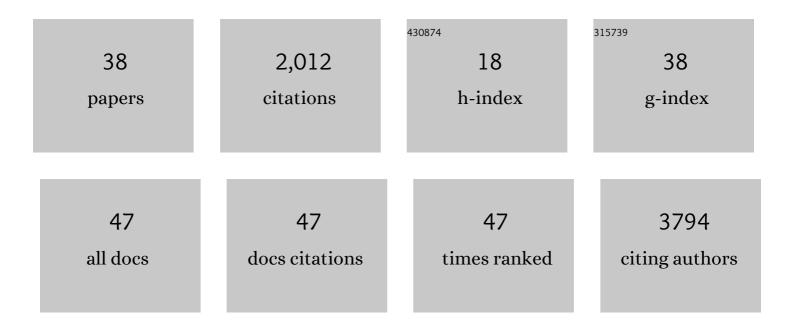
Julia Steinberg

List of Publications by Year in descending order

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IIIIIA STEINBERC

#	Article	IF	CITATIONS
1	A molecular map of long non-coding RNA expression, isoform switching and alternative splicing in osteoarthritis. Human Molecular Genetics, 2022, 31, 2090-2105.	2.9	15
2	Independent evaluation of melanoma polygenic risk scores in <scp>UK</scp> and Australian prospective cohorts*. British Journal of Dermatology, 2022, 186, 823-834.	1.5	10
3	Accurate categorisation of menopausal status for research studies: a step-by-step guide and detailed algorithm considering age, self-reported menopause and factors potentially masking the occurrence of menopause. BMC Research Notes, 2022, 15, 88.	1.4	5
4	Using multivariable Mendelian randomization to estimate the causal effect of bone mineral density on osteoarthritis risk, independently of body mass index. International Journal of Epidemiology, 2022, 51, 1254-1267.	1.9	20
5	Lynch syndrome testing of colorectal cancer patients in a high-income country with universal healthcare: a retrospective study of current practice and gaps in seven australian hospitals. Hereditary Cancer in Clinical Practice, 2022, 20, 18.	1.5	2
6	Evaluating risk factors for lung cancer among never-smoking individuals using two Australian studies. Journal of Cancer Research and Clinical Oncology, 2022, 148, 2827-2840.	2.5	3
7	Projections of smoking-related cancer mortality in Australia to 2044. Journal of Epidemiology and Community Health, 2022, 76, 792-799.	3.7	0
8	An epigenome-wide view of osteoarthritis in primary tissues. American Journal of Human Genetics, 2022, 109, 1255-1271.	6.2	13
9	Large-scale systematic analysis of exposure to multiple cancer risk factors and the associations between exposure patterns and cancer incidence. Scientific Reports, 2021, 11, 2343.	3.3	2
10	How Well Have Projected Lung Cancer Rates Predicted the Actual Observed Rates?. Asian Pacific Journal of Cancer Prevention, 2021, 22, 437-445.	1.2	1
11	A molecular quantitative trait locus map for osteoarthritis. Nature Communications, 2021, 12, 1309.	12.8	53
12	Systematic review and meta-analysis of residential radon and lung cancer in never-smokers. European Respiratory Review, 2021, 30, 200230.	7.1	36
13	Linking chondrocyte and synovial transcriptional profile to clinical phenotype in osteoarthritis. Annals of the Rheumatic Diseases, 2021, 80, 1070-1074.	0.9	25
14	Deciphering osteoarthritis genetics across 826,690 individuals from 9 populations. Cell, 2021, 184, 4784-4818.e17.	28.9	188
15	Accelerating functional gene discovery in osteoarthritis. Nature Communications, 2021, 12, 467.	12.8	33
16	Health system costs and days in hospital for colorectal cancer patients in New South Wales, Australia. PLoS ONE, 2021, 16, e0260088.	2.5	5
17	Lung cancer risk in never-smokers: An overview of environmental and genetic factors. Chinese Journal of Cancer Research: Official Journal of China Anti-Cancer Association, Beijing Institute for Cancer Research, 2021, 33, 548-562.	2.2	13
18	Changes in cancer incidence and mortality in Australia over the period 1996–2015. BMC Research Notes, 2020, 13, 561.	1.4	6

JULIA STEINBERG

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19	ldentification of new therapeutic targets for osteoarthritis through genome-wide analyses of UK Biobank data. Nature Genetics, 2019, 51, 230-236.	21.4	331
20	Towards global elimination of cervical cancer in all groups of women – Authors' reply. Lancet Oncology, The, 2019, 20, e239.	10.7	0
21	Lung cancer mortality in Australia in the twenty-first century: How many lives can be saved with effective tobacco control?. Lung Cancer, 2019, 130, 208-215.	2.0	16
22	Impact of scaled up human papillomavirus vaccination and cervical screening and the potential for global elimination of cervical cancer in 181 countries, 2020–99: a modelling study. Lancet Oncology, The, 2019, 20, 394-407.	10.7	279
23	A novel variant in <i>GLIS3</i> is associated with osteoarthritis. Annals of the Rheumatic Diseases, 2018, 77, 620-623.	0.9	27
24	Genome-wide analyses using UK Biobank data provide insights into the genetic architecture of osteoarthritis. Nature Genetics, 2018, 50, 549-558.	21.4	223
25	Investigation of common, low-frequency and rare genome-wide variation in anorexia nervosa. Molecular Psychiatry, 2018, 23, 1169-1180.	7.9	32
26	Widespread epigenomic, transcriptomic and proteomic differences between hip osteophytic and articular chondrocytes in osteoarthritis. Rheumatology, 2018, 57, 1481-1489.	1.9	19
27	Effects of chronic cobalt and chromium exposure after metalâ€onâ€metal hip resurfacing: An epigenomeâ€wide association pilot study. Journal of Orthopaedic Research, 2017, 35, 2323-2328.	2.3	11
28	Whole-Genome Sequencing Coupled to Imputation Discovers Genetic Signals for Anthropometric Traits. American Journal of Human Genetics, 2017, 100, 865-884.	6.2	131
29	Integrative epigenomics, transcriptomics and proteomics of patient chondrocytes reveal genes and pathways involved in osteoarthritis. Scientific Reports, 2017, 7, 8935.	3.3	90
30	Evaluation of shared genetic aetiology between osteoarthritis and bone mineral density identifies SMAD3 as a novel osteoarthritis risk locus. Human Molecular Genetics, 2017, 26, 3850-3858.	2.9	56
31	Functional genomics in osteoarthritis: Past, present, and future. Journal of Orthopaedic Research, 2016, 34, 1105-1110.	2.3	28
32	Duplications in ADHD patients harbour neurobehavioural genes that are coâ€expressed with genes associated with hyperactivity in the mouse. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 97-107.	1.7	2
33	Haploinsufficiency predictions without study bias. Nucleic Acids Research, 2015, 43, e101-e101.	14.5	54
34	GeneNet Toolbox for MATLAB: a flexible platform for the analysis of gene connectivity in biological networks. Bioinformatics, 2015, 31, 442-444.	4.1	14
35	Gene Networks Underlying Convergent and Pleiotropic Phenotypes in a Large and Systematically-Phenotyped Cohort with Heterogeneous Developmental Disorders. PLoS Genetics, 2015, 11, e1005012.	3.5	14
36	Diagnostically relevant facial gestalt information from ordinary photos. ELife, 2014, 3, e02020.	6.0	129

#	Article	IF	CITATIONS
37	Gene Age Predicts the Strength of Purifying Selection Acting on Gene Expression Variation in Humans. American Journal of Human Genetics, 2014, 95, 660-674.	6.2	35
38	The Roles of FMRP-Regulated Genes in Autism Spectrum Disorder: Single- and Multiple-Hit Genetic Etiologies. American Journal of Human Genetics, 2013, 93, 825-839.	6.2	60