

Julia A Newton-Bishop

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

175 papers	8,609 citations	50 h-index	87 g-index
202 ext. papers	9,994 ext. citations	8.3 avg, IF	5.06 L-index

#	Paper	IF	Citations
175	Cross-cohort gut microbiome associations with immune checkpoint inhibitor response in advanced melanoma.. <i>Nature Medicine</i> , 2022 ,	50.5	14
174	ROR2 has a protective role in melanoma by inhibiting Akt activity, cell-cycle progression, and proliferation. <i>Journal of Biomedical Science</i> , 2021 , 28, 76	13.3	3
173	Tumour gene expression signature in primary melanoma predicts long-term outcomes. <i>Nature Communications</i> , 2021 , 12, 1137	17.4	5
172	MX2 mediates establishment of interferon response profile, regulates XAF1, and can sensitize melanoma cells to targeted therapy. <i>Cancer Medicine</i> , 2021 , 10, 2840-2854	4.8	0
171	Birth cohort-specific trends of sun-related behaviors among individuals from an international consortium of melanoma-prone families. <i>BMC Public Health</i> , 2021 , 21, 692	4.1	2
170	Transcriptional signatures underlying dynamic phenotypic switching and novel disease biomarkers in a linear cellular model of melanoma progression. <i>Neoplasia</i> , 2021 , 23, 439-455	6.4	0
169	Inherited duplications of PPP2R3B predispose to nevi and melanoma via a C21orf91-driven proliferative phenotype. <i>Genetics in Medicine</i> , 2021 , 23, 1636-1647	8.1	0
168	Multiple Primary Melanoma Incidence Trends Over Five Decades: A Nationwide Population-Based Study. <i>Journal of the National Cancer Institute</i> , 2021 , 113, 318-328	9.7	6
167	Cooperation between melanoma cell states promotes metastasis through heterotypic cluster formation. <i>Developmental Cell</i> , 2021 , 56, 2808-2825.e10	10.2	4
166	Factors Affecting Sentinel Node Metastasis in Thin (T1) Cutaneous Melanomas: Development and External Validation of a Predictive Nomogram. <i>Journal of Clinical Oncology</i> , 2020 , 38, 1591-1601	2.2	26
165	Genome-wide association meta-analyses combining multiple risk phenotypes provide insights into the genetic architecture of cutaneous melanoma susceptibility. <i>Nature Genetics</i> , 2020 , 52, 494-504	36.3	39
164	MX 2 is a novel regulator of cell cycle in melanoma cells. <i>Pigment Cell and Melanoma Research</i> , 2020 , 33, 446-457	4.5	6
163	Study of the Female Sex Survival Advantage in Melanoma-A Focus on X-Linked Epigenetic Regulators and Immune Responses in Two Cohorts. <i>Cancers</i> , 2020 , 12,	6.6	7
162	Gene co-expression and histone modification signatures are associated with melanoma progression, epithelial-to-mesenchymal transition, and metastasis. <i>Clinical Epigenetics</i> , 2020 , 12, 127	7.7	2
161	MC1R variants and cutaneous melanoma risk according to histological type, body site, and Breslow thickness: a pooled analysis from the M-SKIP project. <i>Melanoma Research</i> , 2020 , 30, 500-510	3.3	2
160	Application of Circulating Cell-Free Tumor DNA Profiles for Therapeutic Monitoring and Outcome Prediction in Genetically Heterogeneous Metastatic Melanoma. <i>JCO Precision Oncology</i> , 2020 , 3,	3.6	10
159	Environmental Exposures Such as Smoking and Low Vitamin D Are Predictive of Poor Outcome in Cutaneous Melanoma rather than Other Deprivation Measures. <i>Journal of Investigative Dermatology</i> , 2020 , 140, 327-337.e2	4.3	7

158	Gain-of-Function Genetic Alterations of G9a Drive Oncogenesis. <i>Cancer Discovery</i> , 2020 , 10, 980-997	24.4	17
157	Transcriptomic Analysis Reveals Prognostic Molecular Signatures of Stage I Melanoma. <i>Clinical Cancer Research</i> , 2019 , 25, 7424-7435	12.9	16
156	High-Resolution Copy Number Patterns From Clinically Relevant FFPE Material. <i>Scientific Reports</i> , 2019 , 9, 8908	4.9	4
155	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	14.5	8
154	Genetic and Environmental Determinants of Immune Response to Cutaneous Melanoma. <i>Cancer Research</i> , 2019 , 79, 2684-2696	10.1	31
153	Estimating CDKN2A mutation carrier probability among global familial melanoma cases using GenoMELPREDICT. <i>Journal of the American Academy of Dermatology</i> , 2019 , 81, 386-394	4.5	9
152	ctDNA as a noninvasive monitoring tool in metastatic melanoma.. <i>Journal of Clinical Oncology</i> , 2019 , 37, 9548-9548	2.2	
151	Vitamin D-VDR Signaling Inhibits Wnt/ECatenin-Mediated Melanoma Progression and Promotes Antitumor Immunity. <i>Cancer Research</i> , 2019 , 79, 5986-5998	10.1	23
150	Association of the POT1 Germline Missense Variant p.I78T With Familial Melanoma. <i>JAMA Dermatology</i> , 2019 , 155, 604-609	5.1	15
149	25-hydroxyvitamin D serum levels in patients with high risk resected melanoma treated in an adjuvant bevacizumab trial. <i>British Journal of Cancer</i> , 2018 , 119, 793-800	8.7	8
148	Amelanotic melanoma. <i>BMJ, The</i> , 2018 , 360, k826	5.9	5
147	variants as melanoma risk factors independent of at-risk phenotypic characteristics: a pooled analysis from the M-SKIP project. <i>Cancer Management and Research</i> , 2018 , 10, 1143-1154	3.6	32
146	Assessing the Incremental Contribution of Common Genomic Variants to Melanoma Risk Prediction in Two Population-Based Studies. <i>Journal of Investigative Dermatology</i> , 2018 , 138, 2617-2624	4.3	36
145	ECatenin-mediated immune evasion pathway frequently operates in primary cutaneous melanomas. <i>Journal of Clinical Investigation</i> , 2018 , 128, 2048-2063	15.9	46
144	Comparative genomics reveals that loss of lunatic fringe (LFNG) promotes melanoma metastasis. <i>Molecular Oncology</i> , 2018 , 12, 239-255	7.9	12
143	Novel pleiotropic risk loci for melanoma and nevus density implicate multiple biological pathways. <i>Nature Communications</i> , 2018 , 9, 4774	17.4	47
142	Comprehensive Study of the Clinical Phenotype of Germline BAP1 Variant-Carrying Families Worldwide. <i>Journal of the National Cancer Institute</i> , 2018 , 110, 1328-1341	9.7	97
141	Somatic cancer genetics in the UK: real-world data from phase I of the Cancer Research UK Stratified Medicine Programme. <i>ESMO Open</i> , 2018 , 3, e000408	6	2

140	A population-based analysis of germline BAP1 mutations in melanoma. <i>Human Molecular Genetics</i> , 2017 , 26, 717-728	5.6	24
139	Functional characterization of a multi-cancer risk locus on chr5p15.33 reveals regulation of TERT by ZNF148. <i>Nature Communications</i> , 2017 , 8, 15034	17.4	26
138	Germline /P16INK4A mutations contribute to genetic determinism of sarcoma. <i>Journal of Medical Genetics</i> , 2017 , 54, 607-612	5.8	14
137	Rare Variant, Gene-Based Association Study of Hereditary Melanoma Using Whole-Exome Sequencing. <i>Journal of the National Cancer Institute</i> , 2017 , 109,	9.7	16
136	Tests to assist in the staging of cutaneous melanoma: a generic protocol. <i>The Cochrane Library</i> , 2017 ,	5.2	1
135	Germline Variation at CDKN2A and Associations with Nevus Phenotypes among Members of Melanoma Families. <i>Journal of Investigative Dermatology</i> , 2017 , 137, 2606-2612	4.3	10
134	A common intronic variant of PARP1 confers melanoma risk and mediates melanocyte growth via regulation of MITF. <i>Nature Genetics</i> , 2017 , 49, 1326-1335	36.3	36
133	Identification of a gene signature for discriminating metastatic from primary melanoma using a molecular interaction network approach. <i>Scientific Reports</i> , 2017 , 7, 17314	4.9	22
132	Which symptoms are linked to a delayed presentation among melanoma patients? A retrospective study. <i>BMC Cancer</i> , 2017 , 17, 5	4.8	4
131	Genomic expression differences between cutaneous cells from red hair color individuals and black hair color individuals based on bioinformatic analysis. <i>Oncotarget</i> , 2017 , 8, 11589-11599	3.3	5
130	Association of Melanocortin-1 Receptor Variants with Pigmentary Traits in Humans: A Pooled Analysis from the M-Skip Project. <i>Journal of Investigative Dermatology</i> , 2016 , 136, 1914-1917	4.3	12
129	The CDKN2A/p16(INK) (4a) 5'RTR sequence and translational regulation: impact of novel variants predisposing to melanoma. <i>Pigment Cell and Melanoma Research</i> , 2016 , 29, 210-21	4.5	8
128	Consensus of Melanoma Gene Expression Subtypes Converges on Biological Entities. <i>Journal of Investigative Dermatology</i> , 2016 , 136, 2502-2505	4.3	18
127	Wide versus narrow excision margins for high-risk, primary cutaneous melanomas: long-term follow-up of survival in a randomised trial. <i>Lancet Oncology</i> , 2016 , 17, 184-192	21.7	82
126	Phenotypic and Histopathological Tumor Characteristics According to CDKN2A Mutation Status among Affected Members of Melanoma Families. <i>Journal of Investigative Dermatology</i> , 2016 , 136, 1066-1069	4.3	9
125	Loss-of-Function Mutations in the Cell-Cycle Control Gene CDKN2A Impact on Glucose Homeostasis in Humans. <i>Diabetes</i> , 2016 , 65, 527-33	0.9	28
124	Germline TERT promoter mutations are rare in familial melanoma. <i>Familial Cancer</i> , 2016 , 15, 139-44	3	34
123	Psychosocial, clinical and demographic features related to worry in patients with melanoma. <i>Melanoma Research</i> , 2016 , 26, 497-504	3.3	6

122	Diagnosis and treatment of melanoma. European consensus-based interdisciplinary guideline - Update 2016. <i>European Journal of Cancer</i> , 2016 , 63, 201-17	7.5	265
121	Vitamin D, vitamin A, the primary melanoma transcriptome and survival. <i>British Journal of Dermatology</i> , 2016 , 175 Suppl 2, 30-34	4	6
120	Histopathology of melanocytic lesions in a family with an inherited BAP1 mutation. <i>Journal of Cutaneous Pathology</i> , 2016 , 43, 287-9	1.7	8
119	The clinicopathological and gene expression patterns associated with ulceration of primary melanoma. <i>Pigment Cell and Melanoma Research</i> , 2015 , 28, 94-104	4.5	23
118	Nonsense mutations in the shelterin complex genes ACD and TERF2IP in familial melanoma. <i>Journal of the National Cancer Institute</i> , 2015 , 107,	9.7	102
117	Genome-wide meta-analysis identifies five new susceptibility loci for cutaneous malignant melanoma. <i>Nature Genetics</i> , 2015 , 47, 987-995	36.3	162
116	Fine mapping of genetic susceptibility loci for melanoma reveals a mixture of single variant and multiple variant regions. <i>International Journal of Cancer</i> , 2015 , 136, 1351-60	7.5	26
115	Regressing Eruptive Disseminated Spitz Nevi. <i>Pediatric Dermatology</i> , 2015 , 32, e181-3	1.9	5
114	Melanoma: summary of NICE guidance. <i>BMJ, The</i> , 2015 , 351, h3708	5.9	16
113	Prognostic Significance of Promoter Hypermethylation and Diminished Gene Expression of SYNPO2 in Melanoma. <i>Journal of Investigative Dermatology</i> , 2015 , 135, 2328-2331	4.3	10
112	MC1R variants increased the risk of sporadic cutaneous melanoma in darker-pigmented Caucasians: a pooled-analysis from the M-SKIP project. <i>International Journal of Cancer</i> , 2015 , 136, 618-31	7.5	67
111	25-Hydroxyvitamin D2 /D3 levels and factors associated with systemic inflammation and melanoma survival in the Leeds Melanoma Cohort. <i>International Journal of Cancer</i> , 2015 , 136, 2890-9	7.5	42
110	AuthorsReply to: High naevus counts confer a favourable prognosis in patients with melanoma. <i>International Journal of Cancer</i> , 2015 , 137, 3008-9	7.5	
109	High nevus counts confer a favorable prognosis in melanoma patients. <i>International Journal of Cancer</i> , 2015 , 137, 1691-8	7.5	27
108	Development and validation of a melanoma risk score based on pooled data from 16 case-control studies. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 817-24	4	18
107	Histologic features of melanoma associated with CDKN2A genotype. <i>Journal of the American Academy of Dermatology</i> , 2015 , 72, 496-507.e7	4.5	16
106	MGMT promoter methylation is associated with temozolomide response and prolonged progression-free survival in disseminated cutaneous melanoma. <i>International Journal of Cancer</i> , 2015 , 136, 2844-53	7.5	39
105	Independent replication of a melanoma subtype gene signature and evaluation of its prognostic value and biological correlates in a population cohort. <i>Oncotarget</i> , 2015 , 6, 11683-93	3.3	34

104	POT1 loss-of-function variants predispose to familial melanoma. <i>Nature Genetics</i> , 2014 , 46, 478-481	36.3	241
103	Primary melanoma tumors from CDKN2A mutation carriers do not belong to a distinct molecular subclass. <i>Journal of Investigative Dermatology</i> , 2014 , 134, 3000-3003	4.3	5
102	Identification of a melanoma susceptibility locus and somatic mutation in TET2. <i>Carcinogenesis</i> , 2014 , 35, 2097-101	4.6	38
101	Somatic BRAF and NRAS mutations in familial melanomas with known germline CDKN2A status: a GenoMEL study. <i>Journal of Investigative Dermatology</i> , 2014 , 134, 287-290	4.3	15
100	Inherited variation in the PARP1 gene and survival from melanoma. <i>International Journal of Cancer</i> , 2014 , 135, 1625-33	7.5	22
99	An inherited variant in the gene coding for vitamin D-binding protein and survival from cutaneous melanoma: a BioGenoMEL study. <i>Pigment Cell and Melanoma Research</i> , 2014 , 27, 234-43	4.5	21
98	Prevalence and predictors of germline CDKN2A mutations for melanoma cases from Australia, Spain and the United Kingdom. <i>Hereditary Cancer in Clinical Practice</i> , 2014 , 12, 20	2.3	31
97	The effect on melanoma risk of genes previously associated with telomere length. <i>Journal of the National Cancer Institute</i> , 2014 , 106,	9.7	97
96	Targeting human apurinic/apyrimidinic endonuclease 1 (APE1) in phosphatase and tensin homolog (PTEN) deficient melanoma cells for personalized therapy. <i>Oncotarget</i> , 2014 , 5, 3273-86	3.3	37
95	MC1R genotype as a predictor of early-onset melanoma, compared with self-reported and physician-measured traditional risk factors: an Australian case-control-family study. <i>BMC Cancer</i> , 2013 , 13, 406	4.8	24
94	Do vitamin A serum levels moderate outcome or the protective effect of vitamin D on outcome from malignant melanoma?. <i>Clinical Nutrition</i> , 2013 , 32, 1012-6	5.9	6
93	A variant in FTO shows association with melanoma risk not due to BMI. <i>Nature Genetics</i> , 2013 , 45, 428-32, 432e1	36.3	95
92	Association between putative functional variants in the PSMB9 gene and risk of melanoma--re-analysis of published melanoma genome-wide association studies. <i>Pigment Cell and Melanoma Research</i> , 2013 , 26, 392-401	4.5	4
91	Melanoma prone families with CDK4 germline mutation: phenotypic profile and associations with MC1R variants. <i>Journal of Medical Genetics</i> , 2013 , 50, 264-70	5.8	89
90	Association between functional polymorphisms in genes involved in the MAPK signaling pathways and cutaneous melanoma risk. <i>Carcinogenesis</i> , 2013 , 34, 885-92	4.6	6
89	Erythema nodosum-like panniculitis in patients with melanoma treated with vemurafenib. <i>Journal of Clinical Oncology</i> , 2013 , 31, e320-1	2.2	23
88	Evaluation of PAX3 genetic variants and nevus number. <i>Pigment Cell and Melanoma Research</i> , 2013 , 26, 666-76	4.5	5
87	Response to P. Autier and M. Boniol regarding our article--Relationship between sunbed use and melanoma risk in a large case-control study in the United Kingdom. <i>International Journal of Cancer</i> , 2013 , 132, 1960-1	7.5	1

86	Vitamin D and melanoma. <i>Dermato-Endocrinology</i> , 2013 , 5, 121-9		20
85	Identification of differentially expressed genes in matched formalin-fixed paraffin-embedded primary and metastatic melanoma tumor pairs. <i>Pigment Cell and Melanoma Research</i> , 2012 , 25, 284-6	4.5	5
84	Inherited variants in the MC1R gene and survival from cutaneous melanoma: a BioGenoMEL study. <i>Pigment Cell and Melanoma Research</i> , 2012 , 25, 384-94	4.5	50
83	Perceptions of genetic research and testing among members of families with an increased risk of malignant melanoma. <i>European Journal of Cancer</i> , 2012 , 48, 3052-62	7.5	14
82	Diagnosis and treatment of melanoma. European consensus-based interdisciplinary guideline--Update 2012. <i>European Journal of Cancer</i> , 2012 , 48, 2375-90	7.5	199
81	EORTC Melanoma Group achievements. <i>European Journal of Cancer, Supplement</i> , 2012 , 10, 112-119	1.6	
80	Melanocortin-1 receptor, skin cancer and phenotypic characteristics (M-SKIP) project: study design and methods for pooling results of genetic epidemiological studies. <i>BMC Medical Research Methodology</i> , 2012 , 12, 116	4.7	10
79	Relationship between sunbed use and melanoma risk in a large case-control study in the United Kingdom. <i>International Journal of Cancer</i> , 2012 , 130, 3011-3	7.5	12
78	Skin examination behavior: the role of melanoma history, skin type, psychosocial factors, and region of residence in determining clinical and self-conducted skin examination. <i>Archives of Dermatology</i> , 2012 , 148, 1142-51		28
77	Objective assessment of blood and lymphatic vessel invasion and association with macrophage infiltration in cutaneous melanoma. <i>Modern Pathology</i> , 2012 , 25, 493-504	9.8	91
76	Clinicopathologic features of V600E and V600K melanoma--letter. <i>Clinical Cancer Research</i> , 2012 , 18, 6792; author's reply p. 6793	12.9	14
75	Genome-wide association study identifies three new melanoma susceptibility loci. <i>Nature Genetics</i> , 2011 , 43, 1108-13	36.3	203
74	Melanoma and vitamin D. <i>Molecular Oncology</i> , 2011 , 5, 197-214	7.9	50
73	Melanocytic Naevi and Melanoma 2011 , 109.1-109.28		1
72	Relationship between sun exposure and melanoma risk for tumours in different body sites in a large case-control study in a temperate climate. <i>European Journal of Cancer</i> , 2011 , 47, 732-41	7.5	80
71	Infliximab in the treatment of a child with cutaneous granulomas associated with ataxia telangiectasia. <i>Journal of the American Academy of Dermatology</i> , 2011 , 65, 676-677	4.5	21
70	Pathway-based analysis of a melanoma genome-wide association study: analysis of genes related to tumour-immunosuppression. <i>PLoS ONE</i> , 2011 , 6, e29451	3.7	16
69	The determinants of periorbital skin ageing in participants of a melanoma case-control study in the U.K. <i>British Journal of Dermatology</i> , 2011 , 165, 1011-21	4	13

68	The determinants of serum vitamin D levels in participants in a melanoma case-control study living in a temperate climate. <i>Cancer Causes and Control</i> , 2011 , 22, 1471-82	2.8	27
67	Melanoma risk for CDKN2A mutation carriers who are relatives of population-based case carriers in Australia and the UK. <i>Journal of Medical Genetics</i> , 2011 , 48, 266-72	5.8	30
66	Prognosis in patients with sentinel node-positive melanoma is accurately defined by the combined Rotterdam tumor load and Dewar topography criteria. <i>Journal of Clinical Oncology</i> , 2011 , 29, 2206-14	2.2	164
65	A novel recurrent mutation in MITF predisposes to familial and sporadic melanoma. <i>Nature</i> , 2011 , 480, 99-103	50.4	335
64	Genome-wide association study identifies a new melanoma susceptibility locus at 1q21.3. <i>Nature Genetics</i> , 2011 , 43, 1114-8	36.3	126
63	Predictors of sun protection behaviors and severe sunburn in an international online study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010 , 19, 2199-210	4	91
62	Patterns of expression of DNA repair genes and relapse from melanoma. <i>Clinical Cancer Research</i> , 2010 , 16, 5211-21	12.9	48
61	Melanocytic nevi, nevus genes, and melanoma risk in a large case-control study in the United Kingdom. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010 , 19, 2043-54	4	83
60	Reply to P.E. Hutchinson et al. <i>Journal of Clinical Oncology</i> , 2010 , 28, e494-e495	2.2	
59	Diagnosis and treatment of melanoma: European consensus-based interdisciplinary guideline. <i>European Journal of Cancer</i> , 2010 , 46, 270-83	7.5	230
58	Laryngo-onycho-cutaneous syndrome. <i>Ophthalmology</i> , 2010 , 117, 1056-1056.e2	7.3	6
57	Melanoma sentinel node biopsy and prediction models for relapse and overall survival. <i>British Journal of Cancer</i> , 2010 , 103, 1229-36	8.7	43
56	Melanoma risk factors, perceived threat and intentional tanning: an international online survey. <i>European Journal of Cancer Prevention</i> , 2010 , 19, 216-26	2	41
55	IRF4 variants have age-specific effects on nevus count and predispose to melanoma. <i>American Journal of Human Genetics</i> , 2010 , 87, 6-16	11	100
54	Deletion at chromosome arm 9p in relation to BRAF/NRAS mutations and prognostic significance for primary melanoma. <i>Genes Chromosomes and Cancer</i> , 2010 , 49, 425-38	5	40
53	Sun exposure and melanoma risk at different latitudes: a pooled analysis of 5700 cases and 7216 controls. <i>International Journal of Epidemiology</i> , 2009 , 38, 814-30	7.8	187
52	Gene expression profiling of paraffin-embedded primary melanoma using the DASL assay identifies increased osteopontin expression as predictive of reduced relapse-free survival. <i>Clinical Cancer Research</i> , 2009 , 15, 6939-46	12.9	83
51	Serum 25-hydroxyvitamin D3 levels are associated with breslow thickness at presentation and survival from melanoma. <i>Journal of Clinical Oncology</i> , 2009 , 27, 5439-44	2.2	212

50	Cutaneous melanoma during pregnancy: is the controversy over?. <i>Journal of Clinical Oncology</i> , 2009 , 27, e11-2; author reply e13-4	2.2	17
49	A pooled analysis of melanocytic nevus phenotype and the risk of cutaneous melanoma at different latitudes. <i>International Journal of Cancer</i> , 2009 , 124, 420-8	7.5	72
48	Overseas sun exposure, nevus counts, and premature skin aging in young English women: a population-based survey. <i>Journal of Investigative Dermatology</i> , 2009 , 129, 50-9	4.3	34
47	Genome-wide association study identifies variants at 9p21 and 22q13 associated with development of cutaneous nevi. <i>Nature Genetics</i> , 2009 , 41, 915-9	36.3	186
46	Genome-wide association study identifies three loci associated with melanoma risk. <i>Nature Genetics</i> , 2009 , 41, 920-5	36.3	360
45	Vitamin D receptor gene polymorphisms, serum 25-hydroxyvitamin D levels, and melanoma: UK case-control comparisons and a meta-analysis of published VDR data. <i>European Journal of Cancer</i> , 2009 , 45, 3271-81	7.5	108
44	Selection criteria for genetic assessment of patients with familial melanoma. <i>Journal of the American Academy of Dermatology</i> , 2009 , 61, 677.e1-14	4.5	115
43	Spontaneous involution of congenital melanocytic nevi of the scalp. <i>Journal of the American Academy of Dermatology</i> , 2008 , 58, 508-11	4.5	50
42	Environmental risk factors for relapse of melanoma. <i>European Journal of Cancer</i> , 2008 , 44, 1717-25	7.5	13
41	Frequent p16-independent inactivation of p14ARF in human melanoma. <i>Journal of the National Cancer Institute</i> , 2008 , 100, 784-95	9.7	83
40	Trends in prognostic factors and survival from cutaneous melanoma in Yorkshire, UK and New South Wales, Australia between 1993 and 2003. <i>International Journal of Cancer</i> , 2008 , 123, 861-6	7.5	37
39	Sun-protective behaviors in families at increased risk of melanoma. <i>Journal of Investigative Dermatology</i> , 2007 , 127, 1343-50	4.3	13
38	Genetics: what advice for patients who present with a family history of melanoma?. <i>Seminars in Oncology</i> , 2007 , 34, 452-9	5.5	20
37	CDKN2A and CDK4 variants in Latvian melanoma patients: analysis of a clinic-based population. <i>Melanoma Research</i> , 2007 , 17, 185-91	3.3	32
36	The emergence of networks in human genome epidemiology: challenges and opportunities. <i>Epidemiology</i> , 2007 , 18, 1-8	3.1	93
35	A road map for efficient and reliable human genome epidemiology. <i>Nature Genetics</i> , 2006 , 38, 3-5	36.3	198
34	Malignant melanoma in pregnancy. <i>Obstetrics and Gynecology Clinics of North America</i> , 2005 , 32, 559-68	3.3	15
33	A mutation hotspot at the p14ARF splice site. <i>Oncogene</i> , 2005 , 24, 4604-8	9.2	67

32	The effect of sun exposure in determining nevus density in UK adolescent twins. <i>Journal of Investigative Dermatology</i> , 2005 , 124, 56-62	4.3	53
31	Intronic sequence variants of the CDKN2A gene in melanoma pedigrees. <i>Genes Chromosomes and Cancer</i> , 2005 , 43, 128-36	5	29
30	Prevalence of 9p21 deletions in UK melanoma families. <i>Genes Chromosomes and Cancer</i> , 2005 , 44, 292-300	3.8	35
29	A network of investigator networks in human genome epidemiology. <i>American Journal of Epidemiology</i> , 2005 , 162, 302-4	3.8	92
28	No Evidence for BRAF as a melanoma/nevus susceptibility gene. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2005 , 14, 913-8	4	23
27	The genetics of susceptibility to cutaneous melanoma. <i>Drugs of Today</i> , 2005 , 41, 193-203		30
26	Effect of pregnancy on survival in women with cutaneous malignant melanoma. <i>Journal of Clinical Oncology</i> , 2004 , 22, 4369-75	2.2	130
25	The relationship between the epidermal growth factor (EGF) 5'UTR variant A61G and melanoma/nevus susceptibility. <i>Journal of Investigative Dermatology</i> , 2004 , 123, 755-9	4.3	34
24	An assessment of a variant of the DNA repair gene XRCC3 as a possible nevus or melanoma susceptibility genotype. <i>Journal of Investigative Dermatology</i> , 2004 , 122, 429-32	4.3	21
23	A quality-of-life study in high-risk (thickness \geq 2 mm) cutaneous melanoma patients in a randomized trial of 1-cm versus 3-cm surgical excision margins. <i>Journal of Investigative Dermatology Symposium Proceedings</i> , 2004 , 9, 152-9	1.1	35
22	Surgical margins in cutaneous melanoma (2 cm versus 5 cm for lesions measuring less than 2.1-mm thick). <i>Cancer</i> , 2004 , 100, 433-4; author reply 434	6.4	2
21	Excision margins in high-risk malignant melanoma. <i>New England Journal of Medicine</i> , 2004 , 350, 757-66	59.2	341
20	UK guidelines for the management of cutaneous melanoma. <i>Journal of Plastic, Reconstructive and Aesthetic Surgery</i> , 2002 , 55, 46-54		60
19	Management of regional lymph nodes in patients with malignant melanoma: questionnaire survey of UK current practice. <i>Journal of Plastic, Reconstructive and Aesthetic Surgery</i> , 2002 , 55, 372-5		10
18	An assessment of the CDKN2A variant Ala148Thr as a nevus/melanoma susceptibility allele. <i>Journal of Investigative Dermatology</i> , 2002 , 119, 961-5	4.3	29
17	Biallelic mutations in p16(INK4a) confer resistance to Ras- and Ets-induced senescence in human diploid fibroblasts. <i>Molecular and Cellular Biology</i> , 2002 , 22, 8135-43	4.8	108
16	Excision margins in the treatment of primary cutaneous melanoma: a systematic review of randomized controlled trials comparing narrow vs wide excision. <i>Archives of Surgery</i> , 2002 , 137, 1101-5		52
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14	Melanoma. <i>British Journal of Hospital Medicine</i> , 2000 , 61, 103-7		1
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4	Tumour gene expression signature in primary melanoma predicts long-term outcomes: A prospective multicentre study		2
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1	Surgical excision methods for skin cancer involving the nail unit. <i>The Cochrane Library</i> ,	5.2	78