

# Sanjay Shete

## List of Publications by Year in descending order

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130  
papers

5,694  
citations

117625

34  
h-index

82547

72  
g-index

134  
all docs

134  
docs citations

134  
times ranked

8775  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide association scan of tag SNPs identifies a susceptibility locus for lung cancer at 15q25.1. <i>Nature Genetics</i> , 2008, 40, 616-622.	21.4	1,189
2	Genome-wide association study identifies five susceptibility loci for glioma. <i>Nature Genetics</i> , 2009, 41, 899-904.	21.4	713
3	A Risk Model for Prediction of Lung Cancer. <i>Journal of the National Cancer Institute</i> , 2007, 99, 715-726.	6.3	362
4	Exploring joint effects of genes and the clinical efficacy of morphine for cancer pain: OPRM1 and COMT gene. <i>Pain</i> , 2007, 130, 25-30.	4.2	269
5	Chromosome 7p11.2 (EGFR) variation influences glioma risk. <i>Human Molecular Genetics</i> , 2011, 20, 2897-2904.	2.9	158
6	Ignoring Linkage Disequilibrium among Tightly Linked Markers Induces False-Positive Evidence of Linkage for Affected Sib Pair Analysis. <i>American Journal of Human Genetics</i> , 2004, 75, 1106-1112.	6.2	137
7	Genetic Variants on 15q25.1, Smoking, and Lung Cancer: An Assessment of Mediation and Interaction. <i>American Journal of Epidemiology</i> , 2012, 175, 1013-1020.	3.4	128
8	Genetic linkage and imprinting effects on body mass index in children and young adults. <i>European Journal of Human Genetics</i> , 2003, 11, 425-432.	2.8	93
9	Adding Further Power to the Haseman and Elston Method for Detecting Linkage in Larger Sibships: Weighting Sums and Differences. <i>Human Heredity</i> , 2003, 55, 79-85.	0.8	91
10	Chemotherapy-Induced Peripheral Neuropathy as a Predictor of Neuropathic Pain in Breast Cancer Patients Previously Treated With Paclitaxel. <i>Journal of Pain</i> , 2009, 10, 1146-1150.	1.4	88
11	Joint Effects of Germ-Line p53 Mutation and Sex on Cancer Risk in Li-Fraumeni Syndrome. <i>Cancer Research</i> , 2006, 66, 8287-8292.	0.9	86
12	GLIOGENE—an International Consortium to Understand Familial Glioma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 1730-1734.	2.5	74
13	A Genome-Wide Association Study Identifies a Locus on Chromosome 14q21 as a Predictor of Leukocyte Telomere Length and as a Marker of Susceptibility for Bladder Cancer. <i>Cancer Prevention Research</i> , 2011, 4, 514-521.	1.5	73
14	Mediating effects of smoking and chronic obstructive pulmonary disease on the relation between the CHRNA5A3 genetic locus and lung cancer risk. <i>Cancer</i> , 2010, 116, 3458-3462.	4.1	67
15	Testing for Genetic Linkage in Families by a Variance-Components Approach in the Presence of Genomic Imprinting. <i>American Journal of Human Genetics</i> , 2002, 70, 751-757.	6.2	61
16	Cytokine Genes and Pain Severity in Lung Cancer: Exploring the Influence of <i>TNF-<math>\alpha</math>-308 G/A</i> <i>IL6-174G/C</i> and <i>IL8-251T/A</i> . <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 2745-2751.	2.5	60
17	Role of Inflammation Gene Polymorphisms on Pain Severity in Lung Cancer Patients. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 2636-2642.	2.5	57
18	Genetic Variations in Interleukin-8 and Interleukin-10 Are Associated With Pain, Depressed Mood, and Fatigue in Lung Cancer Patients. <i>Journal of Pain and Symptom Management</i> , 2013, 46, 161-172.	1.2	57

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19	Effect of Winsorization on Power and Type 1 Error of Variance Components and Related Methods of QTL Detection. <i>Behavior Genetics</i> , 2004, 34, 153-159.	2.1	56
20	Promoter polymorphism (T>C) in the endothelial nitric oxide synthase gene is associated with risk of sporadic breast cancer in non-Hispanic white women age younger than 55 years. <i>Cancer</i> , 2006, 107, 2245-2253.	4.1	56
21	X-Chromosome Genetic Association Test Accounting for X-Inactivation, Skewed X-Inactivation, and Escape from X-Inactivation. <i>Genetic Epidemiology</i> , 2014, 38, 483-493.	1.3	56
22	Smoking Behaviors in Survivors of Smoking-Related and Non-Smoking-Related Cancers. <i>JAMA Network Open</i> , 2020, 3, e209072.	5.9	52
23	Methylenetetrahydrofolate reductase polymorphisms and risk of squamous cell carcinoma of the head and neck: A case-control analysis. <i>International Journal of Cancer</i> , 2005, 115, 131-136.	5.1	50
24	The Influence of Tumor Necrosis Factor- $\alpha$ 308 G/A and IL-6 174 G/C on Pain and Analgesia Response in Lung Cancer Patients Receiving Supportive Care. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 3262-3267.	2.5	50
25	Using the weighted area under the net benefit curve for decision curve analysis. <i>BMC Medical Informatics and Decision Making</i> , 2016, 16, 94.	3.0	49
26	Symptom clusters of pain, depressed mood, and fatigue in lung cancer: assessing the role of cytokine genes. <i>Supportive Care in Cancer</i> , 2013, 21, 3117-3125.	2.2	47
27	Genomic Imprinting and Linkage Test for Quantitative-Trait Loci in Extended Pedigrees. <i>American Journal of Human Genetics</i> , 2003, 73, 933-938.	6.2	45
28	Genome-Wide High-Density SNP Linkage Search for Glioma Susceptibility Loci: Results from the Gliogene Consortium. <i>Cancer Research</i> , 2011, 71, 7568-7575.	0.9	44
29	Gene network analysis shows immune-signaling and ERK1/2 as novel genetic markers for multiple addiction phenotypes: alcohol, smoking and opioid addiction. <i>BMC Systems Biology</i> , 2015, 9, 25.	3.0	43
30	Blood-Based Biomarker Panel for Personalized Lung Cancer Risk Assessment. <i>Journal of Clinical Oncology</i> , 2022, 40, 876-883.	1.6	43
31	Exposure to Smoking Imagery in the Movies and Experimenting with Cigarettes among Mexican Heritage Youth. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 3435-3443.	2.5	40
32	Declining awareness of HPV and HPV vaccine within the general US population. <i>Human Vaccines and Immunotherapeutics</i> , 2021, 17, 420-427.	3.3	40
33	Genetic and Nongenetic Covariates of Pain Severity in Patients with Adenocarcinoma of the Pancreas: Assessing the Influence of Cytokine Genes. <i>Journal of Pain and Symptom Management</i> , 2009, 38, 894-902.	1.2	38
34	Estimation of odds ratios of genetic variants for the secondary phenotypes associated with primary diseases. <i>Genetic Epidemiology</i> , 2011, 35, 190-200.	1.3	37
35	Trends in HPV Vaccination Initiation and Completion Within Ages 9-12 Years: 2008-2018. <i>Pediatrics</i> , 2021, 147, .	2.1	37
36	Complex segregation analysis reveals a multigene model for lung cancer. <i>Human Genetics</i> , 2005, 116, 121-127.	3.8	36

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37	Genetic Epidemiology and Nonsyndromic Structural Birth Defects. <i>JAMA Pediatrics</i> , 2014, 168, 371.	6.2	36
38	A Test for Genetic Association that Incorporates Information about Deviation from Hardy-Weinberg Proportions in Cases. <i>American Journal of Human Genetics</i> , 2008, 83, 53-63.	6.2	35
39	A Genome-Wide Association Study Identifies Two Novel Susceptible Regions for Squamous Cell Carcinoma of the Head and Neck. <i>Cancer Research</i> , 2020, 80, 2451-2460.	0.9	33
40	Improving the Power of Sib Pair Quantitative Trait Loci Detection by Phenotype Winsorization. <i>Human Heredity</i> , 2002, 53, 59-67.	0.8	32
41	Glioma-related seizures in relation to histopathological subtypes: a report from the glioma international case-control study. <i>Journal of Neurology</i> , 2018, 265, 1432-1442.	3.6	32
42	Description of selected characteristics of familial glioma patients – Results from the Gliogene Consortium. <i>European Journal of Cancer</i> , 2013, 49, 1335-1345.	2.8	30
43	Oral microbiome and onset of oral mucositis in patients with squamous cell carcinoma of the head and neck. <i>Cancer</i> , 2020, 126, 5124-5136.	4.1	30
44	Disparities in Secondhand Smoke Exposure in the United States. <i>JAMA Internal Medicine</i> , 2021, 181, 134.	5.1	30
45	Identifying novel genes and biological processes relevant to the development of cancer therapy-induced mucositis: An informative gene network analysis. <i>PLoS ONE</i> , 2017, 12, e0180396.	2.5	27
46	Assessment of Trends in Cigarette Smoking Cessation After Cancer Diagnosis Among US Adults, 2000 to 2017. <i>JAMA Network Open</i> , 2020, 3, e2012164.	5.9	27
47	Genetic determinants of immune-related adverse events in patients with melanoma receiving immune checkpoint inhibitors. <i>Cancer Immunology, Immunotherapy</i> , 2021, 70, 1939-1949.	4.2	27
48	Influence of Subjective Social Status on the Relationship Between Positive Outcome Expectations and Experimentation with Cigarettes. <i>Journal of Adolescent Health</i> , 2009, 44, 342-348.	2.5	26
49	MAPK1/ERK2 as novel target genes for pain in head and neck cancer patients. <i>BMC Genetics</i> , 2016, 17, 40.	2.7	25
50	Method for Evaluating Multiple Mediators: Mediating Effects of Smoking and COPD on the Association between the CHRNA5-A3 Variant and Lung Cancer Risk. <i>PLoS ONE</i> , 2012, 7, e47705.	2.5	23
51	Using Both Cases and Controls for Testing Hardy-Weinberg Proportions in a Genetic Association Study. <i>Human Heredity</i> , 2010, 69, 212-218.	0.8	22
52	Targeted Sequencing in Chromosome 17q Linkage Region Identifies Familial Glioma Candidates in the Gliogene Consortium. <i>Scientific Reports</i> , 2015, 5, 8278.	3.3	22
53	A call for the introduction of gender-neutral HPV vaccination to national immunisation programmes in Africa. <i>The Lancet Global Health</i> , 2019, 7, e20-e21.	6.3	21
54	Comprehensive functional annotation of susceptibility variants identifies genetic heterogeneity between lung adenocarcinoma and squamous cell carcinoma. <i>Frontiers of Medicine</i> , 2021, 15, 275-291.	3.4	21

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55	Mixed-effects Logistic Approach for Association Following Linkage Scan for Complex Disorders. <i>Annals of Human Genetics</i> , 2007, 71, 230-237.	0.8	20
56	A Variable Age of Onset Segregation Model for Linkage Analysis, with Correction for Ascertainment, Applied to Glioma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 2242-2251.	2.5	20
57	Insight in glioma susceptibility through an analysis of 6p22.3, 12p13.33-12.1, 17q22-23.2 and 18q23 SNP genotypes in familial and non-familial glioma. <i>Human Genetics</i> , 2012, 131, 1507-1517.	3.8	20
58	Testing Departure from Hardy-Weinberg Proportions. <i>Methods in Molecular Biology</i> , 2017, 1666, 83-115.	0.9	20
59	Cancer-Related Risk Perceptions and Beliefs in Texas: Findings from a 2018 Population-Level Survey. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 486-494.	2.5	19
60	Genome-wide association study identifies genes associated with neuropathy in patients with head and neck cancer. <i>Scientific Reports</i> , 2018, 8, 8789.	3.3	18
61	Linkage Analysis of Affected Sib Pairs Allowing for Parental Origin Effects. <i>Annals of Human Genetics</i> , 2005, 69, 113-126.	0.8	16
62	Cigarette Experimentation in Mexican Origin Youth: Psychosocial and Genetic Determinants. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 228-238.	2.5	16
63	Exposure to Court-Ordered Tobacco Industry Antismoking Advertisements Among US Adults. <i>JAMA Network Open</i> , 2019, 2, e196935.	5.9	16
64	Safety Concerns or Adverse Effects as the Main Reason for Human Papillomavirus Vaccine Refusal. <i>JAMA Pediatrics</i> , 2021, 175, 1074.	6.2	16
65	Parametric Approach to Genomic Imprinting Analysis with Applications to Angelman's Syndrome. <i>Human Heredity</i> , 2005, 59, 26-33.	0.8	15
66	Differences in Sun Protection Behaviors Between Rural and Urban Communities in Texas. <i>Journal of Rural Health</i> , 2019, 35, 155-166.	2.9	15
67	Determinants of patient-reported xerostomia among long-term oropharyngeal cancer survivors. <i>Cancer</i> , 2021, 127, 4470-4480.	4.1	14
68	Calculation of exact p-values when SNPs are tested using multiple genetic models. <i>BMC Genetics</i> , 2014, 15, 75.	2.7	12
69	Genome-wide association study suggests common variants within RP11-634B7.4 gene influencing severe pre-treatment pain in head and neck cancer patients. <i>Scientific Reports</i> , 2016, 6, 34206.	3.3	12
70	Selection of X-chromosome Inactivation Model. <i>Cancer Informatics</i> , 2017, 16, 117693511774727.	1.9	12
71	Reasons for not receiving the HPV vaccine among eligible adults: Lack of knowledge and of provider recommendations contribute more than safety and insurance concerns. <i>Cancer Medicine</i> , 2020, 9, 5281-5290.	2.8	12
72	Effects of measured susceptibility genes on cancer risk in family studies. <i>Human Genetics</i> , 2010, 127, 349-357.	3.8	11

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73	Testing Hardy-Weinberg Proportions in a Frequency-Matched Case-Control Genetic Association Study. PLoS ONE, 2011, 6, e27642.	2.5	11
74	Power and type I error results for a bias-correction approach recently shown to provide accurate odds ratios of genetic variants for the secondary phenotypes associated with primary diseases. Genetic Epidemiology, 2011, 35, 739-743.	1.3	11
75	Characteristics of US adults attempting tobacco use cessation using e-cigarettes. Addictive Behaviors, 2020, 100, 106123.	3.0	11
76	Sustained reduction of attentional bias to smoking cues by smartphone-delivered attentional bias modification training for smokers.. Psychology of Addictive Behaviors, 2022, 36, 906-919.	2.1	11
77	A Note on the Optimal Measure of Allelic Association. Annals of Human Genetics, 2003, 67, 189-191.	0.8	10
78	Joint linkage and imprinting analyses of GAW15 rheumatoid arthritis and gene expression data. BMC Proceedings, 2007, 1, S53.	1.6	10
79	A Linkage Disequilibrium-Based Approach to Selecting Disease-Associated Rare Variants. PLoS ONE, 2013, 8, e69226.	2.5	10
80	Examining Rural-Urban Differences in Fatalism and Information Overload: Data from 12 NCI-Designated Cancer Centers. Cancer Epidemiology Biomarkers and Prevention, 2022, 31, 393-403.	2.5	10
81	Individual-Specific Liability Groups in Genetic Linkage, with Applications to Kindreds with Li-Fraumeni Syndrome. American Journal of Human Genetics, 2002, 70, 813-817.	6.2	9
82	Imprinting detection by extending a regression-based QTL analysis method. Human Genetics, 2007, 122, 159-174.	3.8	9
83	Analysis of Secondary Phenotype Involving the Interactive Effect of the Secondary Phenotype and Genetic Variants on the Primary Disease. Annals of Human Genetics, 2012, 76, 484-499.	0.8	9
84	Beliefs About HPV Vaccine's Success at Cervical Cancer Prevention Among Adult US Women. JNCI Cancer Spectrum, 2019, 3, pkz064.	2.9	9
85	Association of dual and poly tobacco use with depressive symptoms and use of antidepressants. Addictive Behaviors, 2021, 115, 106790.	3.0	9
86	A Risk Prediction Model for Smoking Experimentation in Mexican American Youth. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 2165-2174.	2.5	8
87	Demographic, psychosocial, and genetic risk associated with smokeless tobacco use among Mexican heritage youth. BMC Medical Genetics, 2015, 16, 43.	2.1	8
88	An approach to estimate bidirectional mediation effects with application to body mass index and fasting glucose. Annals of Human Genetics, 2018, 82, 396-406.	0.8	7
89	Physician-office vs home uptake of colorectal cancer screening using FOBT/FIT among screening-eligible US adults. Cancer Medicine, 2019, 8, 7408-7418.	2.8	7
90	Genome-wide association study of INDELs identified four novel susceptibility loci associated with lung cancer risk. International Journal of Cancer, 2020, 146, 2855-2864.	5.1	7

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91	Trends in the rates of health-care providersâ€™ recommendation for HPV vaccine from 2012 to 2018: a multi-round cross-sectional analysis of the health information national trends survey. <i>Human Vaccines and Immunotherapeutics</i> , 2021, 17, 3081-3089.	3.3	7
92	Genetic variants in <i>CYP2B6</i> and <i>HSD17B12</i> associated with risk of squamous cell carcinoma of the head and neck. <i>International Journal of Cancer</i> , 2022, 151, 553-564.	5.1	7
93	The influence of parentâ€™s child gender on intentions to refuse HPV vaccination due to safety concerns/side effects, National Immunization Survey â€œ Teen, 2010â€œ2019. <i>Human Vaccines and Immunotherapeutics</i> , 2022, 18, .	3.3	7
94	Variance Components Analysis for Genetic Linkage of Time to Onset for Disease. <i>Genetic Epidemiology</i> , 2001, 21, S768-73.	1.3	6
95	A Novel Approach to Exploring Potential Interactions among Single-Nucleotide Polymorphisms of Inflammation Genes in Gliomagenesis: An Exploratory Case-Only Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 1683-1689.	2.5	6
96	Gaussian graphical models for phenotypes using pedigree data and exploratory analysis using networks with genetic and nongenetic factors based on Genetic Analysis Workshop 18 data. <i>BMC Proceedings</i> , 2014, 8, S99.	1.6	6
97	Estimation of indirect effect when the mediator is a censored variable. <i>Statistical Methods in Medical Research</i> , 2018, 27, 3010-3025.	1.5	6
98	Chronic obstructive pulmonary disease among lung cancer-free smokers: The importance of healthy controls. <i>Respiratory Investigation</i> , 2018, 56, 28-33.	1.8	6
99	Association Analysis of Driver Geneâ€™s Related Genetic Variants Identified Novel Lung Cancer Susceptibility Loci with 20,871 Lung Cancer Cases and 15,971 Controls. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1423-1429.	2.5	6
100	Awareness of Heated Tobacco Products among US Adults â€œ Health Information National Trends Survey, 2020. <i>Substance Abuse</i> , 2022, 43, 1023-1034.	2.3	6
101	Uniformly Minimum Variance Unbiased Estimation of Gene Diversity. , 2003, 94, 421-424.		5
102	Genetic imprinting analysis for alcoholism genes using variance components approach. <i>BMC Genetics</i> , 2005, 6, S161.	2.7	5
103	A Novel Approach to Detect Parent-of-Origin Effects from Pedigree Data with Application to Beckwith-Wiedemann Syndrome. <i>Annals of Human Genetics</i> , 2007, 71, 804-814.	0.8	5
104	Empirical estimation of sequencing error rates using smoothing splines. <i>BMC Bioinformatics</i> , 2016, 17, 177.	2.6	5
105	Prevalence and determinants of cervical cancer screening with a combination of cytology and human papillomavirus testing. <i>Annals of Epidemiology</i> , 2019, 36, 40-47.	1.9	5
106	Mediation analysis in a caseâ€“control study when the mediator is a censored variable. <i>Statistics in Medicine</i> , 2019, 38, 1213-1229.	1.6	5
107	Association of Exposure to Court-Ordered Tobacco Industry Antismoking Advertisements With Intentions and Attempts to Quit Smoking Among US Adults. <i>JAMA Network Open</i> , 2020, 3, e209504.	5.9	5
108	Examining lung cancer screening utilization with public-use data: Opportunities and challenges. <i>Preventive Medicine</i> , 2021, 147, 106503.	3.4	5

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109	Association of Risk Factors With Patient-Reported Voice and Speech Symptoms Among Long-term Survivors of Oropharyngeal Cancer. <i>JAMA Otolaryngology - Head and Neck Surgery</i> , 2021, 147, 615.	2.2	5
110	Are beliefs about the importance of genetics for cancer prevention and early detection associated with high risk cancer genetic testing in the U.S. Population?. <i>Preventive Medicine Reports</i> , 2022, 27, 101781.	1.8	5
111	Response to Zang et Al. and Han et Al.. <i>American Journal of Human Genetics</i> , 2009, 84, 298-300.	6.2	4
112	Exact Statistical Tests for Heterogeneity of Frequencies Based on Extreme Values. <i>Communications in Statistics Part B: Simulation and Computation</i> , 2010, 39, 612-623.	1.2	4
113	Time-varying SMART design and data analysis methods for evaluating adaptive intervention effects. <i>BMC Medical Research Methodology</i> , 2016, 16, 112.	3.1	4
114	A Robust and Powerful Set-Valued Approach to Rare Variant Association Analyses of Secondary Traits in Case-Control Sequencing Studies. <i>Genetics</i> , 2017, 205, 1049-1062.	2.9	4
115	Processing and Analyzing Human Microbiome Data. <i>Methods in Molecular Biology</i> , 2017, 1666, 649-677.	0.9	4
116	TLINKAGE-IMPRINT: A Model-Based Approach to Performing Two-Locus Genetic Imprinting Analysis. <i>Human Heredity</i> , 2006, 62, 145-156.	0.8	3
117	Prevalence of abnormal cervical cancer screening outcomes among screening-compliant women in the United States. <i>American Journal of Obstetrics and Gynecology</i> , 2019, 221, 75-77.	1.3	3
118	An Approach to Analyze Longitudinal Zero-Inflated Microbiome Count Data Using Two-Stage Mixed Effects Models. <i>Statistics in Biosciences</i> , 2021, 13, 267-290.	1.2	3
119	Factors Influencing Discussion of Cancer Genetic Testing with Health-Care Providers in a Population-Based Survey. <i>Public Health Genomics</i> , 2021, 24, 160-170.	1.0	3
120	Association of hearing loss and tinnitus symptoms with health-related quality of life among long-term oropharyngeal cancer survivors. <i>Cancer Medicine</i> , 0, , .	2.8	3
121	Adaptations of Linkage and Association Methods for the Study of Asthma, A Complex Trait. <i>Genetic Epidemiology</i> , 2001, 21, S89-96.	1.3	2
122	Comparison of multilevel modeling and the family-based association test for identifying genetic variants associated with systolic and diastolic blood pressure using Genetic Analysis Workshop 18 simulated data. <i>BMC Proceedings</i> , 2014, 8, S30.	1.6	2
123	Evaluating Methods for Modeling Epistasis Networks with Application to Head and Neck Cancer. <i>Cancer Informatics</i> , 2015, 14s2, CIN.S17289.	1.9	2
124	Mediation model with a categorical exposure and a censored mediator with application to a genetic study. <i>PLoS ONE</i> , 2021, 16, e0257628.	2.5	2
125	Risk factors associated with patient-reported fatigue among long-term oropharyngeal carcinoma survivors. <i>Head and Neck</i> , 2022, 44, 952-963.	2.0	2
126	Genetic susceptibility to patient-reported xerostomia among long-term oropharyngeal cancer survivors. <i>Scientific Reports</i> , 2022, 12, 6662.	3.3	2



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127	Data mining of RNA expression and DNA genotype data: Presentation Group 5 contributions to Genetic Analysis Workshop 15. <i>Genetic Epidemiology</i> , 2007, 31, S43-S50.	1.3	1
128	Prevalence of abnormal cervical cancer screening outcomes among women in the United States: results from the National Health Interview Survey, 2018. <i>American Journal of Obstetrics and Gynecology</i> , 2020, 223, 938-941.	1.3	1
129	Industry-sponsored antismoking advertisements in low-income countries. <i>The Lancet Global Health</i> , 2020, 8, e485-e486.	6.3	1
130	Gender-neutral HPV vaccination in Africa – Authors' reply. <i>The Lancet Global Health</i> , 2019, 7, e564.	6.3	0