

# Hkan L Olsson

## 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.

300  
papers

16,308  
citations

63  
h-index

120  
g-index

317  
ext. papers

19,824  
ext. citations

8.3  
avg, IF

5.6  
L-index

#	Paper	IF	Citations
300	Rare germline copy number variants (CNVs) and breast cancer risk.. <i>Communications Biology</i> , <b>2022</b> , 5, 65	6.7	0
299	Common variants in breast cancer risk loci predispose to distinct tumor subtypes.. <i>Breast Cancer Research</i> , <b>2022</b> , 24, 2	8.3	3
298	Oral Contraceptive Use in BRCA1 and BRCA2 Mutation Carriers: Absolute Cancer Risks and Benefits.. <i>Journal of the National Cancer Institute</i> , <b>2022</b> ,	9.7	1
297	A Genome-Wide Gene-Based Gene-Environment Interaction Study of Breast Cancer in More than 90,000 Women. <i>Cancer Research Communications</i> , <b>2022</b> , 2, 211-219		0
296	Genome-wide interaction analysis of menopausal hormone therapy use and breast cancer risk among 62,370 women.. <i>Scientific Reports</i> , <b>2022</b> , 12, 6199	4.9	
295	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> , <b>2021</b> , 30, 217-228	4	7
294	Germline variants and breast cancer survival in patients with distant metastases at primary breast cancer diagnosis. <i>Scientific Reports</i> , <b>2021</b> , 11, 19787	4.9	0
293	Prevalence of germline pathogenic variants in 22 cancer susceptibility genes in Swedish pediatric cancer patients. <i>Scientific Reports</i> , <b>2021</b> , 11, 5307	4.9	2
292	Improved survival in several cancers with use of H-antihistamines desloratadine and loratadine. <i>Translational Oncology</i> , <b>2021</b> , 14, 101029	4.9	4
291	Birth cohort-specific trends of sun-related behaviors among individuals from an international consortium of melanoma-prone families. <i>BMC Public Health</i> , <b>2021</b> , 21, 692	4.1	2
290	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> , <b>2021</b> , 2, 100042-100042	0.8	2
289	The human melanoma proteome atlas-Defining the molecular pathology. <i>Clinical and Translational Medicine</i> , <b>2021</b> , 11, e473	5.7	3
288	The Human Melanoma Proteome Atlas-Complementing the melanoma transcriptome. <i>Clinical and Translational Medicine</i> , <b>2021</b> , 11, e451	5.7	5
287	Functional annotation of the 2q35 breast cancer risk locus implicates a structural variant in influencing activity of a long-range enhancer element. <i>American Journal of Human Genetics</i> , <b>2021</b> , 108, 1190-1203	11	1
286	Combined Associations of a Polygenic Risk Score and Classical Risk Factors With Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , <b>2021</b> , 113, 329-337	9.7	14
285	Multiple Primary Melanoma Incidence Trends Over Five Decades: A Nationwide Population-Based Study. <i>Journal of the National Cancer Institute</i> , <b>2021</b> , 113, 318-328	9.7	6
284	Common Susceptibility Loci for Male Breast Cancer. <i>Journal of the National Cancer Institute</i> , <b>2021</b> , 113, 453-461	9.7	4

283	CYP3A7*1C allele: linking premenopausal oestrone and progesterone levels with risk of hormone receptor-positive breast cancers. <i>British Journal of Cancer</i> , <b>2021</b> , 124, 842-854	8.7	2
282	A case-only study to identify genetic modifiers of breast cancer risk for BRCA1/BRCA2 mutation carriers. <i>Nature Communications</i> , <b>2021</b> , 12, 1078	17.4	4
281	Oral contraceptive use and ovarian cancer risk for BRCA1/2 mutation carriers: an international cohort study. <i>American Journal of Obstetrics and Gynecology</i> , <b>2021</b> , 225, 51.e1-51.e17	6.4	9
280	Association of germline genetic variants with breast cancer-specific survival in patient subgroups defined by clinic-pathological variables related to tumor biology and type of systemic treatment. <i>Breast Cancer Research</i> , <b>2021</b> , 23, 86	8.3	1
279	Mendelian randomisation study of smoking exposure in relation to breast cancer risk. <i>British Journal of Cancer</i> , <b>2021</b> , 125, 1135-1145	8.7	0
278	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , <b>2021</b> , 596, 393-397	30.4	28
277	Genetic testing in women with early-onset breast cancer: a Traceback pilot study. <i>Breast Cancer Research and Treatment</i> , <b>2021</b> , 190, 307-315	4.4	
276	Breast Cancer Risk Factors and Survival by Tumor Subtype: Pooled Analyses from the Breast Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2021</b> , 30, 623-642	4	4
275	Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. <i>Nature Genetics</i> , <b>2020</b> , 52, 572-581	36.3	76
274	Desloratadine and loratadine stand out among common H-antihistamines for association with improved breast cancer survival. <i>Acta Oncologica</i> , <b>2020</b> , 59, 1103-1109	3.2	10
273	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. <i>Scientific Reports</i> , <b>2020</b> , 10, 9688	4.9	2
272	Analysis of DNA methylation patterns in the tumor immune microenvironment of metastatic melanoma. <i>Molecular Oncology</i> , <b>2020</b> , 14, 933-950	7.9	16
271	Protein Expression in Metastatic Melanoma and the Link to Disease Presentation in a Range of Tumor Phenotypes. <i>Cancers</i> , <b>2020</b> , 12,	6.6	2
270	Desloratadine and loratadine use associated with improved melanoma survival. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , <b>2020</b> , 75, 2096-2099	9.3	12
269	Variations in the Referral Pattern for Genetic Counseling of Patients with Early-Onset Breast Cancer: A Population-Based Study in Southern Sweden. <i>Public Health Genomics</i> , <b>2020</b> , 23, 100-109	1.9	1
268	Transcriptome-wide association study of breast cancer risk by estrogen-receptor status. <i>Genetic Epidemiology</i> , <b>2020</b> , 44, 442-468	2.6	9
267	Tertiary lymphoid structures improve immunotherapy and survival in melanoma. <i>Nature</i> , <b>2020</b> , 577, 561-565	56.4	542
266	Alcohol Consumption, Cigarette Smoking, and Risk of Breast Cancer for and Mutation Carriers: Results from The BRCA1 and BRCA2 Cohort Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2020</b> , 29, 368-378	4	9

265	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , <b>2020</b> , 11, 312	17.4	20
264	Risk-reducing salpingo-oophorectomy, natural menopause, and breast cancer risk: an international prospective cohort of BRCA1 and BRCA2 mutation carriers. <i>Breast Cancer Research</i> , <b>2020</b> , 22, 8	8.3	22
263	Women with a predisposition for diabetes have an increased risk of pregnancy complications, especially in combination with pregestational overweight. <i>BMC Pregnancy and Childbirth</i> , <b>2020</b> , 20, 74	3.2	1
262	The Role of PTEN Loss in Immune Escape, Melanoma Prognosis and Therapy Response. <i>Cancers</i> , <b>2020</b> , 12,	6.6	12
261	Genome-wide association meta-analyses combining multiple risk phenotypes provide insights into the genetic architecture of cutaneous melanoma susceptibility. <i>Nature Genetics</i> , <b>2020</b> , 52, 494-504	36.3	39
260	Women with fair phenotypes seem to confer a survival advantage in a low UV milieu. A nested matched case control study. <i>PLoS ONE</i> , <b>2020</b> , 15, e0228582	3.7	4
259	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , <b>2020</b> , 52, 56-73	36.3	56
258	Tumor genetic heterogeneity analysis of chronic sun-damaged melanoma. <i>Pigment Cell and Melanoma Research</i> , <b>2020</b> , 33, 480-489	4.5	7
257	Increased Cancer Risk in Families with Pediatric Cancer Is Associated with Gender, Age, Diagnosis, and Degree of Relation to the Child. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2020</b> , 29, 2171-2174	4	1
256	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , <b>2020</b> , 107, 837-848	11	12
255	Efficacy of novel immunotherapy regimens in patients with metastatic melanoma with germline mutations. <i>Journal of Medical Genetics</i> , <b>2020</b> , 57, 316-321	5.8	18
254	Genetically Predicted Levels of DNA Methylation Biomarkers and Breast Cancer Risk: Data From 228 951 Women of European Descent. <i>Journal of the National Cancer Institute</i> , <b>2020</b> , 112, 295-304	9.7	18
253	Novel functional proteins coded by the human genome discovered in metastases of melanoma patients. <i>Cell Biology and Toxicology</i> , <b>2020</b> , 36, 261-272	7.4	9
252	Evaluation of associations between genetically predicted circulating protein biomarkers and breast cancer risk. <i>International Journal of Cancer</i> , <b>2020</b> , 146, 2130-2138	7.5	9
251	Two truncating variants in FANCC and breast cancer risk. <i>Scientific Reports</i> , <b>2019</b> , 9, 12524	4.9	2
250	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , <b>2019</b> , 10, 431	17.4	45
249	Rationale for a Swedish cohort consortium. <i>Uppsala Journal of Medical Sciences</i> , <b>2019</b> , 124, 21-28	2.8	1
248	The X-Linked DDX3X RNA Helicase Dictates Translation Reprogramming and Metastasis in Melanoma. <i>Cell Reports</i> , <b>2019</b> , 27, 3573-3586.e7	10.6	36

247	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , <b>2019</b> , 10, 1741	17.4	47
246	Physical activity and survival following breast cancer. <i>European Journal of Cancer Care</i> , <b>2019</b> , 28, e13037	2.4	10
245	Clinical protein science in translational medicine targeting malignant melanoma. <i>Cell Biology and Toxicology</i> , <b>2019</b> , 35, 293-332	7.4	20
244	Improved survival prognostication of node-positive malignant melanoma patients utilizing shotgun proteomics guided by histopathological characterization and genomic data. <i>Scientific Reports</i> , <b>2019</b> , 9, 5154	4.9	9
243	Genome-wide association study of germline variants and breast cancer-specific mortality. <i>British Journal of Cancer</i> , <b>2019</b> , 120, 647-657	8.7	28
242	The association between weight at birth and breast cancer risk revisited using Mendelian randomisation. <i>European Journal of Epidemiology</i> , <b>2019</b> , 34, 591-600	12.1	11
241	Risk factors for subarachnoid haemorrhage: a nationwide cohort of 950000 adults. <i>International Journal of Epidemiology</i> , <b>2019</b> , 48, 2018-2025	7.8	9
240	A Targeted Mass Spectrometry Strategy for Developing Proteomic Biomarkers: A Case Study of Epithelial Ovarian Cancer. <i>Molecular and Cellular Proteomics</i> , <b>2019</b> , 18, 1836-1850	7.6	24
239	Estimating CDKN2A mutation carrier probability among global familial melanoma cases using GenoMELPREDICT. <i>Journal of the American Academy of Dermatology</i> , <b>2019</b> , 81, 386-394	4.5	9
238	The :p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. <i>Npj Breast Cancer</i> , <b>2019</b> , 5, 38	7.8	12
237	A single exercise session improves side-effects of chemotherapy in women with breast cancer: an observational study. <i>BMC Cancer</i> , <b>2019</b> , 19, 1073	4.8	11
236	Evaluation of the contribution of germline variants in BRCA1 and BRCA2 to uveal and cutaneous melanoma. <i>Melanoma Research</i> , <b>2019</b> , 29, 483-490	3.3	7
235	The Hidden Story of Heterogeneous B-raf V600E Mutation Quantitative Protein Expression in Metastatic Melanoma-Association with Clinical Outcome and Tumor Phenotypes. <i>Cancers</i> , <b>2019</b> , 11,	6.6	10
234	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. <i>American Journal of Human Genetics</i> , <b>2019</b> , 104, 21-34	11	363
233	Associations of obesity and circulating insulin and glucose with breast cancer risk: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , <b>2019</b> , 48, 795-806	7.8	52
232	Genetic Data from Nearly 63,000 Women of European Descent Predicts DNA Methylation Biomarkers and Epithelial Ovarian Cancer Risk. <i>Cancer Research</i> , <b>2019</b> , 79, 505-517	10.1	28
231	CM-Score: a validated scoring system to predict germline mutations in melanoma families from Northern Europe. <i>Journal of Medical Genetics</i> , <b>2018</b> , 55, 661-668	5.8	11
230	Current smoking is associated with a larger waist circumference and a more androgenic profile in young healthy women from high-risk breast cancer families. <i>Cancer Causes and Control</i> , <b>2018</b> , 29, 243-251	2.8	6

229	Stellate cells and mesenchymal stem cells in benign mammary stroma are associated with risk factors for breast cancer - an observational study. <i>BMC Cancer</i> , <b>2018</b> , 18, 230	4.8	4
228	Assessment of moderate coffee consumption and risk of epithelial ovarian cancer: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , <b>2018</b> , 47, 450-459	7.8	8
227	Plasma enterolactone and risk of prostate cancer in middle-aged Swedish men. <i>European Journal of Nutrition</i> , <b>2018</b> , 57, 2595-2606	5.2	8
226	Lung cancer in young women in southern Sweden: A descriptive study. <i>Clinical Respiratory Journal</i> , <b>2018</b> , 12, 1565-1571	1.7	7
225	A Transcriptome-Wide Association Study Among 97,898 Women to Identify Candidate Susceptibility Genes for Epithelial Ovarian Cancer Risk. <i>Cancer Research</i> , <b>2018</b> , 78, 5419-5430	10.1	32
224	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , <b>2018</b> , 9, 3166	17.4	70
223	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. <i>Nature Genetics</i> , <b>2018</b> , 50, 968-978	36.3	101
222	Accuracy of self-reported family history of cancer, mutation status and tumor characteristics in patients with early onset breast cancer. <i>Acta Oncologica</i> , <b>2018</b> , 57, 595-603	3.2	14
221	Phenocopies in melanoma-prone families with germ-line CDKN2A mutations. <i>Genetics in Medicine</i> , <b>2018</b> , 20, 1087-1090	8.1	5
220	The Influence of Number and Timing of Pregnancies on Breast Cancer Risk for Women With or Mutations. <i>JNCI Cancer Spectrum</i> , <b>2018</b> , 2, pky078	4.6	10
219	Endogenous expression mapping of malignant melanoma by mass spectrometry imaging. <i>Clinical and Translational Medicine</i> , <b>2018</b> , 7, 22	5.7	6
218	Oral Contraceptive Use and Breast Cancer Risk: Retrospective and Prospective Analyses From a BRCA1 and BRCA2 Mutation Carrier Cohort Study. <i>JNCI Cancer Spectrum</i> , <b>2018</b> , 2, pky023	4.6	13
217	Risks of Breast, Ovarian, and Contralateral Breast Cancer for BRCA1 and BRCA2 Mutation Carriers. <i>JAMA - Journal of the American Medical Association</i> , <b>2017</b> , 317, 2402-2416	27.4	1140
216	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , <b>2017</b> , 49, 680-691	36.3	190
215	NF1-mutated melanoma tumors harbor distinct clinical and biological characteristics. <i>Molecular Oncology</i> , <b>2017</b> , 11, 438-451	7.9	73
214	Occupational sedentariness and breast cancer risk. <i>Acta Oncologica</i> , <b>2017</b> , 56, 75-80	3.2	16
213	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , <b>2017</b> , 551, 92-94	50.4	643
212	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , <b>2017</b> , 49, 1767-1778	36.3	186

211	Predominance of girls with cancer in families with multiple childhood cancer cases. <i>BMC Cancer</i> , <b>2017</b> , 17, 868	4.8	4
210	Germline Variation at CDKN2A and Associations with Nevus Phenotypes among Members of Melanoma Families. <i>Journal of Investigative Dermatology</i> , <b>2017</b> , 137, 2606-2612	4.3	10
209	Cancer risks and survival in patients with multiple primary melanomas: Association with family history of melanoma and germline CDKN2A mutation status. <i>Journal of the American Academy of Dermatology</i> , <b>2017</b> , 77, 893-901	4.5	19
208	Impact of Pregestational Weight and Weight Gain during Pregnancy on Long-Term Risk for Diseases. <i>PLoS ONE</i> , <b>2017</b> , 12, e0168543	3.7	28
207	Correlation of histopathologic characteristics to protein expression and function in malignant melanoma. <i>PLoS ONE</i> , <b>2017</b> , 12, e0176167	3.7	17
206	Multiple rare variants in high-risk pancreatic cancer-related genes may increase risk for pancreatic cancer in a subset of patients with and without germline CDKN2A mutations. <i>Human Genetics</i> , <b>2016</b> , 135, 1241-1249	6.3	19
205	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. <i>Cancer Discovery</i> , <b>2016</b> , 6, 1052-67	24.4	104
204	Multiregion Whole-Exome Sequencing Uncovers the Genetic Evolution and Mutational Heterogeneity of Early-Stage Metastatic Melanoma. <i>Cancer Research</i> , <b>2016</b> , 76, 4765-74	10.1	70
203	Germline CDKN2A Mutation Status and Survival in Familial Melanoma Cases. <i>Journal of the National Cancer Institute</i> , <b>2016</b> , 108,	9.7	29
202	Phenotypic and Histopathological Tumor Characteristics According to CDKN2A Mutation Status among Affected Members of Melanoma Families. <i>Journal of Investigative Dermatology</i> , <b>2016</b> , 136, 1066-1069	4.3	9
201	The absence of aldehyde dehydrogenase 1 A1-positive cells in benign mammary stroma is associated with risk factors for breast cancer. <i>Breast Cancer: Targets and Therapy</i> , <b>2016</b> , 8, 117-24	3.9	1
200	High TERT promoter mutation frequency in non-acral cutaneous metastatic melanoma. <i>Pigment Cell and Melanoma Research</i> , <b>2016</b> , 29, 598-600	4.5	20
199	The BRCA1- $\Delta$ 11q Alternative Splice Isoform Bypasses Germline Mutations and Promotes Therapeutic Resistance to PARP Inhibition and Cisplatin. <i>Cancer Research</i> , <b>2016</b> , 76, 2778-90	10.1	136
198	The clinicopathological and gene expression patterns associated with ulceration of primary melanoma. <i>Pigment Cell and Melanoma Research</i> , <b>2015</b> , 28, 94-104	4.5	23
197	Nonsense mutations in the shelterin complex genes ACD and TERF2IP in familial melanoma. <i>Journal of the National Cancer Institute</i> , <b>2015</b> , 107,	9.7	102
196	Genome-wide meta-analysis identifies five new susceptibility loci for cutaneous malignant melanoma. <i>Nature Genetics</i> , <b>2015</b> , 47, 987-995	36.3	162
195	Fine mapping of genetic susceptibility loci for melanoma reveals a mixture of single variant and multiple variant regions. <i>International Journal of Cancer</i> , <b>2015</b> , 136, 1351-60	7.5	26
194	Tobacco and alcohol in relation to male breast cancer: an analysis of the male breast cancer pooling project consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2015</b> , 24, 520-31	4	15

193	Impact of a paternal origin of germline BRCA1/2 mutations on the age at breast and ovarian cancer diagnosis in a Southern Swedish cohort. <i>Genes Chromosomes and Cancer</i> , <b>2015</b> , 54, 39-50	5	5
192	CDKN2a mutation-negative melanoma families have increased risk exclusively for skin cancers but not for other malignancies. <i>International Journal of Cancer</i> , <b>2015</b> , 137, 2220-6	7.5	13
191	A protein deep sequencing evaluation of metastatic melanoma tissues. <i>PLoS ONE</i> , <b>2015</b> , 10, e0123661	3.7	17
190	Development and validation of a melanoma risk score based on pooled data from 16 case-control studies. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2015</b> , 24, 817-24	4	18
189	Germline mutation in BRCA1 or BRCA2 and ten-year survival for women diagnosed with epithelial ovarian cancer. <i>Clinical Cancer Research</i> , <b>2015</b> , 21, 652-7	12.9	107
188	Molecular stratification of metastatic melanoma using gene expression profiling: Prediction of survival outcome and benefit from molecular targeted therapy. <i>Oncotarget</i> , <b>2015</b> , 6, 12297-309	3.3	102
187	Anthropometric and hormonal risk factors for male breast cancer: male breast cancer pooling project results. <i>Journal of the National Cancer Institute</i> , <b>2014</b> , 106, djt465	9.7	92
186	Feasibility study on measuring selected proteins in malignant melanoma tissue by SRM quantification. <i>Journal of Proteome Research</i> , <b>2014</b> , 13, 1315-26	5.6	7
185	High risk of in-breast tumor recurrence after BRCA1/2-associated breast cancer. <i>Breast Cancer Research and Treatment</i> , <b>2014</b> , 147, 571-8	4.4	34
184	A new look at drugs targeting malignant melanoma--an application for mass spectrometry imaging. <i>Proteomics</i> , <b>2014</b> , 14, 1963-70	4.8	25
183	Primary melanoma tumors from CDKN2A mutation carriers do not belong to a distinct molecular subclass. <i>Journal of Investigative Dermatology</i> , <b>2014</b> , 134, 3000-3003	4.3	5
182	Oral contraceptive use, parity, and constitutional characteristics in soft tissue sarcoma: a Swedish population-based case-control study 1988-2009. <i>Cancer Causes and Control</i> , <b>2014</b> , 25, 1167-77	2.8	2
181	Investigation of a putative melanoma susceptibility locus at chromosome 3q29. <i>Cancer Genetics</i> , <b>2014</b> , 207, 70-4	2.3	3
180	Epithelial and stromal microRNA signatures of columnar cell hyperplasia linking Let-7c to precancerous and cancerous breast cancer cell proliferation. <i>PLoS ONE</i> , <b>2014</b> , 9, e105099	3.7	20
179	Somatic BRAF and NRAS mutations in familial melanomas with known germline CDKN2A status: a GenoMEL study. <i>Journal of Investigative Dermatology</i> , <b>2014</b> , 134, 287-290	4.3	15
178	High risk of tobacco-related cancers in CDKN2A mutation-positive melanoma families. <i>Journal of Medical Genetics</i> , <b>2014</b> , 51, 545-52	5.8	55
177	Inherited variation in the PARP1 gene and survival from melanoma. <i>International Journal of Cancer</i> , <b>2014</b> , 135, 1625-33	7.5	22
176	Molecular and genetic diversity in the metastatic process of melanoma. <i>Journal of Pathology</i> , <b>2014</b> , 233, 39-50	9.4	47



175	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> , <b>2014</b> , 27, 234-43	4.5	21
174	Plasma alkylresorcinol metabolites as biomarkers for whole-grain intake and their association with prostate cancer: a Swedish nested case-control study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2014</b> , 23, 73-83	4	17
173	The effect on melanoma risk of genes previously associated with telomere length. <i>Journal of the National Cancer Institute</i> , <b>2014</b> , 106,	9.7	97
172	Analysis of alpha-synuclein in malignant melanoma - development of a SRM quantification assay. <i>PLoS ONE</i> , <b>2014</b> , 9, e110804	3.7	19
171	Women Who Develop Diabetes Later in Life Have Diabetes-Associated Complications during Preceding Pregnancies. <i>Journal of Diabetes Mellitus</i> , <b>2014</b> , 04, 341-349	0.5	1
170	Prognostic factors in lung cancer in a defined geographical area over two decades with a special emphasis on gender. <i>Clinical Respiratory Journal</i> , <b>2013</b> , 7, 91-100	1.7	9
169	Establishing a Southern Swedish Malignant Melanoma OMICS and biobank clinical capability. <i>Clinical and Translational Medicine</i> , <b>2013</b> , 2, 7	5.7	15
168	Tamoxifen and risk of contralateral breast cancer for BRCA1 and BRCA2 mutation carriers. <i>Journal of Clinical Oncology</i> , <b>2013</b> , 31, 3091-9	2.2	118
167	Autoimmune diseases and hypersensitivities improve the prognosis in ER-negative breast cancer. <i>SpringerPlus</i> , <b>2013</b> , 2, 357		4
166	A variant in FTO shows association with melanoma risk not due to BMI. <i>Nature Genetics</i> , <b>2013</b> , 45, 428-32, 432e1	36.3	95
165	Women with familial risk for breast cancer have an increased frequency of aldehyde dehydrogenase expressing cells in breast ductules. <i>BMC Clinical Pathology</i> , <b>2013</b> , 13, 28	3	12
164	Gene expression profiling of primary male breast cancers reveals two unique subgroups and identifies N-acetyltransferase-1 (NAT1) as a novel prognostic biomarker. <i>Breast Cancer Research</i> , <b>2012</b> , 14, R31	8.3	83
163	Ovarian cancer susceptibility alleles and risk of ovarian cancer in BRCA1 and BRCA2 mutation carriers. <i>Human Mutation</i> , <b>2012</b> , 33, 690-702	4.7	31
162	Breast cancer and spider telangiectasias at diagnosis and its relation to histopathology and prognosis: a population-based study. <i>Breast Cancer Research and Treatment</i> , <b>2012</b> , 131, 177-86	4.4	1
161	Increased incidence of childhood, prostate and breast cancers in relatives of childhood cancer patients. <i>Familial Cancer</i> , <b>2012</b> , 11, 145-55	3	6
160	Association between BRCA1 and BRCA2 mutations and survival in women with invasive epithelial ovarian cancer. <i>JAMA - Journal of the American Medical Association</i> , <b>2012</b> , 307, 382-90	27.4	427
159	Genome-wide association study identifies a common variant in RAD51B associated with male breast cancer risk. <i>Nature Genetics</i> , <b>2012</b> , 44, 1182-4	36.3	84
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17	Head trauma and exposure to prolactin-elevating drugs as risk factors for male breast cancer. <i>Journal of the National Cancer Institute</i> , <b>1988</b> , 80, 679-83	9.7	57
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15	Relationship between cytogenetic findings and histopathology in non-Hodgkin lymphoma. <i>Acta Pathologica, Microbiologica, Et Immunologica Scandinavica Section A, Pathology</i> , <b>1987</b> , 95, 1-5		1
14	Cytogenetic studies in Hodgkin's disease. <i>Acta Pathologica, Microbiologica, Et Immunologica Scandinavica Section A, Pathology</i> , <b>1987</b> , 95, 289-95		4

13	The regulatory and basal phosphorylation sites of hormone-sensitive lipase are dephosphorylated by protein phosphatase-1, 2A and 2C but not by protein phosphatase-2B. <i>FEBS Journal</i> , <b>1987</b> , 168, 399-405		28
12	Proliferative activity and number of clonal chromosome aberrations in non-Hodgkin's lymphomas. <i>Scandinavian Journal of Haematology</i> , <b>1986</b> , 37, 106-10		3
11	Phosphorylation of the basal site of hormone-sensitive lipase by glycogen synthase kinase-4. <i>FEBS Letters</i> , <b>1986</b> , 209, 175-80	3.8	23
10	Cytogenetic studies in non-Hodgkin lymphomas--results from fine-needle aspiration samples. <i>Hereditas</i> , <b>1985</b> , 103, 63-76	2.4	26
9	C-band polymorphism in non-Hodgkin lymphoma. <i>Hereditas</i> , <b>1985</b> , 103, 85-7	2.4	5
8	Hypophyseal tumor and gynecomastia preceding bilateral breast cancer development in a man. <i>Cancer</i> , <b>1984</b> , 53, 1974-7	6.4	24
7	PARTIAL PURIFICATION AND PROPERTIES OF A PROTEIN PHOSPHATASE FROM RAT ADIPOSE TISSUE. <i>Biochemical Society Transactions</i> , <b>1981</b> , 9, 237P-237P	5.1	
6	Common variants in breast cancer risk loci predispose to distinct tumor subtypes		1
5	Polygenic Risk Modelling for Prediction of Epithelial Ovarian Cancer Risk		1
4	Combining genome-wide studies of breast, prostate, ovarian and endometrial cancers maps cross-cancer susceptibility loci and identifies new genetic associations		2
3	Fine-mapping of 150 breast cancer risk regions identifies 178 high confidence target genes		2
2	Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses		2
1	Correlation between p53, c-erbB-2, and topoisomerase II $\alpha$ expression, DNA ploidy, hormonal receptor status and proliferation in 356 node-negative breast carcinomas: prognostic implications		1