

Nuria Malats

List of Publications by Citations

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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.

228
papers

12,501
citations

66
h-index

105
g-index

239
ext. papers

14,875
ext. citations

8.2
avg, IF

5.46
L-index

#	Paper	IF	Citations
228	NAT2 slow acetylation, GSTM1 null genotype, and risk of bladder cancer: results from the Spanish Bladder Cancer Study and meta-analyses. <i>Lancet, The</i> , 2005 , 366, 649-59	40	483
227	Detectable clonal mosaicism and its relationship to aging and cancer. <i>Nature Genetics</i> , 2012 , 44, 651-8	36.3	409
226	A multi-stage genome-wide association study of bladder cancer identifies multiple susceptibility loci. <i>Nature Genetics</i> , 2010 , 42, 978-84	36.3	408
225	Prognosis Research Strategy (PROGRESS) 2: prognostic factor research. <i>PLoS Medicine</i> , 2013 , 10, e1001380	30.6	382
224	Bladder cancer and exposure to water disinfection by-products through ingestion, bathing, showering, and swimming in pools. <i>American Journal of Epidemiology</i> , 2007 , 165, 148-56	3.8	382
223	Comprehensive Transcriptional Analysis of Early-Stage Urothelial Carcinoma. <i>Cancer Cell</i> , 2016 , 30, 27-42	24.3	325
222	A Consensus Molecular Classification of Muscle-invasive Bladder Cancer. <i>European Urology</i> , 2020 , 77, 420-433	10.2	309
221	Prospective study of FGFR3 mutations as a prognostic factor in nonmuscle invasive urothelial bladder carcinomas. <i>Journal of Clinical Oncology</i> , 2006 , 24, 3664-71	2.2	256
220	P53 as a prognostic marker for bladder cancer: a meta-analysis and review. <i>Lancet Oncology, The</i> , 2005 , 6, 678-86	21.7	244
219	Genome-wide association study identifies multiple susceptibility loci for pancreatic cancer. <i>Nature Genetics</i> , 2014 , 46, 994-1000	36.3	226
218	Epidemiology of urinary bladder cancer: from tumor development to patient's death. <i>World Journal of Urology</i> , 2007 , 25, 285-95	4	203
217	Genomic DNA hypomethylation as a biomarker for bladder cancer susceptibility in the Spanish Bladder Cancer Study: a case-control study. <i>Lancet Oncology, The</i> , 2008 , 9, 359-66	21.7	193
216	Recurrent inactivation of STAG2 in bladder cancer is not associated with aneuploidy. <i>Nature Genetics</i> , 2013 , 45, 1464-9	36.3	186
215	Exocrine pancreatic cancer: symptoms at presentation and their relation to tumour site and stage. <i>Clinical and Translational Oncology</i> , 2005 , 7, 189-97	3.6	179
214	PIK3CA mutations are an early genetic alteration associated with FGFR3 mutations in superficial papillary bladder tumors. <i>Cancer Research</i> , 2006 , 66, 7401-4	10.1	175
213	Telomerase reverse transcriptase promoter mutations in bladder cancer: high frequency across stages, detection in urine, and lack of association with outcome. <i>European Urology</i> , 2014 , 65, 360-6	10.2	166
212	Gene expression signatures predict outcome in non-muscle-invasive bladder carcinoma: a multicenter validation study. <i>Clinical Cancer Research</i> , 2007 , 13, 3545-51	12.9	164

211	Polymorphisms in GSTT1, GSTZ1, and CYP2E1, disinfection by-products, and risk of bladder cancer in Spain. <i>Environmental Health Perspectives</i> , 2010 , 118, 1545-50	8.4	162
210	Genome-wide association study identifies two susceptibility loci for osteosarcoma. <i>Nature Genetics</i> , 2013 , 45, 799-803	36.3	156
209	Prognostic factors and risk groups in T1G3 non-muscle-invasive bladder cancer patients initially treated with Bacillus Calmette-Guérin: results of a retrospective multicenter study of 2451 patients. <i>European Urology</i> , 2015 , 67, 74-82	10.2	149
208	Serum concentrations of organochlorine compounds and K-ras mutations in exocrine pancreatic cancer. PANKRAS II Study Group. <i>Lancet, The</i> , 1999 , 354, 2125-9	40	146
207	Genetic variation in the nucleotide excision repair pathway and bladder cancer risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006 , 15, 536-42	4	128
206	GATA6 regulates EMT and tumour dissemination, and is a marker of response to adjuvant chemotherapy in pancreatic cancer. <i>Gut</i> , 2017 , 66, 1665-1676	19.2	125
205	Smoking and bladder cancer in Spain: effects of tobacco type, timing, environmental tobacco smoke, and gender. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006 , 15, 1348-54	4	121
204	Identification of a Three-Biomarker Panel in Urine for Early Detection of Pancreatic Adenocarcinoma. <i>Clinical Cancer Research</i> , 2015 , 21, 3512-21	12.9	120
203	Circulating tumor cells (Ctc) and kras mutant circulating free Dna (cfdna) detection in peripheral blood as biomarkers in patients diagnosed with exocrine pancreatic cancer. <i>BMC Cancer</i> , 2015 , 15, 797	4.8	116
202	Epidemiology of bladder cancer. <i>Hematology/Oncology Clinics of North America</i> , 2015 , 29, 177-89, vii	3.1	109
201	Large-scale evaluation of candidate genes identifies associations between VEGF polymorphisms and bladder cancer risk. <i>PLoS Genetics</i> , 2007 , 3, e29	6	109
200	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
199	Genetic variation in the base excision repair pathway and bladder cancer risk. <i>Human Genetics</i> , 2007 , 121, 233-42	6.3	107
198	FGFR3 and Tp53 mutations in T1G3 transitional bladder carcinomas: independent distribution and lack of association with prognosis. <i>Clinical Cancer Research</i> , 2005 , 11, 5444-50	12.9	106
197	Association of germline variants in the APOBEC3 region with cancer risk and enrichment with APOBEC-signature mutations in tumors. <i>Nature Genetics</i> , 2016 , 48, 1330-1338	36.3	104
196	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. <i>Nature Communications</i> , 2018 , 9, 556	17.4	103
195	Genome-wide association study identifies multiple loci associated with bladder cancer risk. <i>Human Molecular Genetics</i> , 2014 , 23, 1387-98	5.6	101
194	Mosaic uniparental disomies and aneuploidies as large structural variants of the human genome. <i>American Journal of Human Genetics</i> , 2010 , 87, 129-38	11	100

193	Statistical consideration for clinical biomarker research in bladder cancer. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2010 , 28, 389-400	2.8	99
192	Food, nutrient and heterocyclic amine intake and the risk of bladder cancer. <i>European Journal of Cancer</i> , 2007 , 43, 1731-40	7.5	99
191	Polymorphisms in DNA repair genes, smoking, and bladder cancer risk: findings from the international consortium of bladder cancer. <i>Cancer Research</i> , 2009 , 69, 6857-64	10.1	94
190	Searching for urine biomarkers of bladder cancer recurrence using a liquid chromatography-mass spectrometry and capillary electrophoresis-mass spectrometry metabolomics approach. <i>Journal of Chromatography A</i> , 2013 , 1318, 163-70	4.5	91
189	Mosaic loss of chromosome Y is associated with common variation near TCL1A. <i>Nature Genetics</i> , 2016 , 48, 563-8	36.3	87
188	Cystic fibrosis transmembrane regulator (CFTR) DeltaF508 mutation and 5T allele in patients with chronic pancreatitis and exocrine pancreatic cancer. PANKRAS II Study Group. <i>Gut</i> , 2001 , 48, 70-4	19.2	86
187	Resection of pancreatic cancer in Europe and USA: an international large-scale study highlighting large variations. <i>Gut</i> , 2019 , 68, 130-139	19.2	86
186	AUC-RF: a new strategy for genomic profiling with random forest. <i>Human Heredity</i> , 2011 , 72, 121-32	1.1	85
185	Evaluation of genetic variation in the double-strand break repair pathway and bladder cancer risk. <i>Carcinogenesis</i> , 2007 , 28, 1788-93	4.6	83
184	Common genetic polymorphisms modify the effect of smoking on absolute risk of bladder cancer. <i>Cancer Research</i> , 2013 , 73, 2211-20	10.1	82
183	A genome-wide association study of bladder cancer identifies a new susceptibility locus within SLC14A1, a urea transporter gene on chromosome 18q12.3. <i>Human Molecular Genetics</i> , 2011 , 20, 4282-9	5.6	82
182	An unusual suspect: an uncommon human-specific synonymous coding variant within the UGT1A6 gene explains a GWAS signal and protects against bladder cancer. <i>Genome Biology</i> , 2011 , 12,	18.3	78
181	Selenium and bladder cancer risk: a meta-analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010 , 19, 2407-15	4	78
180	Characterization of large structural genetic mosaicism in human autosomes. <i>American Journal of Human Genetics</i> , 2015 , 96, 487-97	11	77
179	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. <i>Human Molecular Genetics</i> , 2014 , 23, 6616-33	5.6	77
178	Occurrence, trends and environment etiology of pancreatic cancer. <i>Scandinavian Journal of Work, Environment and Health</i> , 1998 , 24, 165-74	4.3	77
177	Validation of a DNA Methylation-Mutation Urine Assay to Select Patients with Hematuria for Cystoscopy. <i>Journal of Urology</i> , 2017 , 197, 590-595	2.5	76
176	Bladder cancer risk and genetic variation in AKR1C3 and other metabolizing genes. <i>Carcinogenesis</i> , 2008 , 29, 1955-62	4.6	76

175	Assessing interactions between the associations of common genetic susceptibility variants, reproductive history and body mass index with breast cancer risk in the breast cancer association consortium: a combined case-control study. <i>Breast Cancer Research</i> , 2010 , 12, R110	8.3	74
174	Improving strategies for detecting genetic patterns of disease susceptibility in association studies. <i>Statistics in Medicine</i> , 2008 , 27, 6532-46	2.3	74
173	mbmdr: an R package for exploring gene-gene interactions associated with binary or quantitative traits. <i>Bioinformatics</i> , 2010 , 26, 2198-9	7.2	73
172	The impact of re-transurethral resection on clinical outcomes in a large multicentre cohort of patients with T1 high-grade/Grade 3 bladder cancer treated with bacille Calmette-Guérin. <i>BJU International</i> , 2016 , 118, 44-52	5.6	72
171	Multiple oncogenic mutations and clonal relationship in spatially distinct benign human epidermal tumors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 20780-5	11.5	71
170	Risk of bladder cancer associated with family history of cancer: do low-penetrance polymorphisms account for the increase in risk?. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007 , 16, 1595-600	4	70
169	Common genetic variants in the PSCA gene influence gene expression and bladder cancer risk. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, 4974-9	11.5	69
168	Pancreatic cancer risk and levels of trace elements. <i>Gut</i> , 2012 , 61, 1583-8	19.2	68
167	Risk of pancreatic cancer in breast cancer families from the breast cancer family registry. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013 , 22, 803-11	4	67
166	Winner's Curse Correction and Variable Thresholding Improve Performance of Polygenic Risk Modeling Based on Genome-Wide Association Study Summary-Level Data. <i>PLoS Genetics</i> , 2016 , 12, e1006493	6	67
165	Inflammatory biomarkers and bladder cancer prognosis: a systematic review. <i>European Urology</i> , 2014 , 66, 1078-91	10.2	66
164	Genetic and non-genetic predictors of LINE-1 methylation in leukocyte DNA. <i>Environmental Health Perspectives</i> , 2013 , 121, 650-6	8.4	66
163	Three new pancreatic cancer susceptibility signals identified on chromosomes 1q32.1, 5p15.33 and 8q24.21. <i>Oncotarget</i> , 2016 , 7, 66328-66343	3.3	66
162	Cigarette smoking and K-ras mutations in pancreas, lung and colorectal adenocarcinomas: etiopathogenic similarities, differences and paradoxes. <i>Mutation Research - Reviews in Mutation Research</i> , 2009 , 682, 83-93	7	63
161	Nitrate in drinking water and bladder cancer risk in Spain. <i>Environmental Research</i> , 2015 , 137, 299-307	7.9	62
160	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , 2016 , 7, 11843	17.4	59
159	Mapping of the UGT1A locus identifies an uncommon coding variant that affects mRNA expression and protects from bladder cancer. <i>Human Molecular Genetics</i> , 2012 , 21, 1918-30	5.6	58
158	Ki-ras mutations in exocrine pancreatic cancer: association with clinico-pathological characteristics and with tobacco and alcohol consumption. PANK-ras I Project Investigators. <i>International Journal of Cancer</i> , 1997 , 70, 661-7	7.5	58

157	Transcriptional regulation by NR5A2 links differentiation and inflammation in the pancreas. <i>Nature</i> , 2018 , 554, 533-537	50.4	57
156	Genome-wide association study identifies inversion in the locus to modify risk for alcoholic and non-alcoholic chronic pancreatitis. <i>Gut</i> , 2018 , 67, 1855-1863	19.2	54
155	FGFR3, TERT and OTX1 as a Urinary Biomarker Combination for Surveillance of Patients with Bladder Cancer in a Large Prospective Multicenter Study. <i>Journal of Urology</i> , 2017 , 197, 1410-1418	2.5	53
154	Total fluid and water consumption and the joint effect of exposure to disinfection by-products on risk of bladder cancer. <i>Environmental Health Perspectives</i> , 2007 , 115, 1569-72	8.4	53
153	Polymorphisms in one-carbon metabolism and trans-sulfuration pathway genes and susceptibility to bladder cancer. <i>International Journal of Cancer</i> , 2007 , 120, 2452-8	7.5	53
152	The p53 pathway and outcome among patients with T1G3 bladder tumors. <i>Clinical Cancer Research</i> , 2006 , 12, 6029-36	12.9	53
151	Genetic susceptibility to distinct bladder cancer subphenotypes. <i>European Urology</i> , 2010 , 57, 283-92	10.2	52
150	Prognostic Impact of a 12-gene Progression Score in Non-muscle-invasive Bladder Cancer: A Prospective Multicentre Validation Study. <i>European Urology</i> , 2017 , 72, 461-469	10.2	51
149	Air pollution and risk of urinary bladder cancer in a case-control study in Spain. <i>Occupational and Environmental Medicine</i> , 2008 , 65, 56-60	2.1	51
148	Caffeic acid phenethyl ester induces apoptosis of human pancreatic cancer cells involving caspase and mitochondrial dysfunction. <i>Pancreatology</i> , 2008 , 8, 566-76	3.8	50
147	Association between coffee drinking and K-ras mutations in exocrine pancreatic cancer. PANKRAS II Study Group. <i>Journal of Epidemiology and Community Health</i> , 1999 , 53, 702-9	5.1	50
146	Integration Analysis of Three Omics Data Using Penalized Regression Methods: An Application to Bladder Cancer. <i>PLoS Genetics</i> , 2015 , 11, e1005689	6	49
145	Assessment of copy number variation using the Illumina Infinium 1M SNP-array: a comparison of methodological approaches in the Spanish Bladder Cancer/EPICURO study. <i>Human Mutation</i> , 2011 , 32, 240-8	4.7	49
144	Molecular Markers Increase Precision of the European Association of Urology Non-Muscle-Invasive Bladder Cancer Progression Risk Groups. <i>Clinical Cancer Research</i> , 2018 , 24, 1586-1593	12.9	48
143	Occupation and bladder cancer in a hospital-based case-control study in Spain. <i>Occupational and Environmental Medicine</i> , 2008 , 65, 347-53	2.1	48
142	Assessment of lifetime exposure to trihalomethanes through different routes. <i>Occupational and Environmental Medicine</i> , 2006 , 63, 273-7	2.1	47
141	Occupational exposure to dyes, metals, polycyclic aromatic hydrocarbons and other agents and K-ras activation in human exocrine pancreatic cancer. <i>International Journal of Cancer</i> , 2003 , 107, 635-41	7.5	47
140	Challenges in the Integration of Omics and Non-Omics Data. <i>Genes</i> , 2019 , 10,	4.2	46

139	A single nucleotide polymorphism tags variation in the arylamine N-acetyltransferase 2 phenotype in populations of European background. <i>Pharmacogenetics and Genomics</i> , 2011 , 21, 231-6	1.9	44
138	Genetic variations in the sonic hedgehog pathway affect clinical outcomes in non-muscle-invasive bladder cancer. <i>Cancer Prevention Research</i> , 2010 , 3, 1235-45	3.2	42
137	Gender-related differences in clinical and pathological characteristics and therapy of bladder cancer. <i>European Urology</i> , 2003 , 43, 53-62	10.2	42
136	ARID1A alterations are associated with FGFR3-wild type, poor-prognosis, urothelial bladder tumors. <i>PLoS ONE</i> , 2013 , 8, e62483	3.7	41
135	TGFB1 and TGFBR1 polymorphic variants in relationship to bladder cancer risk and prognosis. <i>International Journal of Cancer</i> , 2009 , 124, 608-13	7.5	41
134	Use of analgesics and nonsteroidal anti-inflammatory drugs, genetic predisposition, and bladder cancer risk in Spain. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006 , 15, 1696-702	4	41
133	Occupational exposure to organic solvents and K-ras mutations in exocrine pancreatic cancer. <i>Carcinogenesis</i> , 2002 , 23, 101-6	4.6	40
132	The efficacy of BCG TICE and BCG Connaught in a cohort of 2,099 patients with T1G3 non-muscle-invasive bladder cancer. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2016 , 34, 484.e19-484.e25	2.8	39
131	Reduced risk of pancreatic cancer associated with asthma and nasal allergies. <i>Gut</i> , 2017 , 66, 314-322	19.2	37
130	Transcriptome analysis of pancreatic cancer reveals a tumor suppressor function for HNF1A. <i>Carcinogenesis</i> , 2014 , 35, 2670-8	4.6	37
129	Screening for bladder cancer: a perspective. <i>World Journal of Urology</i> , 2008 , 26, 13-8	4	35
128	Risk prediction scores for recurrence and progression of non-muscle invasive bladder cancer: an international validation in primary tumours. <i>PLoS ONE</i> , 2014 , 9, e96849	3.7	34
127	Genome-wide interaction study of smoking and bladder cancer risk. <i>Carcinogenesis</i> , 2014 , 35, 1737-44	4.6	33
126	Large-scale pathway-based analysis of bladder cancer genome-wide association data from five studies of European background. <i>PLoS ONE</i> , 2012 , 7, e29396	3.7	33
125	PanGen-Fam: Spanish registry of hereditary pancreatic cancer. <i>European Journal of Cancer</i> , 2015 , 51, 1911-7	7.5	32
124	Coffee consumption, genetic susceptibility and bladder cancer risk. <i>Cancer Causes and Control</i> , 2009 , 20, 121-7	2.8	32
123	Correcting serum concentrations of organochlorine compounds by lipids: alternatives to the organochlorine/total lipids ratio. <i>Environment International</i> , 2009 , 35, 1080-5	12.9	32
122	Evidence for an intensity-dependent interaction of NAT2 acetylation genotype and cigarette smoking in the Spanish Bladder Cancer Study. <i>International Journal of Epidemiology</i> , 2007 , 36, 236-41	7.8	31

121	Validity of the hospital discharge diagnosis in epidemiologic studies of biliopancreatic pathology. PANKRAS II Study Group. <i>European Journal of Epidemiology</i> , 2000 , 16, 533-41	12.1	31
120	A large-scale assessment of two-way SNP interactions in breast cancer susceptibility using 46,450 cases and 42,461 controls from the breast cancer association consortium. <i>Human Molecular Genetics</i> , 2014 , 23, 1934-46	5.6	28
119	Urinary pH, cigarette smoking and bladder cancer risk. <i>Carcinogenesis</i> , 2011 , 32, 843-7	4.6	28
118	Plasma 25-hydroxyvitamin D(3) and bladder cancer risk according to tumor stage and FGFR3 status: a mechanism-based epidemiological study. <i>Journal of the National Cancer Institute</i> , 2012 , 104, 1897-904	9.7	28
117	Confirmation of 5p12 as a susceptibility locus for progesterone-receptor-positive, lower grade breast cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011 , 20, 2222-31	4	27
116	Analysis of molecular intra-patient variation and delineation of a prognostic 12-gene signature in non-muscle invasive bladder cancer; technology transfer from microarrays to PCR. <i>British Journal of Cancer</i> , 2012 , 107, 1392-8	8.7	27
115	Vitamin D metabolic pathway genes and pancreatic cancer risk. <i>PLoS ONE</i> , 2015 , 10, e0117574	3.7	26
114	Family history of cancer and germline BRCA2 mutations in sporadic exocrine pancreatic cancer. <i>Gut</i> , 2002 , 50, 653-7	19.2	26
113	Does increased urination frequency protect against bladder cancer?. <i>International Journal of Cancer</i> , 2008 , 123, 1644-8	7.5	25
112	A Transcriptome-Wide Association Study Identifies Novel Candidate Susceptibility Genes for Pancreatic Cancer. <i>Journal of the National Cancer Institute</i> , 2020 , 112, 1003-1012	9.7	25
111	Identification of new genetic susceptibility loci for breast cancer through consideration of gene-environment interactions. <i>Genetic Epidemiology</i> , 2014 , 38, 84-93	2.6	24
110	An integrated multi-omics analysis identifies prognostic molecular subtypes of non-muscle-invasive bladder cancer. <i>Nature Communications</i> , 2021 , 12, 2301	17.4	24
109	Modification of Occupational Exposures on Bladder Cancer Risk by Common Genetic Polymorphisms. <i>Journal of the National Cancer Institute</i> , 2015 , 107,	9.7	23
108	Timing of blood extraction in epidemiologic and proteomic studies: results and proposals from the PANKRAS II Study. <i>European Journal of Epidemiology</i> , 2007 , 22, 577-88	12.1	23
107	Medical conditions in patients with pancreatic and biliary diseases: validity and agreement between data from questionnaires and medical records. PANKRAS II Study Group. <i>Digestive Diseases and Sciences</i> , 1999 , 44, 2469-77	4	23
106	Food and nutrient intakes and K-ras mutations in exocrine pancreatic cancer. <i>Journal of Epidemiology and Community Health</i> , 2007 , 61, 641-9	5.1	22
105	Lifetime history of tobacco consumption and K-ras mutations in exocrine pancreatic cancer. <i>Pancreas</i> , 2007 , 35, 135-41	2.6	22
104	LINE-1 methylation in granulocyte DNA and trihalomethane exposure is associated with bladder cancer risk. <i>Epigenetics</i> , 2014 , 9, 1532-9	5.7	21

103	Risk of pancreatic cancer associated with family history of cancer and other medical conditions by accounting for smoking among relatives. <i>International Journal of Epidemiology</i> , 2018 , 47, 473-483	7.8	20
102	Identification of a novel susceptibility locus at 13q34 and refinement of the 20p12.2 region as a multi-signal locus associated with bladder cancer risk in individuals of European ancestry. <i>Human Molecular Genetics</i> , 2016 , 25, 1203-14	5.6	20
101	A Multicenter Trial Defining a Serum Protein Signature Associated with Pancreatic Ductal Adenocarcinoma. <i>International Journal of Proteomics</i> , 2015 , 2015, 587250		20
100	Gene-environment interactions in pancreatic cancer. <i>Pancreatology</i> , 2001 , 1, 472-6	3.8	20
99	Diagnostic certainty and potential for misclassification in exocrine pancreatic cancer. PANKRAS I Project Investigations. <i>Journal of Clinical Epidemiology</i> , 1994 , 47, 1069-79	5.7	20
98	Cyclooxygenase-2 expression in bladder cancer and patient prognosis: results from a large clinical cohort and meta-analysis. <i>PLoS ONE</i> , 2012 , 7, e45025	3.7	20
97	Framework for the Integration of Genomics, Epigenomics and Transcriptomics in Complex Diseases. <i>Human Heredity</i> , 2015 , 79, 124-36	1.1	19
96	The 19q12 bladder cancer GWAS signal: association with cyclin E function and aggressive disease. <i>Cancer Research</i> , 2014 , 74, 5808-18	10.1	19
95	Select your SNPs (SYSNPs): a web tool for automatic and massive selection of SNPs. <i>International Journal of Data Mining and Bioinformatics</i> , 2012 , 6, 324-34	0.5	19
94	Recurrence, progression and cancer-specific mortality according to stage at re-TUR in T1G3 bladder cancer patients treated with BCG: not as bad as previously thought. <i>World Journal of Urology</i> , 2018 , 36, 1621-1627	4	18
93	Bladder cancer and seroreactivity to BK, JC and Merkel cell polyomaviruses: the Spanish bladder cancer study. <i>International Journal of Cancer</i> , 2013 , 133, 597-603	7.5	18
92	The influence of lipid and lifestyle factors upon correlations between highly prevalent organochlorine compounds in patients with exocrine pancreatic cancer. <i>Environment International</i> , 2007 , 33, 946-54	12.9	18
91	Learning from case reports: diagnostic issues in an epidemiologic study of pancreatic cancer. <i>Journal of Clinical Epidemiology</i> , 1998 , 51, 1215-21	5.7	17
90	The International Bladder Cancer Bank: proposal for a new study concept. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2004 , 22, 277-84	2.8	17
89	Coffee, pancreatic cancer, and K-ras mutations: updating the research agenda. <i>Journal of Epidemiology and Community Health</i> , 2000 , 54, 656-9	5.1	17
88	Deciphering the complex interplay between pancreatic cancer, diabetes mellitus subtypes and obesity/BMI through causal inference and mediation analyses. <i>Gut</i> , 2021 , 70, 319-329	19.2	16
87	A comprehensive analysis of candidate genes in familial pancreatic cancer families reveals a high frequency of potentially pathogenic germline variants. <i>EBioMedicine</i> , 2020 , 53, 102675	8.8	16
86	Bladder Cancer Genetic Susceptibility. A Systematic Review. <i>Bladder Cancer</i> , 2018 , 4, 215-226	1	16

85	LINE-1 methylation in leukocyte DNA, interaction with phosphatidylethanolamine N-methyltransferase variants and bladder cancer risk. <i>British Journal of Cancer</i> , 2014 , 110, 2123-30	8.7	16
84	Biological and statistical approaches for modeling exposure to specific trihalomethanes and bladder cancer risk. <i>American Journal of Epidemiology</i> , 2013 , 178, 652-60	3.8	16
83	Socioeconomic status and exposure to disinfection by-products in drinking water in Spain. <i>Environmental Health</i> , 2011 , 10, 18	6	16
82	A systems approach identifies time-dependent associations of multimorbidities with pancreatic cancer risk. <i>Annals of Oncology</i> , 2017 , 28, 1618-1624	10.3	15
81	Pancreatic Cancer Risk in Relation to Lifetime Smoking Patterns, Tobacco Type, and Dose-Response Relationships. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020 , 29, 1009-1018	4	15
80	Toward the integration of Omics data in epidemiological studies: still a "long and winding road". <i>Genetic Epidemiology</i> , 2016 , 40, 558-569	2.6	15
79	Next generation modeling in GWAS: comparing different genetic architectures. <i>Human Genetics</i> , 2014 , 133, 1235-53	6.3	15
78	The relative influence of diet and serum concentrations of organochlorine compounds on K-ras mutations in exocrine pancreatic cancer. <i>Chemosphere</i> , 2010 , 79, 686-97	8.4	15
77	Do we believe what patients say about their neoplastic symptoms? An analysis of factors that influence the interviewer's judgement. <i>European Journal of Epidemiology</i> , 1996 , 12, 553-62	12.1	15
76	Review of studies of selected metabolic polymorphisms and cancer. <i>Iarc (international Agency for Research on Cancer) Scientific Publications</i> , 1999 , 323-93		15
75	Genome-wide CNV analysis replicates the association between GSTM1 deletion and bladder cancer: a support for using continuous measurement from SNP-array data. <i>BMC Genomics</i> , 2012 , 13, 326	4.5	14
74	In pancreatic ductal adenocarcinoma blood concentrations of some organochlorine compounds and coffee intake are independently associated with KRAS mutations. <i>Mutagenesis</i> , 2009 , 24, 513-21	2.8	14
73	Ambient air pollution and incident bladder cancer risk: Updated analysis of the Spanish Bladder Cancer Study. <i>International Journal of Cancer</i> , 2019 , 145, 894-900	7.5	14
72	Genetic variation in the TP53 pathway and bladder cancer risk. a comprehensive analysis. <i>PLoS ONE</i> , 2014 , 9, e89952	3.7	13
71	Public health perspective: from personalized medicine to personal health. <i>Personalized Medicine</i> , 2012 , 9, 115-119	2.2	13
70	Bladder cancer and reproductive factors among women in Spain. <i>Cancer Causes and Control</i> , 2009 , 20, 1907-13	2.8	13
69	Application of multi-SNP approaches Bayesian LASSO and AUC-RF to detect main effects of inflammatory-gene variants associated with bladder cancer risk. <i>PLoS ONE</i> , 2013 , 8, e83745	3.7	12
68	Clinical validity of detecting K-ras mutations for the diagnosis of exocrine pancreatic cancer: a prospective study in a clinically-relevant spectrum of patients. <i>European Journal of Epidemiology</i> , 2011 , 26, 229-36	12.1	12

67	Genetic epidemiology of bladder cancer: scaling up in the identification of low-penetrance genetic markers of bladder cancer risk and progression. <i>Scandinavian Journal of Urology and Nephrology</i> , 2008 , 131-40		12
66	Generalizing molecular results arising from incomplete biological samples: expected bias and unexpected findings. <i>Annals of Epidemiology</i> , 2002 , 12, 7-14	6.4	11
65	Whole genome prediction of bladder cancer risk with the Bayesian LASSO. <i>Genetic Epidemiology</i> , 2014 , 38, 467-76	2.6	10
64	Identification and replication of the interplay of four genetic high-risk variants for urinary bladder cancer. <i>Carcinogenesis</i> , 2017 , 38, 1167-1179	4.6	9
63	Lifetime history of alcohol consumption and K-ras mutations in pancreatic ductal adenocarcinoma. <i>Environmental and Molecular Mutagenesis</i> , 2009 , 50, 421-30	3.2	9
62	Estimating dietary intakes from a brief questionnaire: A simulation study of reliability in a molecular epidemiologic study of pancreatic and biliary diseases. <i>European Journal of Epidemiology</i> , 2006 , 21, 417-26	12.1	9
61	Associations between Genetically Predicted Blood Protein Biomarkers and Pancreatic Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020 , 29, 1501-1508	4	9
60	Concentrations of trace elements and KRAS mutations in pancreatic ductal adenocarcinoma. <i>Environmental and Molecular Mutagenesis</i> , 2019 , 60, 693-703	3.2	8
59	Asthma status is associated with decreased risk of aggressive urothelial bladder cancer. <i>International Journal of Cancer</i> , 2018 , 142, 470-476	7.5	8
58	An epistatic interaction between the PAX8 and STK17B genes in papillary thyroid cancer susceptibility. <i>PLoS ONE</i> , 2013 , 8, e74765	3.7	8
57	Determinants of quality of interview and impact on risk estimates in a case-control study of bladder cancer. <i>American Journal of Epidemiology</i> , 2009 , 170, 237-43	3.8	8
56	The inherited genetic component of sporadic pancreatic adenocarcinoma. <i>Pancreatology</i> , 2009 , 9, 206-14	3.8	8
55	CYP1B1 polymorphisms and k-ras mutations in patients with pancreatic ductal adenocarcinoma. <i>Digestive Diseases and Sciences</i> , 2008 , 53, 1417-21	4	8
54	Diesel exhaust and bladder cancer risk by pathologic stage and grade subtypes. <i>Environment International</i> , 2020 , 135, 105346	12.9	8
53	Predictors of oncological outcomes in T1G3 patients treated with BCG who undergo radical cystectomy. <i>World Journal of Urology</i> , 2018 , 36, 1775-1781	4	8
52	DNA Methylation-Derived Immune Cell Profiles, CpG Markers of Inflammation, and Pancreatic Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020 , 29, 1577-1585	4	7
51	Immunohistochemistry-Based Taxonomical Classification of Bladder Cancer Predicts Response to Neoadjuvant Chemotherapy. <i>Cancers</i> , 2020 , 12,	6.6	7
50	Next-generation sequencing of urologic cancers: next is now. <i>European Urology</i> , 2014 , 66, 4-7	10.2	7

49	Environmental and genomic factors as well as interventions influencing smoking cessation: a systematic review of reviews and a proposed working model. <i>Public Health Genomics</i> , 2013 , 16, 159-73	1.9	7
48	Prediction of non-muscle invasive bladder cancer outcomes assessed by innovative multimarker prognostic models. <i>BMC Cancer</i> , 2016 , 16, 351	4.8	7
47	Inflammatory-Related Genetic Variants in Non-Muscle-Invasive Bladder Cancer Prognosis: A Multimarker Bayesian Assessment. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016 , 25, 1144-50	4	7
46	Bringing Greater Accuracy to Europe's Healthcare Systems: The Unexploited Potential of Biomarker Testing in Oncology. <i>Biomedicine Hub</i> , 2020 , 5, 182-223	1.3	6
45	A dynamic model for the risk of bladder cancer progression. <i>Statistics in Medicine</i> , 2012 , 31, 287-300	2.3	6
44	Relationships of hepatic and pancreatic biomarkers with the cholestatic syndrome and tumor stage in pancreatic cancer. <i>Biomarkers</i> , 2012 , 17, 557-65	2.6	6
43	Reply to "Mosaic loss of chromosome Y in leukocytes matters". <i>Nature Genetics</i> , 2019 , 51, 7-9	36.3	6
42	A multilayered post-GWAS assessment on genetic susceptibility to pancreatic cancer. <i>Genome Medicine</i> , 2021 , 13, 15	14.4	6
41	7q21-rs6964587 and breast cancer risk: an extended case-control study by the Breast Cancer Association Consortium. <i>Journal of Medical Genetics</i> , 2011 , 48, 698-702	5.8	5
40	Bladder cancer and apoptosis: matters of life and death. <i>Lancet Oncology</i> , 2007 , 8, 91-2	21.7	5
39	A combination of urinary biomarker panel and PancRISK score for earlier detection of pancreatic cancer: A case-control study. <i>PLoS Medicine</i> , 2020 , 17, e1003489	11.6	5
38	Advantage of using allele-specific copy numbers when testing for association in regions with common copy number variants. <i>PLoS ONE</i> , 2013 , 8, e75350	3.7	5
37	UEG position paper on pancreatic cancer. Bringing pancreatic cancer to the 21st century: Prevent, detect, and treat the disease earlier and better. <i>United European Gastroenterology Journal</i> , 2021 , 9, 860	5.3	5
36	A faecal microbiota signature with high specificity for pancreatic cancer.. <i>Gut</i> , 2022 ,	19.2	5
35	Genome-Wide Gene-Diabetes and Gene-Obesity Interaction Scan in 8,255 Cases and 11,900 Controls from PanScan and PanC4 Consortia. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020 , 29, 1784-1791	4	4
34	Pancreatic cancer and autoimmune diseases: An association sustained by computational and epidemiological case-control approaches. <i>International Journal of Cancer</i> , 2019 , 144, 1540-1549	7.5	4
33	Diagnostic and Prognostic Performance of Secreted Protein Acidic and Rich in Cysteine (SPARC) Assay for Detecting Primary and Recurrent Urinary Bladder Cancer. <i>Proteomics - Clinical Applications</i> , 2019 , 13, e1800148	3.1	4
32	Bringing Onco-Innovation to Europe's Healthcare Systems: The Potential of Biomarker Testing, Real World Evidence, Tumour Agnostic Therapies to Empower Personalised Medicine. <i>Cancers</i> , 2021 , 13,	6.6	4

31	Plasma protein biomarkers for early detection of pancreatic ductal adenocarcinoma. <i>International Journal of Cancer</i> , 2021 , 148, 2048-2058	7.5	4
30	DoriTool: A Bioinformatics Integrative Tool for Post-Association Functional Annotation. <i>Public Health Genomics</i> , 2017 , 20, 126-135	1.9	3
29	Association of patientsPsex with treatment outcomes after intravesical bacillus Calmette-Guérin immunotherapy for T1G3/HG bladder cancer. <i>World Journal of Urology</i> , 2021 , 39, 3337-3344	4	3
28	CD8+ Cytotoxic Immune Infiltrate in Non-Muscle Invasive Bladder Cancer: A Standardized Methodology to Study Association with Clinico-Pathological Features and Prognosis. <i>Bladder Cancer</i> , 2019 , 5, 159-169	1	2
27	Genomics in Primary and Secondary Prevention of Pancreatic Cancer. <i>Public Health Genomics</i> , 2017 , 20, 92-99	1.9	2
26	Cancer Genomics and Public Health. <i>Public Health Genomics</i> , 2017 , 20, 67-69	1.9	2
25	Bratislava Statement: consensus recommendations for improving pancreatic cancer care. <i>ESMO Open</i> , 2020 , 5, e001051	6	2
24	Mendelian Randomization Analysis of n-6 Polyunsaturated Fatty Acid Levels and Pancreatic Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020 , 29, 2735-2739	4	2
23	Associations between pancreatic expression quantitative traits and risk of pancreatic ductal adenocarcinoma. <i>Carcinogenesis</i> , 2021 , 42, 1037-1045	4.6	2
22	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, 1408-1417	7	2
21	Smoking Modifies Pancreatic Cancer Risk Loci on 2q21.3. <i>Cancer Research</i> , 2021 , 81, 3134-3143	10.1	2
20	Response to: Variation of the age at onset of pancreatic cancer according to tobacco smoking and family history. <i>International Journal of Epidemiology</i> , 2018 , 47, 1358-1359	7.8	2
19	Association of Genetic Variants Affecting microRNAs and Pancreatic Cancer Risk. <i>Frontiers in Genetics</i> , 2021 , 12, 693933	4.5	2
18	Genome-Wide Association Study Data Reveal Genetic Susceptibility to Chronic Inflammatory Intestinal Diseases and Pancreatic Ductal Adenocarcinoma Risk. <i>Cancer Research</i> , 2020 , 80, 4004-4013	10.1	1
17	Integrative eQTL analysis of tumor and host omics data in individuals with bladder cancer. <i>Genetic Epidemiology</i> , 2017 , 41, 567-573	2.6	1
16	EU Pancreas: an integrated European platform for pancreas cancer research--from basic science to clinical and public health interventions for a rare disease. <i>Public Health Genomics</i> , 2013 , 16, 305-12	1.9	1
15	Public health genomics and the challenges for epidemiology. <i>European Journal of Public Health</i> , 2011 , 21, 5-6	2.1	1
14	Perspectives on Data Integration in Human Complex Disease Analysis. <i>Advances in Bioinformatics and Biomedical Engineering Book Series</i> , 2015 , 284-322	0.4	1

13	An integrated multi-omics analysis identifies clinically relevant molecular subtypes of non-muscle-invasive bladder cancer		1
12	Somatic Mutation Profiling in the Liquid Biopsy and Clinical Analysis of Hereditary and Familial Pancreatic Cancer Cases Reveals Negativity and a Longer Overall Survival. <i>Cancers</i> , 2021 , 13,	6.6	1
11	A 584 bp deletion in CTRB2 inhibits chymotrypsin B2 activity and secretion and confers risk of pancreatic cancer. <i>American Journal of Human Genetics</i> , 2021 , 108, 1852-1865	11	1
10	Tumor-Infiltrating B- and T-Cell Repertoire in Pancreatic Cancer Associated With Host and Tumor Features. <i>Frontiers in Immunology</i> , 2021 , 12, 730746	8.4	1
9	Disinfection By-Products in Drinking Water and Bladder Cancer: Evaluation of Risk Modification by Common Genetic Polymorphisms in Two Case-Control Studies.. <i>Environmental Health Perspectives</i> , 2022 , 130, 57006	8.4	1
8	Health Literacy: Read All about It <i>Biomedicine Hub</i> , 2017 , 2, 44-47	1.3	0
7	Genome-wide Meta-analysis Identifies Novel Genes Associated with Recurrence and Progression in Non-muscle-invasive Bladder Cancer. <i>European Urology Oncology</i> , 2021 , 5, 70-70	6.7	0
6	Author's reply to: Air pollution and incident bladder cancer: A risk assessment. <i>International Journal of Cancer</i> , 2019 , 145, 3178	7.5	
5	Editorial comment on: FGFR3 mutations indicate better survival in invasive upper urinary tract and bladder tumours. <i>European Urology</i> , 2009 , 55, 658	10.2	
4	Public health genomics in Spain: the status of a non-existing reality. <i>Public Health Genomics</i> , 2012 , 15, 313-21	1.9	
3	Perspectives on Data Integration in Human Complex Disease Analysis 2019 , 1826-1866		
2	Genetic Testing, Genetic Variation, and Genetic Susceptibility 2018 , 629-649		
1	Risk factors for residual disease at re-TUR in a large cohort of T1G3 patients. <i>Actas Urológicas Españolas (English Edition)</i> , 2021 , 45, 473-478		0.1