Lenka Foretova

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.

62 14,456 115 221 h-index g-index citations papers 7.8 17,202 4.7 235 L-index avg, IF ext. citations ext. papers

#	Paper	IF	Citations
221	Occupational exposure to organic dust and risk of lymphoma subtypes in the EPILYMPH case-control study. <i>Scandinavian Journal of Work, Environment and Health</i> , 2021 , 47, 42-51	4.3	
220	Risks of breast and ovarian cancer for women harboring pathogenic missense variants in BRCA1 and BRCA2 compared with those harboring protein truncating variants <i>Genetics in Medicine</i> , 2021 ,	8.1	2
219	Occupational exposure to organic dust and risk of lymphoma subtypes in the EPILYMPH caseflontrol study. <i>Scandinavian Journal of Work, Environment and Health</i> , 2021 , 47, 42-51	4.3	O
218	Neoadjuvant Chemotherapy of Triple-Negative Breast Cancer: Evaluation of Early Clinical Response, Pathological Complete Response Rates, and Addition of Platinum Salts Benefit Based on Real-World Evidence. <i>Cancers</i> , 2021 , 13,	6.6	3
217	Application of two job indices for general occupational demands in a pooled analysis of case-control studies on lung cancer. <i>Scandinavian Journal of Work, Environment and Health</i> , 2021 , 47, 475-481	4.3	
216	The predictive ability of the 313 variant-based polygenic risk score for contralateral breast cancer risk prediction in women of European ancestry with a heterozygous BRCA1 or BRCA2 pathogenic variant. <i>Genetics in Medicine</i> , 2021 , 23, 1726-1737	8.1	2
215	Re: ERCC3, a new ovarian cancer susceptibility gene?. European Journal of Cancer, 2021, 150, 278-280	7.5	
214	Hepcidin-regulating iron metabolism genes and pancreatic ductal adenocarcinoma: a pathway analysis of genome-wide association studies. <i>American Journal of Clinical Nutrition</i> , 2021 , 114, 1408-141	7	2
213	Shared genetic risk between eating disorder- and substance-use-related phenotypes: Evidence from genome-wide association studies. <i>Addiction Biology</i> , 2021 , 26, e12880	4.6	12
212	Lung cancer risk in painters: results from the SYNERGY pooled case-control study consortium. <i>Occupational and Environmental Medicine</i> , 2021 , 78, 269-278	2.1	5
211	Morphological findings in frozen non-neoplastic kidney tissues of patients with kidney cancer from large-scale multicentric studies on renal cancer. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2021 , 478, 1099-1107	5.1	1
210	Genome-wide homozygosity and risk of four non-Hodgkin lymphoma subtypes. <i>Journal of Translational Genetics and Genomics</i> , 2021 , 5, 200-217	1.7	
209	Sexual dimorphism in cancer: insights from transcriptional signatures in kidney tissue and renal cell carcinoma. <i>Human Molecular Genetics</i> , 2021 , 30, 343-355	5.6	3
208	Oral contraceptive use and ovarian cancer risk for BRCA1/2 mutation carriers: an international cohort study. <i>American Journal of Obstetrics and Gynecology</i> , 2021 , 225, 51.e1-51.e17	6.4	9
207	Common Genetic Variation And Age at Onset Of Anorexia Nervosa. <i>Biological Psychiatry Global Open Science</i> , 2021 ,		3
206	Needlestack: an ultra-sensitive variant caller for multi-sample next generation sequencing data. <i>NAR Genomics and Bioinformatics</i> , 2020 , 2, lqaa021	3.7	2
205	Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. <i>Nature Genetics</i> , 2020 , 52, 572-581	36.3	76

(2019-2020)

204	Circulating tumour-derived KRAS mutations in pancreatic cancer cases are predominantly carried by very short fragments of cell-free DNA. <i>EBioMedicine</i> , 2020 , 55, 102462	8.8	5
203	Diesel Engine Exhaust Exposure, Smoking, and Lung Cancer Subtype Risks. A Pooled Exposure-Response Analysis of 14 Case-Control Studies. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2020 , 202, 402-411	10.2	12
202	Respirable Crystalline Silica Exposure, Smoking, and Lung Cancer Subtype Risks. A Pooled Analysis of Case-Control Studies. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2020 , 202, 412-421	10.2	11
201	Characterization of the Cancer Spectrum in Men With Germline BRCA1 and BRCA2 Pathogenic Variants: Results From the Consortium of Investigators of Modifiers of BRCA1/2 (CIMBA). <i>JAMA Oncology</i> , 2020 , 6, 1218-1230	13.4	25
200	Association of ionizing radiation dose from common medical diagnostic procedures and lymphoma risk in the Epilymph case-control study. <i>PLoS ONE</i> , 2020 , 15, e0235658	3.7	1
199	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> , 2020 , 29, 368-378	4	9
198	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> , 2020 , 22, 8	8.3	22
197	Multigene Panel Germline Testing of 1333 Czech Patients with Ovarian Cancer. <i>Cancers</i> , 2020 , 12,	6.6	8
196	Occupational exposure to ionizing radiation and risk of lymphoma subtypes: results of the Epilymph European case-control study. <i>Environmental Health</i> , 2020 , 19, 43	6	O
195	Association of Genomic Domains in and with Prostate Cancer Risk and Aggressiveness. <i>Cancer Research</i> , 2020 , 80, 624-638	10.1	22
194	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020 , 52, 56-73	36.3	56
193	Identification of Germline Mutations in Melanoma Patients with Early Onset, Double Primary Tumors, or Family Cancer History by NGS Analysis of 217 Genes. <i>Biomedicines</i> , 2020 , 8,	4.8	4
192	Polygenic risk scores and breast and epithelial ovarian cancer risks for carriers of BRCA1 and BRCA2 pathogenic variants. <i>Genetics in Medicine</i> , 2020 , 22, 1653-1666	8.1	34
191	Mendelian randomisation study of height and body mass index as modifiers of ovarian cancer risk in 22,588 BRCA1 and BRCA2 mutation carriers. <i>British Journal of Cancer</i> , 2019 , 121, 180-192	8.7	13
190	Associations Between Attention-Deficit/Hyperactivity Disorder and Various Eating Disorders: A Swedish Nationwide Population Study Using Multiple Genetically Informative Approaches. <i>Biological Psychiatry</i> , 2019 , 86, 577-586	7.9	24
189	Identification of deleterious germline CHEK2 mutations and their association with breast and ovarian cancer. <i>International Journal of Cancer</i> , 2019 , 145, 1782-1797	7.5	22
188	Genetically Determined Height and Risk of Non-hodgkin Lymphoma. Frontiers in Oncology, 2019, 9, 153	95.3	1
187	A Plasma-Derived Protein-Metabolite Multiplexed Panel for Early-Stage Pancreatic Cancer. <i>Journal of the National Cancer Institute</i> , 2019 , 111, 372-379	9.7	52

186	Genetic overlap between autoimmune diseases and non-Hodgkin lymphoma subtypes. <i>Genetic Epidemiology</i> , 2019 , 43, 844-863	2.6	15
185	Genome-wide association study identifies eight risk loci and implicates metabo-psychiatric origins for anorexia nervosa. <i>Nature Genetics</i> , 2019 , 51, 1207-1214	36.3	303
184	Sex specific associations in genome wide association analysis of renal cell carcinoma. <i>European Journal of Human Genetics</i> , 2019 , 27, 1589-1598	5.3	15
183	GAPPS - Gastric Adenocarcinoma and Proximal Polyposis of the Stomach Syndrome in 8 Families Tested at Masaryk Memorial Cancer Institute - Prevention and Prophylactic Gastrectomies. <i>Klinicka Onkologie</i> , 2019 , 32, 109-117	2	12
182	BAP1 Syndrome - Predisposition to Malignant Mesothelioma, Skin and Uveal Melanoma, Renal and Other Cancers. <i>Klinicka Onkologie</i> , 2019 , 32, 118-122	2	2
181	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	9.7	16
180	The influence of obesity-related factors in the etiology of renal cell carcinoma-A mendelian randomization study. <i>PLoS Medicine</i> , 2019 , 16, e1002724	11.6	38
179	Functional evaluation of variants of unknown significance in the BRCA2 gene identified in genetic testing. <i>Cancer Biology and Therapy</i> , 2019 , 20, 633-641	4.6	
178	Height and Body Mass Index as Modifiers of Breast Cancer Risk in BRCA1/2 Mutation Carriers: A Mendelian Randomization Study. <i>Journal of the National Cancer Institute</i> , 2019 , 111, 350-364	9.7	22
177	Mutational spectrum in a worldwide study of 29,700 families with BRCA1 or BRCA2 mutations. <i>Human Mutation</i> , 2018 , 39, 593-620	4.7	138
176	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. <i>Nature Communications</i> , 2018 , 9, 556	17.4	103
175	Thorough in silico and in vitro cDNA analysis of 21 putative BRCA1 and BRCA2 splice variants and a complex tandem duplication in BRCA2 allowing the identification of activated cryptic splice donor sites in BRCA2 exon 11. <i>Human Mutation</i> , 2018 , 39, 515-526	4.7	5
174	Genetic Contributions to The Association Between Adult Height and Head and Neck Cancer: A Mendelian Randomization Analysis. <i>Scientific Reports</i> , 2018 , 8, 4534	4.9	3
173	No association between global DNA methylation in peripheral blood and lung cancer risk in nonsmoking women: results from a multicenter study in Eastern and Central Europe. <i>European Journal of Cancer Prevention</i> , 2018 , 27, 1-5	2	8
172	HLA Class I and II Diversity Contributes to the Etiologic Heterogeneity of Non-Hodgkin Lymphoma Subtypes. <i>Cancer Research</i> , 2018 , 78, 4086-4096	10.1	18
171	Lung cancer and socioeconomic status in a pooled analysis of case-control studies. <i>PLoS ONE</i> , 2018 , 13, e0192999	3.7	54
170	The Influence of Number and Timing of Pregnancies on Breast Cancer Risk for Women With or Mutations. <i>JNCI Cancer Spectrum</i> , 2018 , 2, pky078	4.6	10
169	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> , 2018 , 2, pky023	4.6	13

(2016-2018)

Two high-risk susceptibility loci at 6p25.3 and 14q32.13 for Waldenstrth macroglobulinemia. <i>Nature Communications</i> , 2018 , 9, 4182	17.4	8
Validation of CZECANCA (CZEch CAncer paNel for Clinical Application) for targeted NGS-based analysis of hereditary cancer syndromes. <i>PLoS ONE</i> , 2018 , 13, e0195761	3.7	15
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	7·5	24
Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , 2017 , 77, 2789-2799	10.1	49
Young Adult and Usual Adult Body Mass Index and Multiple Myeloma Risk: A Pooled Analysis in the International Multiple Myeloma Consortium (IMMC). <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017 , 26, 876-885	4	21
Significant Locus and Metabolic Genetic Correlations Revealed in Genome-Wide Association Study of Anorexia Nervosa. <i>American Journal of Psychiatry</i> , 2017 , 174, 850-858	11.9	276
Risks of Breast, Ovarian, and Contralateral Breast Cancer for BRCA1 and BRCA2 Mutation Carriers. JAMA - Journal of the American Medical Association, 2017, 317, 2402-2416	27.4	1140
Genome-wide association study identifies multiple risk loci for renal cell carcinoma. <i>Nature Communications</i> , 2017 , 8, 15724	17.4	50
Exposure-Response Analyses of Asbestos and Lung Cancer Subtypes in a Pooled Analysis of Case-Control Studies. <i>Epidemiology</i> , 2017 , 28, 288-299	3.1	41
Loss of chromosome Y leads to down regulation of KDM5D and KDM6C epigenetic modifiers in clear cell renal cell carcinoma. <i>Scientific Reports</i> , 2017 , 7, 44876	4.9	24
Genetic Variants Related to Longer Telomere Length are Associated with Increased Risk of Renal Cell Carcinoma. <i>European Urology</i> , 2017 , 72, 747-754	10.2	27
Lupus-related single nucleotide polymorphisms and risk of diffuse large B-cell lymphoma. <i>Lupus Science and Medicine</i> , 2017 , 4, e000187	4.6	10
Association of breast cancer risk in BRCA1 and BRCA2 mutation carriers with genetic variants showing differential allelic expression: identification of a modifier of breast cancer risk at locus 11q22.3. <i>Breast Cancer Research and Treatment</i> , 2017 , 161, 117-134	4.4	15
A Pooled Analysis of Reproductive Factors, Exogenous Hormone Use, and Risk of Multiple Myeloma among Women in the International Multiple Myeloma Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016 , 25, 217-21	4	4
Occupational prestige, social mobility and the association with lung cancer in men. <i>BMC Cancer</i> , 2016 , 16, 395	4.8	12
Inheritance of deleterious mutations at both BRCA1 and BRCA2 in an international sample of 32,295 women. <i>Breast Cancer Research</i> , 2016 , 18, 112	8.3	25
Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016 , 7, 12675	17.4	53
Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. <i>Nature Communications</i> , 2016 , 7, 10933	17.4	70
	Validation of CZECANCA (CZEch CAncer paNel for Clinical Application) for targeted NGS-based analysis of hereditary cancer syndromes. PLoS ONE, 2018, 13, e0195761 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 Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. Cancer Research, 2017, 77, 2789-2799 Young Adult and Usual Adult Body Mass Index and Multiple Myeloma Risk: A Pooled Analysis in the International Multiple Myeloma Consortium (IMMC). Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 876-885 Significant Locus and Metabolic Genetic Correlations Revealed in Genome-Wide Association Study of Anorexia Nervosa. American Journal of Psychiatry, 2017, 174, 850-858 Risks of Breast, Ovarian, and Contralateral Breast Cancer for BRCA1 and BRCA2 Mutation Carriers. JAMA - Journal of the American Medical Association, 2017, 317, 2402-2416 Genome-wide association study identifies multiple risk loci for renal cell carcinoma. Nature Communications, 2017, 8, 15724 Exposure-Response Analyses of Asbestos and Lung Cancer Subtypes in a Pooled Analysis of Case-Control Studies. Epidemiology, 2017, 28, 288-299 Loss of chromosome Y leads to down regulation of KDM5D and KDM6C epigenetic modifiers in clear cell renal cell carcinoma. Scientific Reports, 2017, 7, 44876 Genetic Variants Related to Longer Telomere Length are Associated with Increased Risk of Renal Cell Carcinoma. European Urology, 2017, 72, 747-754 Lupus-related single nucleotide polymorphisms and risk of diffuse large B-cell lymphoma. Lupus Science and Medicine, 2017, 4, e000187 Association of breast cancer risk in BRCA1 and BRCA2 mutation carriers with genetic variants showing differential allelic expression: identification of a modifier of breast cancer risk at locus 11022.3. Breast Cancer Research and Treatment, 2017, 161, 117-134 A Pooled Analysis of Reproductive Factors, Exposence Hormone U	Validation of CZECANCA (CZEch CAncer paNel for Clinical Application) for targeted NCS-based analysis of hereditary cancer syndromes. PLoS ONE, 2018, 13, e0195761 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 Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. Cancer Research, 2017, 77, 2789-2799 Young Adult and Usual Adult Body Mass Index and Multiple Myeloma Risk: A Pooled Analysis in the International Multiple Myeloma Consortium (IMMC). Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 876-885 Significant Locus and Metabolic Genetic Correlations Revealed in Genome-Wide Association Study of Anorexia Nervosa. American Journal of Psychiatry, 2017, 174, 850-858 Risks of Breast, Ovarian, and Contralateral Breast Cancer for BRCA1 and BRCA2 Mutation Carriers. JAMA - Journal of the American Medical Association, 2017, 317, 2402-2416 Genome-wide association study identifies multiple risk loci for renal cell carcinoma. Nature Communications, 2017, 8, 15724 Exposure-Response Analyses of Asbestos and Lung Cancer Subtypes in a Pooled Analysis of Case-Control Studies. Epidemiology, 2017, 28, 288-299 Loss of chromosome Y leads to down regulation of KDM5D and KDM6C epigenetic modifiers in clear cell crant cell carcinoma. Scientific Reports, 2017, 7, 44876 Genetic Variants Related to Longer Telomere Length are Associated with Increased Risk of Renal Cell Carcinoma. European Urology, 2017, 72, 747-754 Lupus-related single nucleotide polymorphisms and risk of diffuse large B-cell lymphoma. Lupus Science and Medicine, 2017, 4, e000187 Association of breast cancer risk in BRCA1 and BRCA2 mutation carriers with genetic variants showing differential allelic expression: identification of a modifier of breast cancer risk al Locus 11922.3. Breast Cancer Research and Treatment, 2017, 161, 117-134 A Pooled Analysis of Reproductive Factors, Exogenous Hormone U

150	Genetically predicted longer telomere length is associated with increased risk of B-cell lymphoma subtypes. <i>Human Molecular Genetics</i> , 2016 , 25, 1663-76	5.6	39
149	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , 2016 , 48, 374-86	36.3	93
148	BRCA2 Polymorphic Stop Codon K3326X and the Risk of Breast, Prostate, and Ovarian Cancers. Journal of the National Cancer Institute, 2016 , 108,	9.7	65
147	KRAS mutations in blood circulating cell-free DNA: a pancreatic cancer case-control. <i>Oncotarget</i> , 2016 , 7, 78827-78840	3.3	47
146	Occupation and Risk of Non-Hodgkin Lymphoma and Its Subtypes: A Pooled Analysis from the InterLymph Consortium. <i>Environmental Health Perspectives</i> , 2016 , 124, 396-405	8.4	27
145	Identification and Functional Testing of ERCC2 Mutations in a Multi-national Cohort of Patients with Familial Breast- and Ovarian Cancer. <i>PLoS Genetics</i> , 2016 , 12, e1006248	6	16
144	Menstrual and Reproductive Factors, Hormone Use, and Risk of Pancreatic Cancer: Analysis From the International Pancreatic Cancer Case-Control Consortium (PanC4). <i>Pancreas</i> , 2016 , 45, 1401-1410	2.6	10
143	Lung Cancer Among Firefighters: Smoking-Adjusted Risk Estimates in a Pooled Analysis of Case-Control Studies. <i>Journal of Occupational and Environmental Medicine</i> , 2016 , 58, 1137-1143	2	11
142	Meta-analysis of genome-wide association studies reveals genetic overlap between Hodgkin lymphoma and multiple sclerosis. <i>International Journal of Epidemiology</i> , 2016 , 45, 728-40	7.8	16
141	Male breast cancer in BRCA1 and BRCA2 mutation carriers: pathology data from the Consortium of Investigators of Modifiers of BRCA1/2. <i>Breast Cancer Research</i> , 2016 , 18, 15	8.3	58
140	A pooled analysis of cigarette smoking and risk of multiple myeloma from the international multiple myeloma consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 631-4	4	11
139	Effect of occupational exposures on lung cancer susceptibility: a study of gene-environment interaction analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 570-9	4	10
138	Associations of non-Hodgkin Lymphoma (NHL) risk with autoimmune conditions according to putative NHL loci. <i>American Journal of Epidemiology</i> , 2015 , 181, 406-21	3.8	42
137	Common variation at 2p13.3, 3q29, 7p13 and 17q25.1 associated with susceptibility to pancreatic cancer. <i>Nature Genetics</i> , 2015 , 47, 911-6	36.3	171
136	Association of type and location of BRCA1 and BRCA2 mutations with risk of breast and ovarian cancer. <i>JAMA - Journal of the American Medical Association</i> , 2015 , 313, 1347-61	27.4	286
135	Lung cancer risk among cooks when accounting for tobacco smoking: a pooled analysis of case-control studies from Europe, Canada, New Zealand, and China. <i>Journal of Occupational and Environmental Medicine</i> , 2015 , 57, 202-9	2	8
134	Identification of lung cancer histology-specific variants applying Bayesian framework variant prioritization approaches within the TRICL and ILCCO consortia. <i>Carcinogenesis</i> , 2015 , 36, 1314-26	4.6	8
133	Lung cancer risk among bricklayers in a pooled analysis of case-control studies. <i>International Journal of Cancer</i> , 2015 , 136, 360-71	7.5	22

132	Analysis of Heritability and Shared Heritability Based on Genome-Wide Association Studies for Thirteen Cancer Types. <i>Journal of the National Cancer Institute</i> , 2015 , 107, djv279	9.7	107
131	Assessing associations between the AURKA-HMMR-TPX2-TUBG1 functional module and breast cancer risk in BRCA1/2 mutation carriers. <i>PLoS ONE</i> , 2015 , 10, e0120020	3.7	26
130	The 12p13.33/RAD52 locus and genetic susceptibility to squamous cell cancers of upper aerodigestive tract. <i>PLoS ONE</i> , 2015 , 10, e0117639	3.7	10
129	A genome-wide association study of marginal zone lymphoma shows association to the HLA region. <i>Nature Communications</i> , 2015 , 6, 5751	17.4	44
128	A rare truncating BRCA2 variant and genetic susceptibility to upper aerodigestive tract cancer. <i>Journal of the National Cancer Institute</i> , 2015 , 107,	9.7	27
127	A Novel Risk Locus at 6p21.3 for Epstein-Barr Virus-Positive Hodgkin Lymphoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 1838-43	4	16
126	A systematic investigation of the contribution of genetic variation within the MHC region to HPV seropositivity. <i>Human Molecular Genetics</i> , 2015 , 24, 2681-8	5.6	8
125	Lung cancer among coal miners, ore miners and quarrymen: smoking-adjusted risk estimates from the synergy pooled analysis of case-control studies. <i>Scandinavian Journal of Work, Environment and Health</i> , 2015 , 41, 467-77	4.3	26
124	No causal association identified for human papillomavirus infections in lung cancer. <i>Cancer Research</i> , 2014 , 74, 3525-34	10.1	27
123	Effect modification of the association of cumulative exposure and cancer risk by intensity of exposure and time since exposure cessation: a flexible method applied to cigarette smoking and lung cancer in the SYNERGY Study. <i>American Journal of Epidemiology</i> , 2014 , 179, 290-8	3.8	33
122	Variation in genomic landscape of clear cell renal cell carcinoma across Europe. <i>Nature Communications</i> , 2014 , 5, 5135	17.4	123
121	Etiologic heterogeneity among non-Hodgkin lymphoma subtypes: the InterLymph Non-Hodgkin Lymphoma Subtypes Project. <i>Journal of the National Cancer Institute Monographs</i> , 2014 , 2014, 130-44	4.8	199
120	Genome-wide association study identifies multiple susceptibility loci for diffuse large B cell lymphoma. <i>Nature Genetics</i> , 2014 , 46, 1233-8	36.3	108
119	Genome-wide association study identifies five susceptibility loci for follicular lymphoma outside the HLA region. <i>American Journal of Human Genetics</i> , 2014 , 95, 462-71	11	74
118	Physical activity and risk of pancreatic cancer in a central European multicenter case-control study. <i>Cancer Causes and Control</i> , 2014 , 25, 669-81	2.8	13
117	Rare variants of large effect in BRCA2 and CHEK2 affect risk of lung cancer. <i>Nature Genetics</i> , 2014 , 46, 736-41	36.3	228
116	Is previous respiratory disease a risk factor for lung cancer?. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2014 , 190, 549-59	10.2	73
115	Exposure to secondhand tobacco smoke and lung cancer by histological type: a pooled analysis of the International Lung Cancer Consortium (ILCCO). <i>International Journal of Cancer</i> , 2014 , 135, 1918-30	7.5	69

114	Occupational exposure to immunologically active agents and risk for lymphoma: the European Epilymph case-control study. <i>Cancer Epidemiology</i> , 2013 , 37, 378-84	2.8	8
113	Tamoxifen and risk of contralateral breast cancer for BRCA1 and BRCA2 mutation carriers. <i>Journal of Clinical Oncology</i> , 2013 , 31, 3091-9	2.2	118
112	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, 371-84, 384e1-2	36.3	422
111	Multiple myeloma and occupation: a pooled analysis by the International Multiple Myeloma Consortium. <i>Cancer Epidemiology</i> , 2013 , 37, 300-5	2.8	16
110	Genome-wide association study identifies multiple risk loci for chronic lymphocytic leukemia. <i>Nature Genetics</i> , 2013 , 45, 868-76	36.3	147
109	Welding and lung cancer in a pooled analysis of case-control studies. <i>American Journal of Epidemiology</i> , 2013 , 178, 1513-25	3.8	35
108	Genome-wide association study in BRCA1 mutation carriers identifies novel loci associated with breast and ovarian cancer risk. <i>PLoS Genetics</i> , 2013 , 9, e1003212	6	209
107	Lung cancer risk among hairdressers: a pooled analysis of case-control studies conducted between 1985 and 2010. <i>American Journal of Epidemiology</i> , 2013 , 178, 1355-65	3.8	8
106	A pooled analysis of alcohol consumption and risk of multiple myeloma in the international multiple myeloma consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013 , 22, 1620-7	4	17
105	Lung cancer risk among bakers, pastry cooks and confectionary makers: the SYNERGY study. <i>Occupational and Environmental Medicine</i> , 2013 , 70, 810-4	2.1	10
104	Lymphoma risk in livestock farmers: results of the Epilymph study. <i>International Journal of Cancer</i> , 2013 , 132, 2613-8	7.5	9
103	An investigation of risk factors for renal cell carcinoma by histologic subtype in two case-control studies. <i>International Journal of Cancer</i> , 2013 , 132, 2640-7	7.5	50
102	Lymphoma risk and occupational exposure to pesticides: results of the Epilymph study. <i>Occupational and Environmental Medicine</i> , 2013 , 70, 91-8	2.1	69
101	Exposure to UV radiation and risk of Hodgkin lymphoma: a pooled analysis. <i>Blood</i> , 2013 , 122, 3492-9	2.2	29
100	Integrative genome-wide gene expression profiling of clear cell renal cell carcinoma in Czech Republic and in the United States. <i>PLoS ONE</i> , 2013 , 8, e57886	3.7	70
99	Reproductive factors and lymphoid neoplasms in Europe: findings from the EpiLymph case-control study. <i>Cancer Causes and Control</i> , 2012 , 23, 195-206	2.8	16
98	PRRC2A and BCL2L11 gene variants influence risk of non-Hodgkin lymphoma: results from the InterLymph consortium. <i>Blood</i> , 2012 , 120, 4645-8	2.2	23
97	Genome-wide association study of classical Hodgkin lymphoma and Epstein-Barr virus status-defined subgroups. <i>Journal of the National Cancer Institute</i> , 2012 , 104, 240-53	9.7	117

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