M Dawn Teare

List of Publications by Year
in descending order

78	4,633	159585 30	106344 65
papers	citations	h-index	g-index

83	83	83	8423
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Large-scale association analysis identifies new lung cancer susceptibility loci and heterogeneity in genetic susceptibility across histological subtypes. Nature Genetics, 2017, 49, 1126-1132.	21.4	472
2	Sample size requirements to estimate key design parameters from external pilot randomised controlled trials: a simulation study. Trials, 2014, 15, 264.	1.6	407
3	Localization to Xq27 of a susceptibility gene for testicular germ-cell tumours. Nature Genetics, 2000, 24, 197-200.	21.4	260
4	Mutations in the Gene Encoding Capillary Morphogenesis Protein 2 Cause Juvenile Hyaline Fibromatosis and Infantile Systemic Hyalinosis. American Journal of Human Genetics, 2003, 73, 791-800.	6.2	209
5	Genetic linkage studies. Lancet, The, 2005, 366, 1036-1044.	13.7	195
6	The Extent of Linkage Disequilibrium in Four Populations with Distinct Demographic Histories. American Journal of Human Genetics, 2000, 67, 1544-1554.	6.2	192
7	Association of a Common Variant of the CASP8 Gene With Reduced Risk of Breast Cancer. Journal of the National Cancer Institute, 2004, 96, 1866-1869.	6.3	188
8	Contemporary Occupational Carcinogen Exposure and Bladder Cancer. JAMA Oncology, 2015, 1, 1282.	7.1	184
9	Replication of Lung Cancer Susceptibility Loci at Chromosomes 15q25, 5p15, and 6p21: A Pooled Analysis From the International Lung Cancer Consortium. Journal of the National Cancer Institute, 2010, 102, 959-971.	6.3	174
10	A common variant in BRCA2 is associated with both breast cancer risk and prenatal viability. Nature Genetics, 2000, 26, 362-364.	21.4	152
11	Increased risk of lung cancer in individuals with a family history of the disease: A pooled analysis from the International Lung Cancer Consortium. European Journal of Cancer, 2012, 48, 1957-1968.	2.8	143
12	Optimizing the yield and utility of circulating cell-free DNA from plasma and serum. Clinica Chimica Acta, 2009, 404, 100-104.	1.1	136
13	Cannabis smoking and lung cancer risk: Pooled analysis in the <scp>I</scp>nternational <scp>L</scp>ung <scp>C</scp>ancer <scp>C</scp>onsortium. International Journal of Cancer, 2015, 136, 894-903.	5.1	131
14	Atrial fibrillation associated with ivabradine treatment: meta-analysis of randomised controlled trials. Heart, 2014, 100, 1506-1510.	2.9	126
15	Shared heritability and functional enrichment across six solid cancers. Nature Communications, 2019, 10, 431.	12.8	88
16	Body Mass Index (BMI), BMI Change, and Overall Survival in Patients With SCLC and NSCLC: A Pooled Analysis of the International Lung Cancer Consortium. Journal of Thoracic Oncology, 2019, 14, 1594-1607.	1.1	81
17	Obesity, metabolic factors and risk of different histological types of lung cancer: A Mendelian randomization study. PLoS ONE, 2017, 12, e0177875.	2.5	79
18	Cancer in the families of children with soft tissue sarcoma. Cancer, 1990, 66, 2239-2248.	4.1	76

#	Article	IF	Citations
19	Evaluation of linkage of breast cancer to the putative BRCA3 locus on chromosome 13q21 in 128 multiple case families from the Breast Cancer Linkage Consortium. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 827-831.	7.1	73
20	CHRNA5 Risk Variant Predicts Delayed Smoking Cessation and Earlier Lung Cancer Diagnosisâ€"A Meta-Analysis. Journal of the National Cancer Institute, 2015, 107, .	6.3	72
21	The Gene for Juvenile Hyaline Fibromatosis Maps to Chromosome 4q21. American Journal of Human Genetics, 2002, 71, 975-980.	6.2	71
22	Strategies to increase influenza vaccination rates: outcomes of a nationwide cross-sectional survey of UK general practice. BMJ Open, 2012, 2, e000851.	1.9	69
23	Localization of the Gene for Distal Hereditary Motor Neuronopathy VII (dHMN-VII) to Chromosome 2q14. American Journal of Human Genetics, 2001, 68, 1270-1276.	6.2	68
24	Risk Prediction Models for Lung Cancer: AÂSystematic Review. Clinical Lung Cancer, 2016, 17, 95-106.	2.6	64
25	Identification of susceptibility pathways for the role of chromosome 15q25.1 in modifying lung cancer risk. Nature Communications, 2018, 9, 3221.	12.8	60
26	Ehlers-Danlos Syndrome with Severe Early-Onset Periodontal Disease (EDS-VIII) Is a Distinct, Heterogeneous Disorder with One Predisposition Gene at Chromosome 12p13. American Journal of Human Genetics, 2003, 73, 198-204.	6.2	51
27	The inter-regional epidemiological study of childhood cancer (IRESCC): Case-control study of children with central nervous system tumours. British Journal of Neurosurgery, 1990, 4, 17-25.	0.8	44
28	Fine mapping of MHC region in lung cancer highlights independent susceptibility loci by ethnicity. Nature Communications, 2018, 9, 3927.	12.8	43
29	Genetic Risk Can Be Decreased: Quitting Smoking Decreases and Delays Lung Cancer for Smokers With High and Low CHRNA5 Risk Genotypes â€" A Meta-Analysis. EBioMedicine, 2016, 11, 219-226.	6.1	40
30	Absence of evidence for a familial breast cancer susceptibility gene at chromosome 8p12-p22. Oncogene, 2000, 19, 4170-4173.	5.9	35
31	Appropriate statistical methods for analysing partially nested randomised controlled trials with continuous outcomes: a simulation study. BMC Medical Research Methodology, 2018, 18, 105.	3.1	32
32	Mendelian Randomization and mediation analysis of leukocyte telomere length and risk of lung and head and neck cancers. International Journal of Epidemiology, 2019, 48, 751-766.	1.9	32
33	Genome-wide interaction study of smoking behavior and non-small cell lung cancer risk in Caucasian population. Carcinogenesis, 2018, 39, 336-346.	2.8	29
34	AHEAD Study: an observational study of the management of anticoagulated patients who suffer head injury. BMJ Open, 2017, 7, e014324.	1.9	28
35	Genetic modifiers of radon-induced lung cancer risk: a genome-wide interaction study in former uranium miners. International Archives of Occupational and Environmental Health, 2018, 91, 937-950.	2.3	27
36	Lung Cancer Risk in Never-Smokers of European Descent is Associated With Genetic Variation in the 5p15.33 TERT-CLPTM1Ll Region. Journal of Thoracic Oncology, 2019, 14, 1360-1369.	1.1	27

#	Article	IF	Citations
37	Confirmation of a gene locus for medullary cystic kidney disease (MCKD2) on chromosome 16p12. Kidney International, 2001, 60, 1233-1239.	5.2	25
38	Genetic interaction analysis among oncogenesis-related genes revealed novel genes and networks in lung cancer development. Oncotarget, 2019, 10, 1760-1774.	1.8	25
39	RIPOSTE: a framework for improving the design and analysis of laboratory-based research. ELife, 2015, 4, .	6.0	24
40	Immune-mediated genetic pathways resulting in pulmonary function impairment increase lung cancer susceptibility. Nature Communications, 2020, 11, 27.	12.8	23
41	The relationship between body-mass index and overall survival in non-small cell lung cancer by sex, smoking status, and race: A pooled analysis of 20,937 International lung Cancer consortium (ILCCO) patients. Lung Cancer, 2021, 152, 58-65.	2.0	22
42	Cytokine Gene Polymorphisms in Heavy Drinkers With and Without Decompensated Liver Disease: A Case-Control Study. American Journal of Gastroenterology, 2008, 103, 3039-3046.	0.4	20
43	Allele dose association of theC5orf30rs26232 variant with joint damage in rheumatoid arthritis. Arthritis and Rheumatism, 2013, 65, n/a-n/a.	6.7	20
44	Genome-wide association meta-analysis identifies pleiotropic risk loci for aerodigestive squamous cell cancers. PLoS Genetics, 2021, 17, e1009254.	3.5	19
45	Meta-analysis of five genome-wide linkage studies for body mass index reveals significant evidence for linkage to chromosome 8p. International Journal of Obesity, 2005, 29, 413-419.	3.4	17
46	Linkage analysis and the study of Mendelian disease in the era of whole exome and genome sequencing. Briefings in Functional Genomics, 2014, 13, 378-383.	2.7	17
47	Comparison of Nottingham Prognostic Index and Adjuvant Online prognostic tools in young women with breast cancer: review of a single-institution experience. BMJ Open, 2015, 5, e005576-e005576.	1.9	17
48	A global countryâ€level analysis of the relationship between obesity and <scp>COVID</scp>â€19 cases and mortality. Diabetes, Obesity and Metabolism, 2021, 23, 2697-2706.	4.4	17
49	Differential Risk of ST-Segment Elevation Myocardial Infarction in Male and FemaleÂSmokers. Journal of the American College of Cardiology, 2019, 73, 3259-3266.	2.8	16
50	Metaanalysis of the Association of Smoking andPTPN22R620W Genotype on Autoantibody Status and Radiological Erosions in Rheumatoid Arthritis. Journal of Rheumatology, 2013, 40, 1048-1053.	2.0	15
51	Should all anticoagulated patients with head injury receive a CT scan? Decision-analysis modelling of an observational cohort. BMJ Open, 2016, 6, e013742.	1.9	15
52	Pronounced increase in risk of acute ST-segment elevation myocardial infarction in younger smokers. Heart, 2017, 103, 586-591.	2.9	15
53	Clinical and cost-effectiveness of one-session treatment (OST) versus multisession cognitiveâ€“behavioural therapy (CBT) for specific phobias in children: protocol for a non-inferiority randomised controlled trial. BMJ Open, 2018, 8, e025031.	1.9	15
54	Transparent reporting of research results in eLife. ELife, 2016, 5, .	6.0	12

#	Article	IF	Citations
55	Pleiotropy of genetic variants on obesity and smoking phenotypes: Results from the Oncoarray Project of The International Lung Cancer Consortium. PLoS ONE, 2017, 12, e0185660.	2.5	11
56	Unit of analysis issues in laboratory-based research. ELife, 2018, 7, .	6.0	11
57	Statistical design and analysis in trials of proportionate interventions: a systematic review. Trials, 2019, 20, 151.	1.6	10
58	Unbiased Detection of Somatic Copy Number Aberrations in cfDNA of Lung Cancer Cases and High-Risk Controls with Low Coverage Whole Genome Sequencing. Advances in Experimental Medicine and Biology, 2016, 924, 29-32.	1.6	9
59	Genome-Wide Analysis of Circulating Cell-Free DNA Copy Number Detects Active Melanoma and Predicts Survival. Clinical Chemistry, 2018, 64, 1338-1346.	3.2	9
60	The Recognition of STEMI by Paramedics and the Effect of Computer inTerpretation (RESPECT): a randomised crossover feasibility study. Emergency Medicine Journal, 2016, 33, 471-476.	1.0	8
61	A randomized controlled trial of a proportionate universal parenting program delivery model (E-SEE) Tj ETQq1 1 0.784314 rgBT /Overl	2.5	8
62	A Candidate Gene Association Study of Bone Mineral Density in an Iranian Population. Frontiers in Endocrinology, 2016, 7, 141.	3.5	7
63	Integration of multiomic annotation data to prioritize and characterize inflammation and immuneâ€related risk variants in squamous cell lung cancer. Genetic Epidemiology, 2021, 45, 99-114.	1.3	7
64	Linkage Analysis. Methods in Molecular Biology, 2011, 760, 19-33.	0.9	5
65	The Potential of Adaptive Design in Animal Studies. International Journal of Molecular Sciences, 2015, 16, 24048-24058.	4.1	5
66	Systematic analyses of regulatory variants in DNase I hypersensitive sites identified two novel lung cancer susceptibility loci. Carcinogenesis, 2019, 40, 432-440.	2.8	5
67	Enhancing Social-Emotional Outcomes in Early Years (E-SEE): Randomized Pilot Study of Incredible Years Infant and Toddler Programs. Journal of Child and Family Studies, 2021, 30, 1933-1949.	1.3	5
68	Comparing Methods for Mapping cis Acting Polymorphisms Using Allelic Expression Ratios. PLoS ONE, 2011, 6, e28636.	2.5	5
69	Cost-utility analysis of LEGO based therapy for school children and young people with autism spectrum disorder: results from a randomised controlled trial. BMJ Open, 2022, 12, e056347.	1.9	5
70	Repeat coronary angiography with previously normal arteries: A futile exercise?. Catheterization and Cardiovascular Interventions, 2015, 85, 401-405.	1.7	4
71	Multinational Survey of Treatment Practices of Clinicians Managing Subclinical Hypothyroidism in Older People in 2019. European Thyroid Journal, 2021, 10, 330-338.	2.4	4
72	BIOlogical Factors that Limit sustAined Remission in rhEumatoid arthritis (the BIO-FLARE study): protocol for a non-randomised longitudinal cohort study. BMC Rheumatology, 2021, 5, 22.	1.6	4

#	Article	IF	Citations
73	Stakeholder Perspectives on Clinical Decision Support Tools to Inform Clinical Artificial Intelligence Implementation: Protocol for a Framework Synthesis for Qualitative Evidence. JMIR Research Protocols, 2022, 11, e33145.	1.0	4
74	Iam hiQâ€”a novel pair of accuracy indices for imputed genotypes. BMC Bioinformatics, 2022, 23, 50.	2.6	2
75	A proportionate, universal parenting programme to enhance social-emotional well-being in infants and toddlers in England: the E-SEE Steps RCT. Public Health Research, 2022, 10, 1-162.	1.3	2
76	Circulating cell-free DNA: a potential biomarker in lung cancer. Lung Cancer Management, 2013, 2, 407-422.	1.5	0
77	A reply to â€œLung cancer outcomes: Are BMI and race clinically relevant?â€• Lung Cancer, 2021, 154, 225-226.	2.0	0
78	Methodology over metrics: response to Van Calster et al. Journal of Clinical Epidemiology, 2021, , .	5.0	0