Jonathan M Kocarnik

List of Publications by Year
in descending order

21 papers	**1,236** citations	623734 **14** h-index	713466 **21** g-index
22 all docs	**22** docs citations	**22** times ranked	**3759** citing authors

#	Article	IF	Citations
1	Genetic analyses of diverse populations improves discovery for complex traits. Nature, 2019, 570, 514-518.	27.8	679
2	Molecular phenotypes of colorectal cancer and potential clinical applications. Gastroenterology Report, 2015, 3, gov046.	1.3	105
3	Strategies for Enriching Variant Coverage in Candidate Disease Loci on a Multiethnic Genotyping Array. PLoS ONE, 2016, 11, e0167758.	2.5	72
4	Timing of Aspirin and Other Nonsteroidal Anti-Inflammatory Drug Use Among Patients With Colorectal Cancer in Relation to Tumor Markers and Survival. Journal of Clinical Oncology, 2017, 35, 2806-2813.	1.6	57
5	Genome-Wide Interaction Analyses between Genetic Variants and Alcohol Consumption and Smoking for Risk of Colorectal Cancer. PLoS Genetics, 2016, 12, e1006296.	3.5	38
6	Returning Pleiotropic Results From Genetic Testing to Patients and Research Participants. JAMA - Journal of the American Medical Association, 2014, 311, 795.	7.4	32
7	Multiancestral Analysis of Inflammation-Related Genetic Variants and C-Reactive Protein in the Population Architecture Using Genomics and Epidemiology Study. Circulation: Cardiovascular Genetics, 2014, 7, 178-188.	5.1	31
8	Relationship of prediagnostic body mass index with survival after colorectal cancer: Stage–specific associations. International Journal of Cancer, 2016, 139, 1065-1072.	5.1	26
9	Replication of Associations between GWAS SNPs and Melanoma Risk in the Population Architecture Using Genomics and Epidemiology (PAGE) Study. Journal of Investigative Dermatology, 2014, 134, 2049-2052.	0.7	21
10	Long–term weight loss after colorectal cancer diagnosis is associated with lower survival: The Colon Cancer Family Registry. Cancer, 2017, 123, 4701-4708.	4.1	20
11	Pleiotropic and Sex-Specific Effects of Cancer GWAS SNPs on Melanoma Risk in the Population Architecture Using Genomics and Epidemiology (PAGE) Study. PLoS ONE, 2015, 10, e0120491.	2.5	19
12	Cross-cancer pleiotropic analysis of endometrial cancer: PAGE and E2C2 consortia. Carcinogenesis, 2014, 35, 2068-2073.	2.8	18
13	Diagnostics for Pleiotropy in Mendelian Randomization Studies: Global and Individual Tests for Direct Effects. American Journal of Epidemiology, 2018, 187, 2672-2680.	3.4	18
14	Leptin gene variants and colorectal cancer risk: Sex-specific associations. PLoS ONE, 2018, 13, e0206519.	2.5	17
15	Pleiotropy of Cancer Susceptibility Variants on the Risk of Non-Hodgkin Lymphoma: The PAGE Consortium. PLoS ONE, 2014, 9, e89791.	2.5	16
16	Discovery, fine-mapping, and conditional analyses of genetic variants associated with C-reactive protein in multiethnic populations using the Metabochip in the Population Architecture using Genomics and Epidemiology (PAGE) study. Human Molecular Genetics, 2018, 27, 2940-2953.	2.9	16
17	No Evidence of Gene–Calcium Interactions from Genome-Wide Analysis of Colorectal Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 2971-2976.	2.5	9
18	Association of family history and survival in patients with colorectal cancer: a pooled analysis of eight epidemiologic studies. Cancer Medicine, 2018, 7, 2192-2199.	2.8	9

#	Article	IF	Citations
19	Cancer's global epidemiological transition and growth. Lancet, The, 2020, 395, 757-758.	13.7	7
20	Generalization and fine mapping of red blood cell trait genetic associations to multiâ€ethnic populations: The PAGE study. American Journal of Hematology, 2018, 93, 1061-1073.	4.1	5
21	Genetic analysis of hsCRP in American Indians: The Strong Heart Family Study. PLoS ONE, 2019, 14, e0223574.	2.5	5