frontiers in Genetics</i> , 2012, 3, 142.	1.1	11
129	Histologic findings in normal breast tissues: comparison to reduction mammoplasty and benign breast disease tissues. <i>Breast Cancer Research and Treatment</i> , 2012, 133, 169-177.	1.1	64
130	Response surface methodology to determine optimal measles-specific cytokine responses in human peripheral blood mononuclear cells. <i>Journal of Immunological Methods</i> , 2012, 382, 220-223.	0.6	2
131	European American Stratification in Ovarian Cancer Case Control Data: The Utility of Genome-Wide Data for Inferring Ancestry. <i>PLoS ONE</i> , 2012, 7, e35235.	1.1	3
132	Alcohol Intake and Colorectal Cancer Risk by Molecularly Defined Subtypes in a Prospective Study of Older Women. <i>Cancer Prevention Research</i> , 2011, 4, 2035-2043.	0.7	17
133	Progesterone receptor gene polymorphisms and risk of endometriosis: results from an international collaborative effort. <i>Fertility and Sterility</i> , 2011, 95, 40-45.	0.5	20
134	Human leukocyte antigen associations with humoral and cellular immunity following a second dose of measles-containing vaccine: Persistence, dampening, and extinction of associations found after a first dose. <i>Vaccine</i> , 2011, 29, 7982-7991.	1.7	17
135	Associations between single nucleotide polymorphisms and haplotypes in cytokine and cytokine receptor genes and immunity to measles vaccination. <i>Vaccine</i> , 2011, 29, 7883-7895.	1.7	62
136	Genetic polymorphisms in host antiviral genes: Associations with humoral and cellular immunity to measles vaccine. <i>Vaccine</i> , 2011, 29, 8988-8997.	1.7	64
137	Correlations Between Vaccinia-Specific Immune Responses Within a Cohort of Armed Forces Members. <i>Viral Immunology</i> , 2011, 24, 415-420.	0.6	16
138	Functional Polymorphisms in the TERT Promoter Are Associated with Risk of Serous Epithelial Ovarian and Breast Cancers. <i>PLoS ONE</i> , 2011, 6, e24987.	1.1	48
139	p16INK4a Expression and Breast Cancer Risk in Women with Atypical Hyperplasia. <i>Cancer Prevention Research</i> , 2011, 4, 1953-1960.	0.7	22
140	Centrosome-related genes, genetic variation, and risk of breast cancer. <i>Breast Cancer Research and Treatment</i> , 2011, 125, 221-228.	1.1	42
141	Effect of aspirin and other NSAIDs on postmenopausal breast cancer incidence by hormone receptor status: results from a prospective cohort study. <i>Breast Cancer Research and Treatment</i> , 2011, 126, 149-155.	1.1	82
142	The role of polymorphisms in Toll-like receptors and their associated intracellular signaling genes in measles vaccine immunity. <i>Human Genetics</i> , 2011, 130, 547-61.	1.8	60
143	Xenobioticâ€Metabolizing gene polymorphisms and ovarian cancer risk. <i>Molecular Carcinogenesis</i> , 2011, 50, 397-402.	1.3	29
144	Inherited Variants in Mitochondrial Biogenesis Genes May Influence Epithelial Ovarian Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 1131-1145.	1.1	62

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145	Platinum Sensitivity-Related Germline Polymorphism Discovered via a Cell-Based Approach and Analysis of Its Association with Outcome in Ovarian Cancer Patients. <i>Clinical Cancer Research</i> , 2011, 17, 5490-5500.	3.2	57
146	Assessment of Hepatocyte Growth Factor in Ovarian Cancer Mortality. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 1638-1648.	1.1	31
147	Prostate Cancer Susceptibility Polymorphism rs2660753 Is Not Associated with Invasive Ovarian Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 1028-1031.	1.1	0
148	The Role of KRAS rs61764370 in Invasive Epithelial Ovarian Cancer: Implications for Clinical Testing. <i>Clinical Cancer Research</i> , 2011, 17, 3742-3750.	3.2	47
149	Risk factors for meningioma in postmenopausal women: results from the Iowa Women's Health Study. <i>Neuro-Oncology</i> , 2011, 13, 1011-1019.	0.6	50
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