

# Irene Orlow

## List of Publications by Year in descending order

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151  
papers

8,575  
citations

57631

44  
h-index

49773

87  
g-index

154  
all docs

154  
docs citations

154  
times ranked

13753  
citing authors

#	ARTICLE	IF	CITATIONS
1	The Ink4a Tumor Suppressor Gene Product, p19Arf, Interacts with MDM2 and Neutralizes MDM2's Inhibition of p53. <i>Cell</i> , 1998, 92, 713-723.	13.5	1,412
2	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
3	GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 362-370.	9.4	326
4	Tumor-Infiltrating Lymphocyte Grade in Primary Melanomas Is Independently Associated With Melanoma-Specific Survival in the Population-Based Genes, Environment and Melanoma Study. <i>Journal of Clinical Oncology</i> , 2013, 31, 4252-4259.	0.8	232
5	Common variation at 2p13.3, 3q29, 7p13 and 17q25.1 associated with susceptibility to pancreatic cancer. <i>Nature Genetics</i> , 2015, 47, 911-916.	9.4	224
6	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015, 47, 164-171.	9.4	221
7	Lifetime Risk of Melanoma in CDKN2A Mutation Carriers in a Population-Based Sample. <i>Journal of the National Cancer Institute</i> , 2005, 97, 1507-1515.	3.0	200
8	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. <i>Nature Communications</i> , 2018, 9, 556.	5.8	188
9	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , 2018, 9, 3166.	5.8	178
10	Association Between <i>NRAS</i> and <i>BRAF</i> Mutational Status and Melanoma-Specific Survival Among Patients With Higher-Risk Primary Melanoma. <i>JAMA Oncology</i> , 2015, 1, 359.	3.4	164
11	Deletions of the INK4A Gene Occur in Malignant Peripheral Nerve Sheath Tumors but not in Neurofibromas. <i>American Journal of Pathology</i> , 1999, 155, 1855-1860.	1.9	161
12	Previous Lung Diseases and Lung Cancer Risk: A Pooled Analysis From the International Lung Cancer Consortium. <i>American Journal of Epidemiology</i> , 2012, 176, 573-585.	1.6	160
13	Genomic and Mutational Profiling to Assess Clonal Relationships Between Multiple Non-Small Cell Lung Cancers. <i>Clinical Cancer Research</i> , 2009, 15, 5184-5190.	3.2	151
14	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
15	Increased risk of lung cancer in individuals with a family history of the disease: A pooled analysis from the International Lung Cancer Consortium. <i>European Journal of Cancer</i> , 2012, 48, 1957-1968.	1.3	143
16	Comparison of Clinicopathologic Features and Survival of Histopathologically Amelanotic and Pigmented Melanomas. <i>JAMA Dermatology</i> , 2014, 150, 1306.	2.0	142
17	Cannabis smoking and lung cancer risk: Pooled analysis in the International Lung Cancer Consortium. <i>International Journal of Cancer</i> , 2015, 136, 894-903.	2.3	131
18	Deletions of the INK4A Gene in Superficial Bladder Tumors. <i>American Journal of Pathology</i> , 1999, 155, 105-113.	1.9	121

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19	The Prevalence of CDKN2A Germ-Line Mutations and Relative Risk for Cutaneous Malignant Melanoma: An International Population-Based Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 1520-1525.	1.1	105
20	Variants in Estrogen Biosynthesis Genes, Sex Steroid Hormone Levels, and Endometrial Cancer: A HuGE Review. <i>American Journal of Epidemiology</i> , 2007, 165, 235-245.	1.6	102
21	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. <i>American Journal of Human Genetics</i> , 2015, 96, 487-497.	2.6	101
22	Identification and molecular characterization of a new ovarian cancer susceptibility locus at 17q21.31. <i>Nature Communications</i> , 2013, 4, 1627.	5.8	98
23	Molecular analyses of the mitotic checkpoint components hMAD2, hBUB1 and hBUB3 in human cancer. <i>International Journal of Cancer</i> , 2001, 95, 223-227.	2.3	92
24	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , 2016, 7, 11843.	5.8	86
25	Analysis of Genetic Variants in Never-Smokers with Lung Cancer Facilitated by an Internet-Based Blood Collection Protocol: A Preliminary Report. <i>Clinical Cancer Research</i> , 2010, 16, 755-763.	3.2	82
26	Genome-wide significant risk associations for mucinous ovarian carcinoma. <i>Nature Genetics</i> , 2015, 47, 888-897.	9.4	78
27	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016, 7, 12675.	5.8	78
28	Asthma and lung cancer risk: a systematic investigation by the International Lung Cancer Consortium. <i>Carcinogenesis</i> , 2012, 33, 587-597.	1.3	69
29	The oral microbiota in patients with pancreatic cancer, patients with IPMNs, and controls: a pilot study. <i>Cancer Causes and Control</i> , 2017, 28, 959-969.	0.8	69
30	Phase II Study of Extended-Dose Temozolomide in Patients With Melanoma. <i>Journal of Clinical Oncology</i> , 2008, 26, 2299-2304.	0.8	66
31	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
32	Vitamin D receptor polymorphisms in patients with cutaneous melanoma. <i>International Journal of Cancer</i> , 2012, 130, 405-418.	2.3	61
33	A Transcriptome-Wide Association Study Identifies Novel Candidate Susceptibility Genes for Pancreatic Cancer. <i>Journal of the National Cancer Institute</i> , 2020, 112, 1003-1012.	3.0	59
34	Allergies, variants in IL-4 and IL-4R $\alpha$ genes, and risk of pancreatic cancer. <i>Cancer Detection and Prevention</i> , 2007, 31, 345-351.	2.1	58
35	The Obesity-Associated Polymorphisms FTO rs9939609 and MC4R rs17782313 and Endometrial Cancer Risk in Non-Hispanic White Women. <i>PLoS ONE</i> , 2011, 6, e16756.	1.1	58
36	DNA Damage and Repair Capacity in Patients With Lung Cancer: Prediction of Multiple Primary Tumors. <i>Journal of Clinical Oncology</i> , 2008, 26, 3560-3566.	0.8	56

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37	Prognostic Significance of Transcription Factor E2F-1 in Bladder Cancer: Genotypic and Phenotypic Characterization. <i>Journal of the National Cancer Institute</i> , 1999, 91, 874-881.	3.0	54
38	Vitamin D receptor polymorphisms and survival in patients with cutaneous melanoma: a population-based study. <i>Carcinogenesis</i> , 2016, 37, 30-38.	1.3	54
39	CDKN2A Germline Mutations in Individuals with Cutaneous Malignant Melanoma. <i>Journal of Investigative Dermatology</i> , 2007, 127, 1234-1243.	0.3	50
40	<i>APOE</i> polymorphisms and cognitive functions in patients with brain tumors. <i>Neurology</i> , 2014, 83, 320-327.	1.5	49
41	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
42	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
43	Analysis of Heritability and Genetic Architecture of Pancreatic Cancer: A PanC4 Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 1238-1245.	1.1	48
44	Aspirin and NSAID use and lung cancer risk: a pooled analysis in the International Lung Cancer Consortium (ILCCO). <i>Cancer Causes and Control</i> , 2011, 22, 1709-1720.	0.8	47
45	A metastasis or a second independent cancer? Evaluating the clonal origin of tumors using array copy number data. <i>Statistics in Medicine</i> , 2010, 29, 1608-1621.	0.8	46
46	Associations of Cumulative Sun Exposure and Phenotypic Characteristics with Histologic Solar Elastosis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 2932-2941.	1.1	45
47	Clinicopathologic Features of Incident and Subsequent Tumors in Patients with Multiple Primary Cutaneous Melanomas. <i>Annals of Surgical Oncology</i> , 2012, 19, 1024-1033.	0.7	45
48	<i>COMT</i> , <i>BDNF</i> , and <i>DTNBP1</i> polymorphisms and cognitive functions in patients with brain tumors. <i>Neuro-Oncology</i> , 2016, 18, 1425-1433.	0.6	45
49	The hCOMET project: International database comparison of results with the comet assay in human biomonitoring. Baseline frequency of DNA damage and effect of main confounders. <i>Mutation Research - Reviews in Mutation Research</i> , 2021, 787, 108371.	2.4	45
50	Matrix Metalloproteinase-9 (MMP-9) polymorphisms in patients with cutaneous malignant melanoma. <i>BMC Medical Genetics</i> , 2007, 8, 10.	2.1	44
51	Common Genetic Variation In Cellular Transport Genes and Epithelial Ovarian Cancer (EOC) Risk. <i>PLoS ONE</i> , 2015, 10, e0128106.	1.1	44
52	Risk of Non-Melanoma Cancers in First-Degree Relatives of CDKN2A Mutation Carriers. <i>Journal of the National Cancer Institute</i> , 2012, 104, 953-956.	3.0	42
53	Genome-wide association study of endometrial cancer in E2C2. <i>Human Genetics</i> , 2014, 133, 211-224.	1.8	42
54	A Replication Study and Genome-Wide Scan of Single-Nucleotide Polymorphisms Associated with Pancreatic Cancer Risk and Overall Survival. <i>Clinical Cancer Research</i> , 2012, 18, 3942-3951.	3.2	40

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55	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
56	Contralateral breast cancers: Independent cancers or metastases?. <i>International Journal of Cancer</i> , 2018, 142, 347-356.	2.3	37
57	Inherited Genetic Variants Associated with Occurrence of Multiple Primary Melanoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 992-997.	1.1	36
58	Vitamin D Receptor Gene Haplotypes and Polymorphisms and Risk of Breast Cancer: A Nested Caseâ€“Control Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 1856-1867.	1.1	35
59	<i>MITF</i> E318K's effect on melanoma risk independent of, but modified by, other risk factors. <i>Pigment Cell and Melanoma Research</i> , 2014, 27, 485-488.	1.5	35
60	Alcohol and lung cancer risk among never smokers: A pooled analysis from the international lung cancer consortium and the SYNERGY study. <i>International Journal of Cancer</i> , 2017, 140, 1976-1984.	2.3	35
61	Microsatellite instability and deletion analysis of chromosome 10 in human prostate cancer. , 1996, 69, 110-113.		34
62	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
63	Genotypic and phenotypic characterization of the histoblood group ABO(H) in primary bladder tumors. , 1998, 75, 819-824.		31
64	Genetic factors associated with naevus count and dermoscopic patterns: preliminary results from the Study of Nevi in Children ( <i>SONIC</i> ). <i>British Journal of Dermatology</i> , 2015, 172, 1081-1089.	1.4	31
65	Variants in hormone biosynthesis genes and risk of endometrial cancer. <i>Cancer Causes and Control</i> , 2008, 19, 955-963.	0.8	29
66	Vitamin D Metabolic Pathway Genes and Pancreatic Cancer Risk. <i>PLoS ONE</i> , 2015, 10, e0117574.	1.1	29
67	Association Between Aryl Hydrocarbon Receptor Genotype and Survival in Soft Tissue Sarcoma. <i>Journal of Clinical Oncology</i> , 2004, 22, 3997-4001.	0.8	28
68	Sun exposure, vitamin D receptor polymorphisms FokI and BsmI and risk of multiple primary melanoma. <i>Cancer Epidemiology</i> , 2011, 35, e105-e110.	0.8	28
69	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
70	Association of Interferon Regulatory Factor-4 Polymorphism rs12203592 With Divergent Melanoma Pathways. <i>Journal of the National Cancer Institute</i> , 2016, 108, djw004.	3.0	28
71	Menstrual and reproductive factors and lung cancer risk: A pooled analysis from the international lung cancer consortium. <i>International Journal of Cancer</i> , 2017, 141, 309-323.	2.3	28
72	Evaluation of the Clonal Origin of Multiple Primary Melanomas Using Molecular Profiling. <i>Journal of Investigative Dermatology</i> , 2009, 129, 1972-1982.	0.3	27

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73	Inherited variation at <i>MC1R</i> and <i>ASIP</i> and association with melanoma-specific survival. <i>International Journal of Cancer</i> , 2015, 136, 2659-2667.	2.3	27
74	Lung Cancer Risk in Never-Smokers of European Descent is Associated With Genetic Variation in the 5p15.33 TERT-CLPTM1L Region. <i>Journal of Thoracic Oncology</i> , 2019, 14, 1360-1369.	0.5	27
75	Sun Exposure and Melanoma Survival: A GEM Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 2145-2152.	1.1	26
76	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
77	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
78	Patterns of Persistent DNA Damage Associated with Sun Exposure and the Glutathione <i>S</i> -transferase M1 Genotype in Melanoma Patients. <i>Photochemistry and Photobiology</i> , 2009, 85, 379-386.	1.3	23
79	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
80	Variants in autophagy-related genes and clinical characteristics in melanoma: a population-based study. <i>Cancer Medicine</i> , 2016, 5, 3336-3345.	1.3	23
81	MRI background parenchymal enhancement, breast density and serum hormones in postmenopausal women. <i>International Journal of Cancer</i> , 2018, 143, 823-830.	2.3	23
82	Epithelial-Mesenchymal Transition (EMT) Gene Variants and Epithelial Ovarian Cancer (EOC) Risk. <i>Genetic Epidemiology</i> , 2015, 39, 689-697.	0.6	22
83	Inherited Variation at <i>MC1R</i> and Histological Characteristics of Primary Melanoma. <i>PLoS ONE</i> , 2015, 10, e0119920.	1.1	22
84	Body mass index, weight change, and risk of second primary breast cancer in the <i>WECARE</i> study: influence of estrogen receptor status of the first breast cancer. <i>Cancer Medicine</i> , 2016, 5, 3282-3291.	1.3	22
85	Alcohol consumption and lung cancer risk: A pooled analysis from the International Lung Cancer Consortium and the SYNERGY study. <i>Cancer Epidemiology</i> , 2019, 58, 25-32.	0.8	22
86	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
87	Genetic variants and cognitive functions in patients with brain tumors. <i>Neuro-Oncology</i> , 2019, 21, 1297-1309.	0.6	21
88	Agnostic Pathway/Gene Set Analysis of Genome-Wide Association Data Identifies Associations for Pancreatic Cancer. <i>Journal of the National Cancer Institute</i> , 2019, 111, 557-567.	3.0	21
89	The use of hierarchical models for estimating relative risks of individual genetic variants: An application to a study of melanoma. <i>Statistics in Medicine</i> , 2008, 27, 1973-1992.	0.8	20
90	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

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91	Sun Exposure, Vitamin D Receptor Genetic Variants, and Risk of Breast Cancer in the Agricultural Health Study. <i>Environmental Health Perspectives</i> , 2014, 122, 165-171.	2.8	20
92	Gastrointestinal Stromal Tumors, Somatic Mutations and Candidate Genetic Risk Variants. <i>PLoS ONE</i> , 2013, 8, e62119.	1.1	19
93	GWAS meta-analysis of 16 852 women identifies new susceptibility locus for endometrial cancer. <i>Human Molecular Genetics</i> , 2016, 25, ddw092.	1.4	19
94	Association of Incident Amelanotic Melanoma With Phenotypic Characteristics, <i>MC1R</i> Status, and Prior Amelanotic Melanoma. <i>JAMA Dermatology</i> , 2017, 153, 1026.	2.0	19
95	Molecular analysis of the INK4A and INK4B gene loci in human breast cancer cell lines and primary carcinomas. <i>Cancer Genetics and Cytogenetics</i> , 2001, 125, 131-138.	1.0	18
96	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , 2016, 141, 386-401.	0.6	18
97	Genetic analyses of gynecological disease identify genetic relationships between uterine fibroids and endometrial cancer, and a novel endometrial cancer genetic risk region at the WNT4 1p36.12 locus. <i>Human Genetics</i> , 2021, 140, 1353-1365.	1.8	18
98	Validation of Denaturing High Performance Liquid Chromatography as a Rapid Detection Method for the Identification of Human INK4A Gene Mutations. <i>Journal of Molecular Diagnostics</i> , 2001, 3, 158-163.	1.2	17
99	Including Additional Controls from Public Databases Improves the Power of a Genome-Wide Association Study. <i>Human Heredity</i> , 2011, 72, 21-34.	0.4	17
100	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.	1.4	17
101	Identification of gene expression levels in primary melanoma associated with clinically meaningful characteristics. <i>Melanoma Research</i> , 2018, 28, 380-389.	0.6	17
102	Consortium analysis of gene and gene-folate 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
103	MC1R variants in childhood and adolescent melanoma: a retrospective pooled analysis of a multicentre cohort. <i>The Lancet Child and Adolescent Health</i> , 2019, 3, 332-342.	2.7	16
104	Alterations of cell cycle regulators affecting the RB pathway in nonfamilial retinoblastoma. <i>Human Pathology</i> , 2001, 32, 537-544.	1.1	15
105	Identifying Etiologically Distinct Subtypes of Cancer: A Demonstration Project Involving Breast Cancer. <i>Cancer Medicine</i> , 2015, 4, 1432-1439.	1.3	15
106	Evaluating the ovarian cancer gonadotropin hypothesis: A candidate gene study. <i>Gynecologic Oncology</i> , 2015, 136, 542-548.	0.6	15
107	Identification of lung cancer histology-specific variants applying Bayesian framework variant prioritization approaches within the TRICL and ILCCO consortia. <i>Carcinogenesis</i> , 2015, 36, 1314-1326.	1.3	15
108	Personalized electro-acupuncture versus auricular-acupuncture comparative effectiveness (PEACE): A protocol of a randomized controlled trial for chronic musculoskeletal pain in cancer survivors. <i>Medicine (United States)</i> , 2020, 99, e20085.	0.4	14

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109	Functional polymorphisms in the promoter regions of MMP2 and MMP3 are not associated with melanoma progression. <i>Journal of Negative Results in BioMedicine</i> , 2007, 6, 9.	1.4	13
110	Genome-wide analysis of the role of copy-number variation in pancreatic cancer risk. <i>Frontiers in Genetics</i> , 2014, 5, 29.	1.1	13
111	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
112	Sunburn, sun exposure, and sun sensitivity in the Study of Nevi in Children. <i>Annals of Epidemiology</i> , 2015, 25, 839-843.e4.	0.9	13
113	The interaction between vitamin D receptor polymorphisms and sun exposure around time of diagnosis influences melanoma survival. <i>Pigment Cell and Melanoma Research</i> , 2018, 31, 287-296.	1.5	13
114	Inherited variants affecting RNA editing may contribute to ovarian cancer susceptibility: results from a large-scale collaboration. <i>Oncotarget</i> , 2016, 7, 72381-72394.	0.8	13
115	Exome-Wide Association Study of Endometrial Cancer in a Multiethnic Population. <i>PLoS ONE</i> , 2014, 9, e97045.	1.1	12
116	Investigation of the Effect of MDM2 SNP309 and TP53 Arg72Pro Polymorphisms on the Age of Onset of Cutaneous Melanoma. <i>Journal of Investigative Dermatology</i> , 2012, 132, 1471-1478.	0.3	11
117	Nevus count associations with pigmentary phenotype, histopathological melanoma characteristics and survival from melanoma. <i>International Journal of Cancer</i> , 2016, 139, 1217-1222.	2.3	11
118	Associations of MC1R Genotype and Patient Phenotypes with BRAF and NRAS Mutations in Melanoma. <i>Journal of Investigative Dermatology</i> , 2017, 137, 2588-2598.	0.3	11
119	CYP2D6 phenotype, tamoxifen, and risk of contralateral breast cancer in the WECARE Study. <i>Breast Cancer Research</i> , 2018, 20, 149.	2.2	11
120	Alterations in the retinoblastoma pathway of cell cycle control in parathyroid tumors.. <i>Oncology Reports</i> , 2000, 7, 421-5.	1.2	10
121	Gastrointestinal stromal tumors: a case-only analysis of single nucleotide polymorphisms and somatic mutations. <i>Clinical Sarcoma Research</i> , 2013, 3, 12.	2.3	10
122	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
123	Inherited Genetic Variants Associated with Melanoma BRAF/NRAS Subtypes. <i>Journal of Investigative Dermatology</i> , 2018, 138, 2398-2404.	0.3	9
124	Variants in genes encoding small GTPases and association with epithelial ovarian cancer susceptibility. <i>PLoS ONE</i> , 2018, 13, e0197561.	1.1	9
125	Genome-Wide Association Study for Ovarian Cancer Susceptibility Using Pooled DNA. <i>Twin Research and Human Genetics</i> , 2012, 15, 615-623.	0.3	8
126	Accuracy of Self-reported Smoking Exposure Among Bladder Cancer Patients Undergoing Surveillance at a Tertiary Referral Center. <i>European Urology Focus</i> , 2016, 2, 441-444.	1.6	8



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127	No association between prediagnosis exercise and survival in patients with high-risk primary melanoma: A population-based study. <i>Pigment Cell and Melanoma Research</i> , 2017, 30, 424-427.	1.5	8
128	Interaction of CDKN2A and Sun Exposure in the Etiology of Melanoma in the General Population. <i>Journal of Investigative Dermatology</i> , 2011, 131, 2500-2503.	0.3	7
129	No prognostic value added by vitamin D pathway SNPs to current prognostic system for melanoma survival. <i>PLoS ONE</i> , 2017, 12, e0174234.	1.1	7
130	Association of Known Melanoma Risk Factors with Primary Melanoma of the Scalp and Neck. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 2203-2210.	1.1	6
131	Effects of acupuncture versus cognitive behavioral therapy on brain-derived neurotrophic factor in cancer survivors with insomnia: an exploratory analysis. <i>Acupuncture in Medicine</i> , 2021, 39, 637-645.	0.4	6
132	Association of a Pathway-Specific Genetic Risk Score With Risk of Radiation-Associated Contralateral Breast Cancer. <i>JAMA Network Open</i> , 2019, 2, e1912259.	2.8	5
133	Genome-Wide Gene-Diabetes and Gene-Obesity Interaction Scan in 8,255 Cases and 11,900 Controls from PanScan and PanC4 Consortia. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1784-1791.	1.1	5
134	Genetic Predictors of Response to Acupuncture or Cognitive Behavioral Therapy for Insomnia in Cancer Survivors: An Exploratory Analysis. <i>Journal of Pain and Symptom Management</i> , 2021, 62, e192-e199.	0.6	5
135	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
136	Prospective evaluation of functional brain activity and oxidative damage in breast cancer: changes in task-induced deactivation during a working memory task. <i>Brain Imaging and Behavior</i> , 2021, 15, 1364-1373.	1.1	4
137	Interpretation of Melanoma Risk Feedback in First-Degree Relatives of Melanoma Patients. <i>Journal of Cancer Epidemiology</i> , 2012, 2012, 1-7.	0.5	3
138	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
139	Comparison of community pathologists with expert dermatopathologists evaluating Breslow thickness and histopathologic subtype in a large international population-based study of melanoma. <i>JAAD International</i> , 2021, 4, 25-27.	1.1	3
140	Genotypic and phenotypic characterization of the histoblood group ABO(H) in primary bladder tumors. , 1998, 75, 819.		2
141	Disease-Associated Risk Variants in <i>ANRIL</i> Are Associated with Tumor-Infiltrating Lymphocyte Presence in Primary Melanomas in the Population-Based GEM Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 2309-2316.	1.1	2
142	Characterization of the tobacco glycoprotein surface binding property of heart and skeletal muscle cells. <i>Archives of Toxicology</i> , 1995, 69, 149-159.	1.9	1
143	Evaluation of Alterations in the Tumor Suppressor Genes INK4A and INK4B in Human Bladder Tumors. , 2002, 179, 043-059.		1
144	Inherited Melanoma Risk Variants Associated with Histopathologically Amelanotic Melanoma. <i>Journal of Investigative Dermatology</i> , 2020, 140, 918-922.e7.	0.3	1

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145	Differences in Melanoma Between Canada and New South Wales, Australia: A Population-Based Genes, Environment, and Melanoma (GEM) Study. <i>JID Innovations</i> , 2021, 1, 100002.	1.2	1
146	Association of Melanoma-Risk Variants with Primary Melanoma Tumor Prognostic Characteristics and Melanoma-Specific Survival in the GEM Study. <i>Current Oncology</i> , 2021, 28, 4756-4771.	0.9	1
147	Patterns and sources of information about family melanoma risk among melanoma survivors. <i>Melanoma Management</i> , 2016, 3, 105-111.	0.1	0
148	Relationship of Chromosome Arm 10q Variants to Occurrence of Multiple Primary Melanoma in the Population-Based Genes, Environment, and Melanoma (GEM) Study. <i>Journal of Investigative Dermatology</i> , 2019, 139, 1410-1412.	0.3	0
149	Human genes differ by their UV sensitivity estimated through analysis of UV-induced silent mutations in melanoma. <i>Human Mutation</i> , 2020, 41, 1751-1760.	1.1	0
150	The p.Ser64Leu and p.Pro104Leu missense variants of PALB2 identified in familial pancreatic cancer patients compromise the DNA damage response. <i>Human Mutation</i> , 2021, 42, 150-163.	1.1	0
151	Minimally invasive micro biopsy for genetic profiling of melanocytic lesions: A case series. <i>Journal of the American Academy of Dermatology</i> , 2021, , .	0.6	0