Amit D Joshi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1236031/publications.pdf

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56	5,376	25 h-index	54
papers	citations		g-index
59	59	59	11281 citing authors
all docs	docs citations	times ranked	

#	Article	IF	CITATIONS
1	Frequency of Bowel Movements and Risk of Diverticulitis. Clinical Gastroenterology and Hepatology, 2022, 20, 325-333.e5.	4.4	7
2	Obesity, Adiposity, and Risk of Symptomatic Gallstone Disease According to Genetic Susceptibility. Clinical Gastroenterology and Hepatology, 2022, 20, e1083-e1120.	4.4	5
3	Cardiovascular disease related circulating biomarkers and cancer incidence and mortality: is there an association?. Cardiovascular Research, 2022, 118, 2317-2328.	3.8	15
4	Gall Bladder Disease and the Risk of Small Bowel Cancerâ€"Results from a Nationwide Swedish Cohort Study. Cancers, 2022, 14, 469.	3.7	0
5	Adherence to 2020 to 2025 Dietary Guidelines for Americans and the Risk of New-Onset Female Gout. JAMA Internal Medicine, 2022, 182, 254.	5.1	21
6	Impact of adiposity on risk of female gout among those genetically predisposed: sex-specific prospective cohort study findings over >32 years. Annals of the Rheumatic Diseases, 2022, 81, 556-563.	0.9	14
7	Beyond GWAS of Colorectal Cancer: Evidence of Interaction with Alcohol Consumption and Putative Causal Variant for the 10q24.2 Region. Cancer Epidemiology Biomarkers and Prevention, 2022, 31, 1077-1089.	2.5	6
8	OUP accepted manuscript. Journal of the National Cancer Institute, 2022, , .	6.3	0
9	Genetic architectures of proximal and distal colorectal cancer are partly distinct. Gut, 2021, 70, 1325-1334.	12.1	44
10	Symptom clusters in COVID-19: A potential clinical prediction tool from the COVID Symptom Study app. Science Advances, 2021, 7, .	10.3	115
11	Response to Li and Hopper. American Journal of Human Genetics, 2021, 108, 527-529.	6.2	5
12	Response to Li and Hopper. American Journal of Human Genetics, 2021, 108, 527-529. Attributes and predictors of long COVID. Nature Medicine, 2021, 27, 626-631.	6.2 30.7	1,613
12	Attributes and predictors of long COVID. Nature Medicine, 2021, 27, 626-631. Genetically Predicted Circulating C-Reactive Protein Concentration and Colorectal Cancer Survival: A Mendelian Randomization Consortium Study. Cancer Epidemiology Biomarkers and Prevention, 2021,	30.7	1,613
12	Attributes and predictors of long COVID. Nature Medicine, 2021, 27, 626-631. Genetically Predicted Circulating C-Reactive Protein Concentration and Colorectal Cancer Survival: A Mendelian Randomization Consortium Study. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 1349-1358.	30.7 2.5	1,613 6
12 13 14	Attributes and predictors of long COVID. Nature Medicine, 2021, 27, 626-631. Genetically Predicted Circulating C-Reactive Protein Concentration and Colorectal Cancer Survival: A Mendelian Randomization Consortium Study. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 1349-1358. Genomic Risk Prediction for Breast Cancer in Older Women. Cancers, 2021, 13, 3533. Genetic Obesity Variants and Risk of Conventional Adenomas and Serrated Polyps. Digestive Diseases	30.7 2.5 3.7	1,613 6 6
12 13 14 15	Attributes and predictors of long COVID. Nature Medicine, 2021, 27, 626-631. Genetically Predicted Circulating C-Reactive Protein Concentration and Colorectal Cancer Survival: A Mendelian Randomization Consortium Study. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 1349-1358. Genomic Risk Prediction for Breast Cancer in Older Women. Cancers, 2021, 13, 3533. Genetic Obesity Variants and Risk of Conventional Adenomas and Serrated Polyps. Digestive Diseases and Sciences, 2021, 1.	30.7 2.5 3.7 2.3	1,613 6 6

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19	Cumulative Burden of Colorectal Cancer–Associated Genetic Variants Is More Strongly Associated With Early-Onset vs Late-Onset Cancer. Gastroenterology, 2020, 158, 1274-1286.e12.	1.3	110
20	Genetic Variant Associated With Survival of Patients With Stage II-III Colon Cancer. Clinical Gastroenterology and Hepatology, 2020, 18, 2717-2723.e3.	4.4	7
21	Exploratory Genome-Wide Interaction Analysis of Nonsteroidal Anti-inflammatory Drugs and Predicted Gene Expression on Colorectal Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 1800-1808.	2.5	1
22	Genome-wide Modeling of Polygenic Risk Score in Colorectal Cancer Risk. American Journal of Human Genetics, 2020, 107, 432-444.	6.2	124
23	Circulating bilirubin levels and risk of colorectal cancer: serological and Mendelian randomization analyses. BMC Medicine, 2020, 18, 229.	5.5	28
24	Rapid implementation of mobile technology for real-time epidemiology of COVID-19. Science, 2020, 368, 1362-1367.	12.6	313
25	Functional informed genomeâ€wide interaction analysis of body mass index, diabetes and colorectal cancer risk. Cancer Medicine, 2020, 9, 3563-3573.	2.8	7
26	Telomere Maintenance Variants and Survival after Colorectal Cancer: Smoking- and Sex-Specific Associations. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 1817-1824.	2.5	5
27	Mortality Risk in Irritable Bowel Syndrome: Results From a Nationwide Prospective Cohort Study. American Journal of Gastroenterology, 2020, 115, 746-755.	0.4	17
28	Racial Differences in Epigenetic Aging of the Colon: Implications for Colorectal Cancer. Journal of the National Cancer Institute, 2020, , .	6.3	4
29	Genetic Variants in the Regulatory T cell–Related Pathway and Colorectal Cancer Prognosis. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 2719-2728.	2.5	1
30	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. Journal of the National Cancer Institute, 2019, 111, 146-157.	6.3	129
31	Metabolomics Analytics Workflow for Epidemiological Research: Perspectives from the Consortium of Metabolomics Studies (COMETS). Metabolites, 2019, 9, 145.	2.9	30
32	Shared heritability and functional enrichment across six solid cancers. Nature Communications, 2019, 10, 431.	12.8	88
33	Effect of video monitor size on polyp detection: a prospective, randomized, controlled trial. Gastrointestinal Endoscopy, 2019, 90, 254-258.e2.	1.0	3
34	Discovery of common and rare genetic risk variants for colorectal cancer. Nature Genetics, 2019, 51, 76-87.	21.4	377
35	Addition of a polygenic risk score, mammographic density, and endogenous hormones to existing breast cancer risk prediction models: A nested case–control study. PLoS Medicine, 2018, 15, e1002644.	8.4	91
36	<i>IRGM</i> Gene Variants Modify the Relationship Between Visceral Adipose Tissue and NAFLD in Patients With Crohn's Disease. Inflammatory Bowel Diseases, 2018, 24, 2247-2257.	1.9	19

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37	Hierarchical modeling of melanocortin 1 receptor variants with skin cancer risk. Genetic Epidemiology, 2018, 42, 571-586.	1.3	5
38	Evidence for large-scale gene-by-smoking interaction effects on pulmonary function. International Journal of Epidemiology, 2017, 46, dyw318.	1.9	36
39	Association Between Proton Pump Inhibitor Use and Cognitive Function in Women. Gastroenterology, 2017, 153, 971-979.e4.	1.3	70
40	Physical activity from menarche to first pregnancy and risk of breast cancer. International Journal of Cancer, 2016, 139, 1223-1230.	5.1	26
41	Breast Cancer Risk From Modifiable and Nonmodifiable Risk Factors Among White Women in the United States. JAMA Oncology, 2016, 2, 1295.	7.1	285
42	Red meat, poultry, and fish intake and breast cancer risk among Hispanic and Non-Hispanic white women: The Breast Cancer Health Disparities Study. Cancer Causes and Control, 2016, 27, 527-543.	1.8	18
43	Four Susceptibility Loci for Gallstone Disease Identified in a Meta-analysis of Genome-Wide Association Studies. Gastroenterology, 2016, 151, 351-363.e28.	1.3	74
44	Telomere structure and maintenance gene variants and risk of five cancer types. International Journal of Cancer, 2016, 139, 2655-2670.	5.1	43
45	Interactions between breast cancer susceptibility loci and menopausal hormone therapy in relationship to breast cancer in the Breast and Prostate Cancer Cohort Consortium. Breast Cancer Research and Treatment, 2016, 155, 531-540.	2.5	2
46	A Cross-Cancer Genetic Association Analysis of the DNA Repair and DNA Damage Signaling Pathways for Lung, Ovary, Prostate, Breast, and Colorectal Cancer. Cancer Epidemiology Biomarkers and Prevention, 2016, 25, 193-200.	2.5	66
47	Response to Day etÂal American Journal of Human Genetics, 2016, 98, 394-395.	6.2	1
48	Adjusting for Heritable Covariates Can Bias Effect Estimates in Genome-Wide Association Studies. American Journal of Human Genetics, 2015, 96, 329-339.	6.2	230
49	Association of breast cancer risk <i>loci</i> with breast cancer survival. International Journal of Cancer, 2015, 137, 2837-2845.	5.1	33
50	Vitamin D–Associated Genetic Variation and Risk of Breast Cancer in the Breast and Prostate Cancer Cohort Consortium (BPC3). Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 627-630.	2.5	20
51	Genetic risk variants associated with in situ breast cancer. Breast Cancer Research, 2015, 17, 82.	5.0	25
52	Testing calibration of risk models at extremes of disease risk. Biostatistics, 2015, 16, 143-154.	1.5	26
53	Joint Effects of Colorectal Cancer Susceptibility Loci, Circulating 25-Hydroxyvitamin D and Risk of Colorectal Cancer. PLoS ONE, 2014, 9, e92212.	2.5	12
54	Post-GWAS gene–environment interplay in breast cancer: results from the Breast and Prostate Cancer Cohort Consortium and a meta-analysis on 79 000 women. Human Molecular Genetics, 2014, 23, 5260-5270.	2.9	37

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#	Article	IF	CITATION
55	Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. Nature Genetics, 2013, 45, 385-391.	21.4	492
56	Identification of seven new prostate cancer susceptibility loci through a genome-wide association study. Nature Genetics, 2009, 41, 1116-1121.	21.4	389