Irene Orlow

List of Publications by Year
in descending order

151	8,575	57719	49868
papers	citations	44 h-index	87 g-index

154	154	154	13753
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Prospective evaluation of functional brain activity and oxidative damage in breast cancer: changes in task-induced deactivation during a working memory task. Brain Imaging and Behavior, 2021, 15, 1364-1373.	1.1	4
2	The p.Ser64Leu and p.Pro104Leu missense variants of PALB2 identified in familial pancreatic cancer patients compromise the DNA damage response. Human Mutation, 2021, 42, 150-163.	1.1	0
3	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. Mutation Research - Reviews in Mutation Research, 2021, 787, 108371.	2.4	45
4	Effects of acupuncture versus cognitive behavioral therapy on brain-derived neurotrophic factor in cancer survivors with insomnia: an exploratory analysis. Acupuncture in Medicine, 2021, 39, 637-645.	0.4	6
5	Differences in Melanoma Between Canada and New South Wales, Australia: A Population-Based Genes, Environment, and Melanoma (GEM) Study. JID Innovations, 2021, 1, 100002.	1.2	1
6	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. Human Genetics, 2021, 140, 1353-1365.	1.8	18
7	Genetic Predictors of Response to Acupuncture or Cognitive Behavioral Therapy for Insomnia in Cancer Survivors: An Exploratory Analysis. Journal of Pain and Symptom Management, 2021, 62, e192-e199.	0.6	5
8	Comparison of community pathologists with expert dermatopathologists evaluating Breslow thickness and histopathologic subtype in a large international population-based study of melanoma. JAAD International, 2021, 4, 25-27.	1.1	3
9	Disease-Associated Risk Variants in ANRIL Are Associated with Tumor-Infiltrating Lymphocyte Presence in Primary Melanomas in the Population-Based GEM Study. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 2309-2316.	1.1	2
10	Association of Melanoma-Risk Variants with Primary Melanoma Tumor Prognostic Characteristics and Melanoma-Specific Survival in the GEM Study. Current Oncology, 2021, 28, 4756-4771.	0.9	1
11	Minimally invasive microbiopsy for genetic profiling of melanocytic lesions: A case series. Journal of the American Academy of Dermatology, 2021, , .	0.6	0
12	Inherited Melanoma Risk Variants Associated with Histopathologically Amelanotic Melanoma. Journal of Investigative Dermatology, 2020, 140, 918-922.e7.	0.3	1
13	A Transcriptome-Wide Association Study Identifies Novel Candidate Susceptibility Genes for Pancreatic Cancer. Journal of the National Cancer Institute, 2020, 112, 1003-1012.	3.0	59
14	Association of Known Melanoma Risk Factors with Primary Melanoma of the Scalp and Neck. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 2203-2210.	1.1	6
15	Personalized electro-acupuncture versus auricular-acupuncture comparative effectiveness (PEACE): A protocol of a randomized controlled trial for chronic musculoskeletal pain in cancer survivors. Medicine (United States), 2020, 99, e20085.	0.4	14
16	Genome-Wide Geneâ€“Diabetes and Geneâ€“Obesity Interaction Scan in 8,255 Cases and 11,900 Controls from PanScan and PanC4 Consortia. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 1784-1791.	1.1	5
17	Human genes differ by their UV sensitivity estimated through analysis of UVâ€induced silent mutations in melanoma. Human Mutation, 2020, 41, 1751-1760.	1.1	0
18	Lung Cancer Risk in Never-Smokers of European Descent is Associated With Genetic Variation in the 5p15.33 TERT-CLPTM1Ll Region. Journal of Thoracic Oncology, 2019, 14, 1360-1369.	0.5	27

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19	Association of a Pathway-Specific Genetic Risk Score With Risk of Radiation-Associated Contralateral Breast Cancer. JAMA Network Open, 2019, 2, e1912259.	2.8	5
20	Genetic variants and cognitive functions in patients with brain tumors. Neuro-Oncology, 2019, 21, 1297-1309.	0.6	21
21	Analysis of Heritability and Genetic Architecture of Pancreatic Cancer: A PanC4 Study. Cancer Epidemiology Biomarkers and Prevention, 2019, 28, 1238-1245.	1.1	48
22	MC1R variants in childhood and adolescent melanoma: a retrospective pooled analysis of a multicentre cohort. The Lancet Child and Adolescent Health, 2019, 3, 332-342.	2.7	16
23	Relationship of Chromosome Arm 10q Variants toÂOccurrence of Multiple Primary Melanoma in theÂPopulation-Based Genes, Environment, andÂMelanoma (GEM) Study. Journal of Investigative Dermatology, 2019, 139, 1410-1412.	0.3	0
24	Agnostic Pathway/Gene Set Analysis of Genome-Wide Association Data Identifies Associations for Pancreatic Cancer. Journal of the National Cancer Institute, 2019, 111, 557-567.	3.0	21
25	Alcohol consumption and lung cancer risk: A pooled analysis from the International Lung Cancer Consortium and the SYNERGY study. Cancer Epidemiology, 2019, 58, 25-32.	0.8	22
26	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. Nature Communications, 2018, 9, 556.	5.8	188
27	MRI background parenchymal enhancement, breast density and serum hormones in postmenopausal women. International Journal of Cancer, 2018, 143, 823-830.	2.3	23
28	The interaction between vitamin D receptor polymorphisms and sun exposure around time of diagnosis influences melanoma survival. Pigment Cell and Melanoma Research, 2018, 31, 287-296.	1.5	13
29	Contralateral breast cancers: Independent cancers or metastases?. International Journal of Cancer, 2018, 142, 347-356.	2.3	37
30	CYP2D6 phenotype, tamoxifen, and risk of contralateral breast cancer in the WECARE Study. Breast Cancer Research, 2018, 20, 149.	2.2	11
31	Identification of gene expression levels in primary melanoma associated with clinically meaningful characteristics. Melanoma Research, 2018, 28, 380-389.	0.6	17
32	Inherited Genetic Variants Associated with Melanoma BRAF/NRAS Subtypes. Journal of Investigative Dermatology, 2018, 138, 2398-2404.	0.3	9
33	Variants in genes encoding small GTPases and association with epithelial ovarian cancer susceptibility. PLoS ONE, 2018, 13, e0197561.	1.1	9
34	Identification of nine new susceptibility loci for endometrial cancer. Nature Communications, 2018, 9, 3166.	5.8	178
35	rs495139 in the TYMS-ENOSF1 Region and Risk of Ovarian Carcinoma of Mucinous Histology. International Journal of Molecular Sciences, 2018, 19, 2473.	1.8	3
36	Alcohol and lung cancer risk among never smokers: A pooled analysis from the international lung cancer consortium and the SYNERGY study. International Journal of Cancer, 2017, 140, 1976-1984.	2.3	35

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37	No association between prediagnosis exercise and survival in patients with highâ€risk primary melanoma: A populationâ€based study. Pigment Cell and Melanoma Research, 2017, 30, 424-427.	1.5	8
38	Menstrual and reproductive factors and lung cancer risk: A pooled analysis from the international lung cancer consortium. International Journal of Cancer, 2017, 141, 309-323.	2.3	28
39	Associations of MC1R Genotype and Patient Phenotypes with BRAF and NRAS Mutations in Melanoma. Journal of Investigative Dermatology, 2017, 137, 2588-2598.	0.3	11
40	Association of Incident Amelanotic Melanoma With Phenotypic Characteristics, MC1R Status, and Prior Amelanotic Melanoma. JAMA Dermatology, 2017, 153, 1026.	2.0	19
41	The oral microbiota in patients with pancreatic cancer, patients with IPMNs, and controls: a pilot study. Cancer Causes and Control, 2017, 28, 959-969.	0.8	69
42	No prognostic value added by vitamin D pathway SNPs to current prognostic system for melanoma survival. PLoS ONE, 2017, 12, e0174234.	1.1	7
43	Exome genotyping arrays to identify rare and low frequency variants associated with epithelial ovarian cancer risk. Human Molecular Genetics, 2016, 25, 3600-3612.	1.4	17
44	Nevus count associations with pigmentary phenotype, histopathological melanoma characteristics and survival from melanoma. International Journal of Cancer, 2016, 139, 1217-1222.	2.3	11
45	COMT,BDNF, andDTNBP1polymorphisms and cognitive functions in patients with brain tumors. Neuro-Oncology, 2016, 18, 1425-1433.	0.6	45
46	Body mass index, weight change, and risk of second primary breast cancer in the <scp>WECARE</scp> study: influence of estrogen receptor status of the first breast cancer. Cancer Medicine, 2016, 5, 3282-3291.	1.3	22
47	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. Nature Communications, 2016, 7, 11843.	5.8	86
48	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breastâ€“ovarian cancer susceptibility locus. Nature Communications, 2016, 7, 12675.	5.8	78
49	Variants in autophagyâ€related genes and clinical characteristics in melanoma: a populationâ€based study. Cancer Medicine, 2016, 5, 3336-3345.	1.3	23
50	Patterns and sources of information about family melanoma risk among melanoma survivors. Melanoma Management, 2016, 3, 105-111.	0.1	0
51	Accuracy of Self-reported Smoking Exposure Among Bladder Cancer Patients Undergoing Surveillance at a Tertiary Referral Center. European Urology Focus, 2016, 2, 441-444.	1.6	8
52	GWAS meta-analysis of 16 852 women identifies new susceptibility locus for endometrial cancer. Human Molecular Genetics, 2016, 25, ddw092.	1.4	19
53	Association of Interferon Regulatory Factor-4 Polymorphism rs12203592 With Divergent Melanoma Pathways. Journal of the National Cancer Institute, 2016, 108, djw004.	3.0	28
54	Investigation of Exomic Variants Associated with Overall Survival in Ovarian Cancer. Cancer Epidemiology Biomarkers and Prevention, 2016, 25, 446-454.	1.1	9

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55	Vitamin D receptor polymorphisms and survival in patients with cutaneous melanoma: a population-based study. Carcinogenesis, 2016, 37, 30-38.	1.3	54
56	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. Gynecologic Oncology, 2016, 141, 386-401.	0.6	18
57	Assessment of variation in immunosuppressive pathway genes reveals TGFBR2 to be associated with risk of clear cell ovarian cancer. Oncotarget, 2016, 7, 69097-69110.	0.8	5
58	Inherited variants affecting RNA editing may contribute to ovarian cancer susceptibility: results from a large-scale collaboration. Oncotarget, 2016, 7, 72381-72394.	0.8	13
59	Identifying Etiologically Distinct Subâ€Types of Cancer: A Demonstration Project Involving Breast Cancer. Cancer Medicine, 2015, 4, 1432-1439.	1.3	15
60	Inherited variation at MC1R and ASIP and association with melanomaâ€specific survival. International Journal of Cancer, 2015, 136, 2659-2667.	2.3	27
61	Epithelialâ€Mesenchymal Transition (EMT) Gene Variants and Epithelial Ovarian Cancer (EOC) Risk. Genetic Epidemiology, 2015, 39, 689-697.	0.6	22
62	Vitamin D Metabolic Pathway Genes and Pancreatic Cancer Risk. PLoS ONE, 2015, 10, e0117574.	1.1	29
63	Inherited Variation at MC1R and Histological Characteristics of Primary Melanoma. PLoS ONE, 2015, 10, e0119920.	1.1	22
64	Common Genetic Variation In Cellular Transport Genes and Epithelial Ovarian Cancer (EOC) Risk. PLoS ONE, 2015, 10, e0128106.	1.1	44
65	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. American Journal of Human Genetics, 2015, 96, 487-497.	2.6	101
66	Association BetweenNRASandBRAFMutational Status and Melanoma-Specific Survival Among Patients With Higher-Risk Primary Melanoma. JAMA Oncology, 2015, 1, 359.	3.4	164
67	Cell-type-specific enrichment of risk-associated regulatory elements at ovarian cancer susceptibility loci. Human Molecular Genetics, 2015, 24, 3595-3607.	1.4	40
68	Genetic factors associated with naevus count and dermoscopic patterns: preliminary results from the Study of Nevi in Children (<scp>SONIC</scp>). British Journal of Dermatology, 2015, 172, 1081-1089.	1.4	31
69	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. Nature Genetics, 2015, 47, 164-171.	9.4	221
70	Genome-wide significant risk associations for mucinous ovarian carcinoma. Nature Genetics, 2015, 47, 888-897.	9.4	78
71	Network-Based Integration of GWAS and Gene Expression Identifies a HOX-Centric Network Associated with Serous Ovarian Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1574-1584.	1.1	28
72	Genome-wide Analysis Identifies Novel Loci Associated with Ovarian Cancer Outcomes: Findings from the Ovarian Cancer Association Consortium. Clinical Cancer Research, 2015, 21, 5264-5276.	3.2	33

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73	Common variation at 2p13.3, 3q29, 7p13 and 17q25.1 associated with susceptibility to pancreatic cancer. Nature Genetics, 2015, 47, 911-916.	9.4	224
74	Evaluating the ovarian cancer gonadotropin hypothesis: A candidate gene study. Gynecologic Oncology, 2015, 136, 542-548.	0.6	15
75	Inherited Genetic Variants Associated with Occurrence of Multiple Primary Melanoma. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 992-997.	1.1	36
76	Cis-eQTL analysis and functional validation of candidate susceptibility genes for high-grade serous ovarian cancer. Nature Communications, 2015, 6, 8234.	5.8	63
77	Identification of lung cancer histology-specific variants applying Bayesian framework variant prioritization approaches within the TRICL and ILCCO consortia. Carcinogenesis, 2015, 36, 1314-1326.	1.3	15
78	Common variants at theCHEK2gene locus and risk of epithelial ovarian cancer. Carcinogenesis, 2015, 36, 1341-1353.	1.3	24
79	Sunburn, sun exposure, and sun sensitivity in the Study of Nevi in Children. Annals of Epidemiology, 2015, 25, 839-843.e4.	0.9	13
80	Cannabis smoking and lung cancer risk: Pooled analysis in the <scp>I</scp>nternational <scp>L</scp>ung <scp>C</scp>ancer <scp>C</scp>onsortium. International Journal of Cancer, 2015, 136, 894-903.	2.3	131
81	Common Genetic Variation in Circadian Rhythm Genes and Risk of Epithelial Ovarian Cancer (EOC). Journal of Genetics and Genome Research, 2015, 2, .	0.3	25
82	Exome-Wide Association Study of Endometrial Cancer in a Multiethnic Population. PLoS ONE, 2014, 9, e97045.	1.1	12
83	Genome-wide analysis of the role of copy-number variation in pancreatic cancer risk. Frontiers in Genetics, 2014, 5, 29.	1.1	13
84	Sun Exposure and Melanoma Survival: A GEM Study. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 2145-2152.	1.1	26
85	Comparison of Clinicopathologic Features and Survival of Histopathologically Amelanotic and Pigmented Melanomas. JAMA Dermatology, 2014, 150, 1306.	2.0	142
86	Variation in NF-ÎºB Signaling Pathways and Survival in Invasive Epithelial Ovarian Cancer. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 1421-1427.	1.1	13
87	<scp>MITF</scp> E318K's effect on melanoma risk independent of, but modified by, other risk factors. Pigment Cell and Melanoma Research, 2014, 27, 485-488.	1.5	35
88	Risk of Ovarian Cancer and the NF-ÎºB Pathway: Genetic Association with IL1A and TNFSF10. Cancer Research, 2014, 74, 852-861.	0.4	48
89	Sun Exposure, Vitamin D Receptor Genetic Variants, and Risk of Breast Cancer in the Agricultural Health Study. Environmental Health Perspectives, 2014, 122, 165-171.	2.8	20
90	Large-Scale Evaluation of Common Variation in Regulatory T Cellâ€“Related Genes and Ovarian Cancer Outcome. Cancer Immunology Research, 2014, 2, 332-340.	1.6	21

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91	Genome-wide association study of endometrial cancer in E2C2. Human Genetics, 2014, 133, 211-224.	1.8	42
92	Genome-wide association study of subtype-specific epithelial ovarian cancer risk alleles using pooled DNA. Human Genetics, 2014, 133, 481-497.	1.8	23
93	Consortium analysis of gene and geneâ€"folate interactions in purine and pyrimidine metabolism pathways with ovarian carcinoma risk. Molecular Nutrition and Food Research, 2014, 58, 2023-2035.	1.5	16
94	APOE polymorphisms and cognitive functions in patients with brain tumors. Neurology, 2014, 83, 320-327.	1.5	49
95	GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. Nature Genetics, 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. Nature Genetics, 2013, 45, 371-384.	9.4	493
97	Tumor-Infiltrating Lymphocyte Grade in Primary Melanomas Is Independently Associated With Melanoma-Specific Survival in the Population-Based Genes, Environment and Melanoma Study. Journal of Clinical Oncology, 2013, 31, 4252-4259.	0.8	232
98	Gastrointestinal stromal tumors: a case-only analysis of single nucleotide polymorphisms and somatic mutations. Clinical Sarcoma Research, 2013, 3, 12.	2.3	10
99	Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. Nature Communications, 2013, 4, 1628.	5.8	144
100	Analysis of Over 10,000 Cases Finds No Association between Previously Reported Candidate Polymorphisms and Ovarian Cancer Outcome. Cancer Epidemiology Biomarkers and Prevention, 2013, 22, 987-992.	1.1	20
101	Identification and molecular characterization of a new ovarian cancer susceptibility locus at 17q21.31. Nature Communications, 2013, 4, 1627.	5.8	98
102	Gastrointestinal Stromal Tumors, Somatic Mutations and Candidate Genetic Risk Variants. PLoS ONE, 2013, 8, e62119.	1.1	19
103	Investigation of the Effect of MDM2 SNP309 and TP53 Arg72Pro Polymorphisms on the Age of Onset of Cutaneous Melanoma. Journal of Investigative Dermatology, 2012, 132, 1471-1478.	0.3	11
104	Asthma and lung cancer risk: a systematic investigation by the International Lung Cancer Consortium. Carcinogenesis, 2012, 33, 587-597.	1.3	69
105	Risk of Non-Melanoma Cancers in First-Degree Relatives of CDKN2A Mutation Carriers. Journal of the National Cancer Institute, 2012, 104, 953-956.	3.0	42
106	Interpretation of Melanoma Risk Feedback in First-Degree Relatives of Melanoma Patients. Journal of Cancer Epidemiology, 2012, 2012, 1-7.	0.5	3
107	Genome-Wide Association Study for Ovarian Cancer Susceptibility Using Pooled DNA. Twin Research and Human Genetics, 2012, 15, 615-623.	0.3	8
108	Previous Lung Diseases and Lung Cancer Risk: A Pooled Analysis From the International Lung Cancer Consortium. American Journal of Epidemiology, 2012, 176, 573-585.	1.6	160

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109	A Replication Study and Genome-Wide Scan of Single-Nucleotide Polymorphisms Associated with Pancreatic Cancer Risk and Overall Survival. Clinical Cancer Research, 2012, 18, 3942-3951.	3.2	40
110	Vitamin D Receptor Gene Haplotypes and Polymorphisms and Risk of Breast Cancer: A Nested Caseâ€“Control Study. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 1856-1867.	1.1	35
111	Increased risk of lung cancer in individuals with a family history of the disease: A pooled analysis from the International Lung Cancer Consortium. European Journal of Cancer, 2012, 48, 1957-1968.	1.3	143
112	Clinicopathologic Features of Incident and Subsequent Tumors in Patients with Multiple Primary Cutaneous Melanomas. Annals of Surgical Oncology, 2012, 19, 1024-1033.	0.7	45
113	Vitamin D receptor polymorphisms in patients with cutaneous melanoma. International Journal of Cancer, 2012, 130, 405-418.	2.3	61
114	Sun exposure, vitamin D receptor polymorphisms FokI and BsmI and risk of multiple primary melanoma. Cancer Epidemiology, 2011, 35, e105-e110.	0.8	28
115	Functional Polymorphisms in the TERT Promoter Are Associated with Risk of Serous Epithelial Ovarian and Breast Cancers. PLoS ONE, 2011, 6, e24987.	1.1	48
116	Aspirin and NSAID use and lung cancer risk: a pooled analysis in the International Lung Cancer Consortium (ILCCO). Cancer Causes and Control, 2011, 22, 1709-1720.	0.8	47
117	Including Additional Controls from Public Databases Improves the Power of a Genome-Wide Association Study. Human Heredity, 2011, 72, 21-34.	0.4	17
118	Interaction of CDKN2A and Sun Exposure in the Etiology of Melanoma in the General Population. Journal of Investigative Dermatology, 2011, 131, 2500-2503.	0.3	7
119	The Obesity-Associated Polymorphisms FTO rs9939609 and MC4R rs17782313 and Endometrial Cancer Risk in Non-Hispanic White Women. PLoS ONE, 2011, 6, e16756.	1.1	58
120	A metastasis or a second independent cancer? Evaluating the clonal origin of tumors using array copy number data. Statistics in Medicine, 2010, 29, 1608-1621.	0.8	46
121	Associations of Cumulative Sun Exposure and Phenotypic Characteristics with Histologic Solar Elastosis. Cancer Epidemiology Biomarkers and Prevention, 2010, 19, 2932-2941.	1.1	45
122	Analysis of Genetic Variants in Never-Smokers with Lung Cancer Facilitated by an Internet-Based Blood Collection Protocol: A Preliminary Report. Clinical Cancer Research, 2010, 16, 755-763.	3.2	82
123	Genomic and Mutational Profiling to Assess Clonal Relationships Between Multiple Nonâ€“Small Cell Lung Cancers. Clinical Cancer Research, 2009, 15, 5184-5190.	3.2	151
124	Evaluation of the Clonal Origin of Multiple Primary Melanomas Using Molecular Profiling. Journal of Investigative Dermatology, 2009, 129, 1972-1982.	0.3	27
125	Patterns of Persistent DNA Damage Associated with Sun Exposure and the Glutathione Sâ€transferase M1 Genotype in Melanoma Patients. Photochemistry and Photobiology, 2009, 85, 379-386.	1.3	23
126	Variants in hormone biosynthesis genes and risk of endometrial cancer. Cancer Causes and Control, 2008, 19, 955-963.	0.8	29

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127	The use of hierarchical models for estimating relative risks of individual genetic variants: An application to a study of melanoma. Statistics in Medicine, 2008, 27, 1973-1992.	0.8	20
128	Phase II Study of Extended-Dose Temozolomide in Patients With Melanoma. Journal of Clinical Oncology, 2008, 26, 2299-2304.	0.8	66
129	DNA Damage and Repair Capacity in Patients With Lung Cancer: Prediction of Multiple Primary Tumors. Journal of Clinical Oncology, 2008, 26, 3560-3566.	0.8	56
130	Variants in Estrogen Biosynthesis Genes, Sex Steroid Hormone Levels, and Endometrial Cancer: A HuGE Review. American Journal of Epidemiology, 2007, 165, 235-245.	1.6	102
131	Matrix Metalloproteinase-9 (MMP-9) polymorphisms in patients with cutaneous malignant melanoma. BMC Medical Genetics, 2007, 8, 10.	2.1	44
132	Functional polymorphisms in the promoter regions of MMP2 and MMP3 are not associated with melanoma progression. Journal of Negative Results in BioMedicine, 2007, 6, 9.	1.4	13
133	CDKN2A Germline Mutations in Individuals with Cutaneous Malignant Melanoma. Journal of Investigative Dermatology, 2007, 127, 1234-1243.	0.3	50
134	Allergies, variants in IL-4 and IL-4RÎ± genes, and risk of pancreatic cancer. Cancer Detection and Prevention, 2007, 31, 345-351.	2.1	58
135	The Prevalence of CDKN2A Germ-Line Mutations and Relative Risk for Cutaneous Malignant Melanoma: An International Population-Based Study. Cancer Epidemiology Biomarkers and Prevention, 2006, 15, 1520-1525.	1.1	105
136	Lifetime Risk of Melanoma in CDKN2A Mutation Carriers in a Population-Based Sample. Journal of the National Cancer Institute, 2005, 97, 1507-1515.	3.0	200
137	Association Between Aryl Hydrocarbon Receptor Genotype and Survival in Soft Tissue Sarcoma. Journal of Clinical Oncology, 2004, 22, 3997-4001.	0.8	28
138	Evaluation of Alterations in the Tumor Suppressor Genes INK4A and INK4B in Human Bladder Tumors. , 2002, 179, 043-059.		1
139	Alterations of cell cycle regulators affecting the RB pathway in nonfamilial retinoblastoma. Human Pathology, 2001, 32, 537-544.	1.1	15
140	Validation of Denaturing High Performance Liquid Chromatography as a Rapid Detection Method for the Identification of Human INK4A Gene Mutations. Journal of Molecular Diagnostics, 2001, 3, 158-163.	1.2	17
141	Molecular analyses of the mitotic checkpoint componentshsMAD2, hBUB1 andhBUB3 in human cancer. International Journal of Cancer, 2001, 95, 223-227.	2.3	92
142	Molecular analysis of the INK4A and INK4B gene loci in human breast cancer cell lines and primary carcinomas. Cancer Genetics and Cytogenetics, 2001, 125, 131-138.	1.0	18
143	Alterations in the retinoblastoma pathway of cell cycle control in parathyroid tumors.. Oncology Reports, 2000, 7, 421-5.	1.2	10
144	Prognostic Significance of Transcription Factor E2F-1 in Bladder Cancer: Genotypic and Phenotypic Characterization. Journal of the National Cancer Institute, 1999, 91, 874-881.	3.0	54

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145	Deletions of the INK4A Gene in Superficial Bladder Tumors. American Journal of Pathology, 1999, 155, 105-113.	1.9	121
146	Deletions of the INK4A Gene Occur in Malignant Peripheral Nerve Sheath Tumors but not in Neurofibromas. American Journal of Pathology, 1999, 155, 1855-1860.	1.9	161
147	Genotypic and phenotypic characterization of the histoblood group ABO(H) in primary bladder tumors. , 1998, 75, 819-824.		31
148	The Ink4a Tumor Suppressor Gene Product, p19Arf, Interacts with MDM2 and Neutralizes MDM2's Inhibition of p53. Cell, 1998, 92, 713-723.	13.5	1,412
149	Genotypic and phenotypic characterization of the histoblood group ABO(H) in primary bladder tumors. International Journal of Cancer, 1998, 75, 819-824.	2.3	2
150	Microsatellite instability and deletion analysis of chromosome 10 in human prostate cancer. , 1996, 69, 110-113.		34
151	Characterization of the tobacco glycoprotein surface binding property of heart and skeletal muscle cells. Archives of Toxicology, 1995, 69, 149-159.	1.9	1