

Paul D P Pharoah

List of Publications by Year in descending order

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Version: 2024-02-01

829
papers

78,883
citations

418

132
h-index

872

243
g-index

892
all docs

892
docs citations

892
times ranked

62856
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 1 | Risk-Adjusted Cancer Screening and Prevention (RiskAP): Complementing Screening for Early Disease Detection by a Learning Screening Based on Risk Factors. <i>Breast Care</i> , 2022, 17, 208-223. | 0.8 | 6 |
| 2 | Comprehensive epithelial tubo-ovarian cancer risk prediction model incorporating genetic and epidemiological risk factors. <i>Journal of Medical Genetics</i> , 2022, 59, 632-643. | 1.5 | 33 |
| 3 | MCM3 is a novel proliferation marker associated with longer survival for patients with tubo-ovarian high-grade serous carcinoma. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2022, 480, 855-871. | 1.4 | 8 |
| 4 | High Prediagnosis Inflammation-Related Risk Score Associated with Decreased Ovarian Cancer Survival. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 443-452. | 1.1 | 2 |
| 5 | Functional analysis of the 1p34.3 risk locus implicates GNL2 in high-grade serous ovarian cancer. <i>American Journal of Human Genetics</i> , 2022, 109, 116-135. | 2.6 | 3 |
| 6 | Reproductive factors do not influence survival with ovarian cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, , cebp.1091.2021. | 1.1 | 1 |
| 7 | Rare germline copy number variants (CNVs) and breast cancer risk. <i>Communications Biology</i> , 2022, 5, 65. | 2.0 | 6 |
| 8 | Polygenic risk modeling for prediction of epithelial ovarian cancer risk. <i>European Journal of Human Genetics</i> , 2022, 30, 349-362. | 1.4 | 23 |
| 9 | Common variants in breast cancer risk loci predispose to distinct tumor subtypes. <i>Breast Cancer Research</i> , 2022, 24, 2. | 2.2 | 15 |
| 10 | Pathology of Tumors Associated With Pathogenic Germline Variants in 9 Breast Cancer Susceptibility Genes. <i>JAMA Oncology</i> , 2022, 8, e216744. | 3.4 | 51 |
| 11 | Concerns About Methods Used in Modeling Study of Risk-Stratified Screening for Breast Cancer. <i>JAMA Oncology</i> , 2022, , . | 3.4 | 3 |
| 12 | Validated biomarker assays confirm that <i>ARID1A</i> loss is confounded with <i>MMR</i> deficiency, <i>CD8</i> ⁺ TIL infiltration, and provides no independent prognostic value in endometriosis-associated ovarian carcinomas. <i>Journal of Pathology</i> , 2022, 256, 388-401. | 2.1 | 15 |
| 13 | Multi-omic machine learning predictor of breast cancer therapy response. <i>Nature</i> , 2022, 601, 623-629. | 13.7 | 187 |
| 14 | Large-scale Integrated Analysis of Genetics and Metabolomic Data Reveals Potential Links Between Lipids and Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 1216-1226. | 1.1 | 3 |
| 15 | A multi-level investigation of the genetic relationship between endometriosis and ovarian cancer histotypes. <i>Cell Reports Medicine</i> , 2022, 3, 100542. | 3.3 | 26 |
| 16 | Germline BRCA variants, lifestyle and ovarian cancer survival. <i>Gynecologic Oncology</i> , 2022, , . | 0.6 | 2 |
| 17 | A Genome-Wide Gene-Based Gene-Environment Interaction Study of Breast Cancer in More than 90,000 Women. <i>Cancer Research Communications</i> , 2022, 2, 211-219. | 0.7 | 6 |
| 18 | Genome-wide and transcriptome-wide association studies of mammographic density phenotypes reveal novel loci. <i>Breast Cancer Research</i> , 2022, 24, 27. | 2.2 | 15 |

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|----|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|-----------|
| 19 | Beyond GWAS of Colorectal Cancer: Evidence of Interaction with Alcohol Consumption and Putative Causal Variant for the 10q24.2 Region. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 1077-1089. | 1.1 | 6 |
| 20 | Genetic Regulation of DNA Methylation Yields Novel Discoveries in GWAS of Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 1068-1076. | 1.1 | 1 |
| 21 | Polygenic risk scores for prediction of breast cancer risk in women of African ancestry: a cross-ancestry approach. <i>Human Molecular Genetics</i> , 2022, 31, 3133-3143. | 1.4 | 11 |
| 22 | Relevance of the MHC region for breast cancer susceptibility in Asians. <i>Breast Cancer</i> , 2022, 29, 869-879. | 1.3 | 1 |
| 23 | Breast cancer risks associated with missense variants in breast cancer susceptibility genes. <i>Genome Medicine</i> , 2022, 14, 51. | 3.6 | 19 |
| 24 | Distinct Reproductive Risk Profiles for Intrinsic-Like Breast Cancer Subtypes: Pooled Analysis of Population-Based Studies. <i>Journal of the National Cancer Institute</i> , 2022, 114, 1706-1719. | 3.0 | 14 |
| 25 | Associations of a breast cancer polygenic risk score with tumor characteristics and survival.. <i>Journal of Clinical Oncology</i> , 2022, 40, 563-563. | 0.8 | 1 |
| 26 | Molecular Subclasses of Clear Cell Ovarian Carcinoma and Their Impact on Disease Behavior and Outcomes. <i>Clinical Cancer Research</i> , 2022, 28, 4947-4956. | 3.2 | 22 |
| 27 | chromMAGMA: regulatory element-centric interrogation of risk variants. <i>Life Science Alliance</i> , 2022, 5, e202201446. | 1.3 | 1 |
| 28 | Uncovering the Contribution of Moderate-Penetrance Susceptibility Genes to Breast Cancer by Whole-Exome Sequencing and Targeted Enrichment Sequencing of Candidate Genes in Women of European Ancestry. <i>Cancers</i> , 2022, 14, 3363. | 1.7 | 2 |
| 29 | Germline allelic expression of genes at 17q22 locus associates with risk of breast cancer. <i>European Journal of Cancer</i> , 2022, 172, 146-157. | 1.3 | 0 |
| 30 | Combined Associations of a Polygenic Risk Score and Classical Risk Factors With Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2021, 113, 329-337. | 3.0 | 45 |
| 31 | Population-based targeted sequencing of 54 candidate genes identifies <i>PALB2</i> as a susceptibility gene for high-grade serous ovarian cancer. <i>Journal of Medical Genetics</i> , 2021, 58, 305-313. | 1.5 | 26 |
| 32 | Common Susceptibility Loci for Male Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2021, 113, 453-461. | 3.0 | 12 |
| 33 | Expanding Our Understanding of Ovarian Cancer Risk: The Role of Incomplete Pregnancies. <i>Journal of the National Cancer Institute</i> , 2021, 113, 301-308. | 3.0 | 8 |
| 34 | Refined cut-off for TP53 immunohistochemistry improves prediction of TP53 mutation status in ovarian mucinous tumors: implications for outcome analyses. <i>Modern Pathology</i> , 2021, 34, 194-206. | 2.9 | 21 |
| 35 | Identifying Novel Susceptibility Genes for Colorectal Cancer Risk From a Transcriptome-Wide Association Study of 125,478 Subjects. <i>Gastroenterology</i> , 2021, 160, 1164-1178.e6. | 0.6 | 36 |
| 36 | Genetically predicted circulating protein biomarkers and ovarian cancer risk. <i>Gynecologic Oncology</i> , 2021, 160, 506-513. | 0.6 | 12 |

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|----|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 37 | A comprehensive re-assessment of the association between vitamin D and cancer susceptibility using Mendelian randomization. <i>Nature Communications</i> , 2021, 12, 246. | 5.8 | 39 |
| 38 | CYP3A7*1C allele: linking premenopausal oestrogen and progesterone levels with risk of hormone receptor-positive breast cancers. <i>British Journal of Cancer</i> , 2021, 124, 842-854. | 2.9 | 5 |
| 39 | Genetically predicted circulating concentrations of micronutrients and risk of colorectal cancer among individuals of European descent: a Mendelian randomization study. <i>American Journal of Clinical Nutrition</i> , 2021, 113, 1490-1502. | 2.2 | 27 |
| 40 | NHS announces a pilot of a blood test for early detection of many cancers. <i>Journal of Medical Screening</i> , 2021, 28, 1-2. | 1.1 | 2 |
| 41 | A case-only study to identify genetic modifiers of breast cancer risk for BRCA1/BRCA2 mutation carriers. <i>Nature Communications</i> , 2021, 12, 1078. | 5.8 | 19 |
| 42 | Genetic architectures of proximal and distal colorectal cancer are partly distinct. <i>Gut</i> , 2021, 70, 1325-1334. | 6.1 | 44 |
| 43 | Meta-analysis of the association between emphysematous change on thoracic computerized tomography scan and recurrent pneumothorax. <i>QJM - Monthly Journal of the Association of Physicians</i> , 2021, , . | 0.2 | 0 |
| 44 | Breast Cancer Risk Genes Association Analysis in More than 113,000 Women. <i>New England Journal of Medicine</i> , 2021, 384, 428-439. | 13.9 | 532 |
| 45 | Germline and Somatic Genetic Variants in the p53 Pathway Interact to Affect Cancer Risk, Progression, and Drug Response. <i>Cancer Research</i> , 2021, 81, 1667-1680. | 0.4 | 32 |
| 46 | Benefit, Harm, and Cost-effectiveness Associated With Magnetic Resonance Imaging Before Biopsy in Age-based and Risk-stratified Screening for Prostate Cancer. <i>JAMA Network Open</i> , 2021, 4, e2037657. | 2.8 | 34 |
| 47 | Evaluating Polygenic Risk Scores for Breast Cancer in Women of African Ancestry. <i>Journal of the National Cancer Institute</i> , 2021, 113, 1168-1176. | 3.0 | 41 |
| 48 | Response to Li and Hopper. <i>American Journal of Human Genetics</i> , 2021, 108, 527-529. | 2.6 | 5 |
| 49 | Gene-Environment Interactions Relevant to Estrogen and Risk of Breast Cancer: Can Gene-Environment Interactions Be Detected Only among Candidate SNPs from Genome-Wide Association Studies?. <i>Cancers</i> , 2021, 13, 2370. | 1.7 | 4 |
| 50 | Joint IARC/NCI International Cancer Seminar Series Report: expert consensus on future directions for ovarian carcinoma research. <i>Carcinogenesis</i> , 2021, 42, 785-793. | 1.3 | 6 |
| 51 | Pleiotropy-guided transcriptome imputation from normal and tumor tissues identifies candidate susceptibility genes for breast and ovarian cancer. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100042. | 1.0 | 6 |
| 52 | Identification of a Locus Near <i>ULK1</i> Associated With Progression-Free Survival in Ovarian Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 1669-1680. | 1.1 | 5 |
| 53 | Cross-ancestry GWAS meta-analysis identifies six breast cancer loci in African and European ancestry women. <i>Nature Communications</i> , 2021, 12, 4198. | 5.8 | 24 |
| 54 | A systematic review assessing the existence of pneumothorax-only variants of FLCN. Implications for lifelong surveillance of renal tumours. <i>European Journal of Human Genetics</i> , 2021, 29, 1595-1600. | 1.4 | 12 |

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|----|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 55 | Functional annotation of the 2q35 breast cancer risk locus implicates a structural variant in influencing activity of a long-range enhancer element. <i>American Journal of Human Genetics</i> , 2021, 108, 1190-1203. | 2.6 | 6 |
| 56 | Genetic analyses of gynecological disease identify genetic relationships between uterine fibroids and endometrial cancer, and a novel endometrial cancer genetic risk region at the WNT4 1p36.12 locus. <i>Human Genetics</i> , 2021, 140, 1353-1365. | 1.8 | 18 |
| 57 | Large-scale cross-cancer fine-mapping of the 5p15.33 region reveals multiple independent signals. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100041. | 1.0 | 6 |
| 58 | Association of germline genetic variants with breast cancer-specific survival in patient subgroups defined by clinic-pathological variables related to tumor biology and type of systemic treatment. <i>Breast Cancer Research</i> , 2021, 23, 86. | 2.2 | 7 |
| 59 | Mendelian randomisation study of smoking exposure in relation to breast cancer risk. <i>British Journal of Cancer</i> , 2021, 125, 1135-1145. | 2.9 | 9 |
| 60 | Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021, 596, 393-397. | 13.7 | 183 |
| 61 | Genomic risk prediction of coronary artery disease in women with breast cancer: a prospective cohort study. <i>Breast Cancer Research</i> , 2021, 23, 94. | 2.2 | 4 |
| 62 | Clinical Impact of the Predict Prostate Risk Communication Tool in Men Newly Diagnosed with Nonmetastatic Prostate Cancer: A Multicentre Randomised Controlled Trial. <i>European Urology</i> , 2021, 80, 661-669. | 0.9 | 7 |
| 63 | Breast Cancer Risk Factors and Survival by Tumor Subtype: Pooled Analyses from the Breast Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 623-642. | 1.1 | 19 |
| 64 | Long-term opiate use and risk of cardiovascular mortality: results from the Golestan Cohort Study. <i>European Journal of Preventive Cardiology</i> , 2021, 28, 98-106. | 0.8 | 13 |
| 65 | A Combined Proteomics and Mendelian Randomization Approach to Investigate the Effects of Aspirin-Targeted Proteins on Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 564-575. | 1.1 | 10 |
| 66 | Germline variants and breast cancer survival in patients with distant metastases at primary breast cancer diagnosis. <i>Scientific Reports</i> , 2021, 11, 19787. | 1.6 | 2 |
| 67 | Care of men with cancer-predisposing BRCA variants. <i>BMJ</i> , The, 2021, 375, n2376. | 3.0 | 1 |
| 68 | Salicylic Acid and Risk of Colorectal Cancer: A Two-Sample Mendelian Randomization Study. <i>Nutrients</i> , 2021, 13, 4164. | 1.7 | 3 |
| 69 | Meta-analysis of up to 622,409 individuals identifies 40 novel smoking behaviour associated genetic loci. <i>Molecular Psychiatry</i> , 2020, 25, 2392-2409. | 4.1 | 83 |
| 70 | Radiogenomics Consortium Genome-Wide Association Study Meta-Analysis of Late Toxicity After Prostate Cancer Radiotherapy. <i>Journal of the National Cancer Institute</i> , 2020, 112, 179-190. | 3.0 | 71 |
| 71 | Genetically Predicted Levels of DNA Methylation Biomarkers and Breast Cancer Risk: Data From 228,951 Women of European Descent. <i>Journal of the National Cancer Institute</i> , 2020, 112, 295-304. | 3.0 | 35 |
| 72 | A prospective study of tea drinking temperature and risk of esophageal squamous cell carcinoma. <i>International Journal of Cancer</i> , 2020, 146, 18-25. | 2.3 | 57 |

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|----|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|-----------|
| 73 | Assessment of interactions between 205 breast cancer susceptibility loci and 13 established risk factors in relation to breast cancer risk, in the Breast Cancer Association Consortium. <i>International Journal of Epidemiology</i> , 2020, 49, 216-232. | 0.9 | 21 |
| 74 | Evaluation of associations between genetically predicted circulating protein biomarkers and breast cancer risk. <i>International Journal of Cancer</i> , 2020, 146, 2130-2138. | 2.3 | 13 |
| 75 | Identification of novel epithelial ovarian cancer loci in women of African ancestry. <i>International Journal of Cancer</i> , 2020, 146, 2987-2998. | 2.3 | 18 |
| 76 | Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73. | 9.4 | 120 |
| 77 | Cumulative Burden of Colorectal Cancer-associated Genetic Variants Is More Strongly Associated With Early-Onset vs Late-Onset Cancer. <i>Gastroenterology</i> , 2020, 158, 1274-1286.e12. | 0.6 | 110 |
| 78 | A Mendelian randomization analysis of circulating lipid traits and breast cancer risk. <i>International Journal of Epidemiology</i> , 2020, 49, 1117-1131. | 0.9 | 41 |
| 79 | Identification of Novel Loci and New Risk Variant in Known Loci for Colorectal Cancer Risk in East Asians. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 477-486. | 1.1 | 25 |
| 80 | Cancer Risks Associated With Germline <i>PALB2</i> Pathogenic Variants: An International Study of 524 Families. <i>Journal of Clinical Oncology</i> , 2020, 38, 674-685. | 0.8 | 270 |
| 81 | Association Between Levels of Sex Hormones and Risk of Esophageal Adenocarcinoma and Barrett's Esophagus. <i>Clinical Gastroenterology and Hepatology</i> , 2020, 18, 2701-2709.e3. | 2.4 | 12 |
| 82 | Ovarian Cancer Risk Variants Are Enriched in Histotype-Specific Enhancers and Disrupt Transcription Factor Binding Sites. <i>American Journal of Human Genetics</i> , 2020, 107, 622-635. | 2.6 | 14 |
| 83 | Cancer Screening, Surrogates of Survival, and the Soma. <i>Cancer Cell</i> , 2020, 38, 433-437. | 7.7 | 14 |
| 84 | Breast cancer risk factors and their effects on survival: a Mendelian randomisation study. <i>BMC Medicine</i> , 2020, 18, 327. | 2.3 | 40 |
| 85 | Genome-wide Modeling of Polygenic Risk Score in Colorectal Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 432-444. | 2.6 | 124 |
| 86 | European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. <i>Nature Communications</i> , 2020, 11, 3833. | 5.8 | 88 |
| 87 | Cancer therapy shapes the fitness landscape of clonal hematopoiesis. <i>Nature Genetics</i> , 2020, 52, 1219-1226. | 9.4 | 367 |
| 88 | Rare Germline Genetic Variants and the Risks of Epithelial Ovarian Cancer. <i>Cancers</i> , 2020, 12, 3046. | 1.7 | 22 |
| 89 | Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 837-848. | 2.6 | 39 |
| 90 | Circulating bilirubin levels and risk of colorectal cancer: serological and Mendelian randomization analyses. <i>BMC Medicine</i> , 2020, 18, 229. | 2.3 | 28 |

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|-----|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 91 | Sex-Specific Genetic Associations for Barrett's Esophagus and Esophageal Adenocarcinoma. <i>Gastroenterology</i> , 2020, 159, 2065-2076.e1. | 0.6 | 16 |
| 92 | Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. <i>Nature Genetics</i> , 2020, 52, 572-581. | 9.4 | 265 |
| 93 | The challenge of early detection in cancer. <i>Science</i> , 2020, 368, 589-590. | 6.0 | 70 |
| 94 | Usual physical activity and subsequent hospital usage over 20 years in a general population: the EPIC-Norfolk cohort. <i>BMC Geriatrics</i> , 2020, 20, 165. | 1.1 | 10 |
| 95 | The effect of sample size on polygenic hazard models for prostate cancer. <i>European Journal of Human Genetics</i> , 2020, 28, 1467-1475. | 1.4 | 14 |
| 96 | Comparative performance and external validation of the multivariable PREDICT Prostate tool for non-metastatic prostate cancer: a study in 69,206 men from Prostate Cancer data Base Sweden (PCBaSe). <i>BMC Medicine</i> , 2020, 18, 139. | 2.3 | 10 |
| 97 | Personalized early detection and prevention of breast cancer: ENVISION consensus statement. <i>Nature Reviews Clinical Oncology</i> , 2020, 17, 687-705. | 12.5 | 178 |
| 98 | Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. <i>Scientific Reports</i> , 2020, 10, 9688. | 1.6 | 2 |
| 99 | Development and Validation of the Gene Expression Predictor of High-grade Serous Ovarian Carcinoma Molecular SubTYPE (ProTYPE). <i>Clinical Cancer Research</i> , 2020, 26, 5411-5423. | 3.2 | 43 |
| 100 | Household Fuel Use and the Risk of Gastrointestinal Cancers: The Golestan Cohort Study. <i>Environmental Health Perspectives</i> , 2020, 128, 67002. | 2.8 | 19 |
| 101 | Clinical and pathological associations of PTEN expression in ovarian cancer: a multicentre study from the Ovarian Tumour Tissue Analysis Consortium. <i>British Journal of Cancer</i> , 2020, 123, 793-802. | 2.9 | 35 |
| 102 | Mendelian Randomization of Circulating Polyunsaturated Fatty Acids and Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 860-870. | 1.1 | 26 |
| 103 | Identification of novel breast cancer susceptibility loci in meta-analyses conducted among Asian and European descendants. <i>Nature Communications</i> , 2020, 11, 1217. | 5.8 | 46 |
| 104 | Functional informed genome-wide interaction analysis of body mass index, diabetes and colorectal cancer risk. <i>Cancer Medicine</i> , 2020, 9, 3563-3573. | 1.3 | 7 |
| 105 | A Genetic Risk Score to Personalize Prostate Cancer Screening, Applied to Population Data. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1731-1738. | 1.1 | 27 |
| 106 | Menopausal hormone therapy prior to the diagnosis of ovarian cancer is associated with improved survival. <i>Gynecologic Oncology</i> , 2020, 158, 702-709. | 0.6 | 15 |
| 107 | Assessment of polygenic architecture and risk prediction based on common variants across fourteen cancers. <i>Nature Communications</i> , 2020, 11, 3353. | 5.8 | 75 |
| 108 | Urinary TERT promoter mutations are detectable up to 10 years prior to clinical diagnosis of bladder cancer: Evidence from the Golestan Cohort Study. <i>EBioMedicine</i> , 2020, 53, 102643. | 2.7 | 51 |

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|-----|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|-----------|
| 109 | Ovarian and Breast Cancer Risks Associated With Pathogenic Variants in <i>RAD51C</i> and <i>RAD51D</i> . <i>Journal of the National Cancer Institute</i> , 2020, 112, 1242-1250. | 3.0 | 106 |
| 110 | Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468. | 0.6 | 32 |
| 111 | A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , 2020, 11, 312. | 5.8 | 30 |
| 112 | Opium use and subsequent incidence of cancer: results from the Golestan Cohort Study. <i>The Lancet Global Health</i> , 2020, 8, e649-e660. | 2.9 | 59 |
| 113 | Prediction of contralateral breast cancer: external validation of risk calculators in 20 international cohorts. <i>Breast Cancer Research and Treatment</i> , 2020, 181, 423-434. | 1.1 | 14 |
| 114 | Combining measures of immune infiltration shows additive effect on survival prediction in high-grade serous ovarian carcinoma. <i>British Journal of Cancer</i> , 2020, 122, 1803-1810. | 2.9 | 23 |
| 115 | Genome-wide association meta-analyses combining multiple risk phenotypes provide insights into the genetic architecture of cutaneous melanoma susceptibility. <i>Nature Genetics</i> , 2020, 52, 494-504. | 9.4 | 138 |
| 116 | Physical activity and risks of breast and colorectal cancer: a Mendelian randomisation analysis. <i>Nature Communications</i> , 2020, 11, 597. | 5.8 | 193 |
| 117 | Genetic Data from Nearly 63,000 Women of European Descent Predicts DNA Methylation Biomarkers and Epithelial Ovarian Cancer Risk. <i>Cancer Research</i> , 2019, 79, 505-517. | 0.4 | 49 |
| 118 | Novel Common Genetic Susceptibility Loci for Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2019, 111, 146-157. | 3.0 | 129 |
| 119 | Sex specific associations in genome wide association analysis of renal cell carcinoma. <i>European Journal of Human Genetics</i> , 2019, 27, 1589-1598. | 1.4 | 27 |
| 120 | The <i>FANCM</i> :p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. <i>Npj Breast Cancer</i> , 2019, 5, 38. | 2.3 | 28 |
| 121 | Two truncating variants in <i>FANCC</i> and breast cancer risk. <i>Scientific Reports</i> , 2019, 9, 12524. | 1.6 | 5 |
| 122 | Evidence of a Causal Association Between Cancer and Alzheimer's Disease: a Mendelian Randomization Analysis. <i>Scientific Reports</i> , 2019, 9, 13548. | 1.6 | 26 |
| 123 | Understanding of prognosis in non-metastatic prostate cancer: a randomised comparative study of clinician estimates measured against the PREDICT prostate prognostic model. <i>British Journal of Cancer</i> , 2019, 121, 715-718. | 2.9 | 12 |
| 124 | Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431. | 5.8 | 88 |
| 125 | Overestimation of the Benefit-to-Harm Ratio of Risk-Based Mammography Screening in the United Kingdom—Reply. <i>JAMA Oncology</i> , 2019, 5, 428. | 3.4 | 1 |
| 126 | Models predicting survival to guide treatment decision-making in newly diagnosed primary non-metastatic prostate cancer: a systematic review. <i>BMJ Open</i> , 2019, 9, e029149. | 0.8 | 15 |

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|-----|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 127 | A combination of the immunohistochemical markers CK7 and SATB2 is highly sensitive and specific for distinguishing primary ovarian mucinous tumors from colorectal and appendiceal metastases. <i>Modern Pathology</i> , 2019, 32, 1834-1846. | 2.9 | 54 |
| 128 | Authors'™ response: Associations of obesity and circulating insulin and glucose with breast cancer risk. <i>International Journal of Epidemiology</i> , 2019, 48, 1016-1017. | 0.9 | 1 |
| 129 | The functional ALDH2 polymorphism is associated with breast cancer risk: A pooled analysis from the Breast Cancer Association Consortium. <i>Molecular Genetics & Genomic Medicine</i> , 2019, 7, e707. | 0.6 | 9 |
| 130 | Going to extremes: determinants of extraordinary response and survival in patients with cancer. <i>Nature Reviews Cancer</i> , 2019, 19, 339-348. | 12.8 | 35 |
| 131 | Association analyses identify 31 new risk loci for colorectal cancer susceptibility. <i>Nature Communications</i> , 2019, 10, 2154. | 5.8 | 172 |
| 132 | Evaluation of vitamin D biosynthesis and pathway target genes reveals UGT2A1/2 and EGFR polymorphisms associated with epithelial ovarian cancer in African American Women. <i>Cancer Medicine</i> , 2019, 8, 2503-2513. | 1.3 | 6 |
| 133 | A transcriptome-wide association study of high-grade serous epithelial ovarian cancer identifies new susceptibility genes and splice variants. <i>Nature Genetics</i> , 2019, 51, 815-823. | 9.4 | 89 |
| 134 | Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , 2019, 10, 1741. | 5.8 | 90 |
| 135 | Individual and Combined Effects of Environmental Risk Factors for Esophageal Cancer Based on Results From the Golestan Cohort Study. <i>Gastroenterology</i> , 2019, 156, 1416-1427. | 0.6 | 123 |
| 136 | Genome-wide association studies identify susceptibility loci for epithelial ovarian cancer in east Asian women. <i>Gynecologic Oncology</i> , 2019, 153, 343-355. | 0.6 | 28 |
| 137 | Individual prognosis at diagnosis in nonmetastatic prostate cancer: Development and external validation of the PREDICT Prostate multivariable model. <i>PLoS Medicine</i> , 2019, 16, e1002758. | 3.9 | 56 |
| 138 | Dynamics of breast-cancer relapse reveal late-recurring ER-positive genomic subgroups. <i>Nature</i> , 2019, 567, 399-404. | 13.7 | 239 |
| 139 | Combined quantitative measures of ER, PR, HER2, and KI67 provide more prognostic information than categorical combinations in luminal breast cancer. <i>Modern Pathology</i> , 2019, 32, 1244-1256. | 2.9 | 51 |
| 140 | Body mass index and the association between low-density lipoprotein cholesterol as predicted by HMGCR genetic variants and breast cancer risk. <i>International Journal of Epidemiology</i> , 2019, 48, 1727-1730. | 0.9 | 3 |
| 141 | No Association Between Vitamin D Status and Risk of Barrett's Esophagus or Esophageal Adenocarcinoma: A Mendelian Randomization Study. <i>Clinical Gastroenterology and Hepatology</i> , 2019, 17, 2227-2235.e1. | 2.4 | 16 |
| 142 | Genome-wide association study of germline variants and breast cancer-specific mortality. <i>British Journal of Cancer</i> , 2019, 120, 647-657. | 2.9 | 52 |
| 143 | The association between weight at birth and breast cancer risk revisited using Mendelian randomisation. <i>European Journal of Epidemiology</i> , 2019, 34, 591-600. | 2.5 | 16 |
| 144 | Genetic variant predictors of gene expression provide new insight into risk of colorectal cancer. <i>Human Genetics</i> , 2019, 138, 307-326. | 1.8 | 44 |

| # | ARTICLE | IF | CITATIONS |
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| 145 | Prediction and clinical utility of a contralateral breast cancer risk model. <i>Breast Cancer Research</i> , 2019, 21, 144. | 2.2 | 24 |
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