# Agnieszka Dansonka-Mieszkowska

List of Publications by Year
in descending order



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|---|---|---|---|
| 52 papers | 3,657 citations | 201674 27 h-index | 175258 52 g-index |
| 52 all docs | 52 docs citations | 52 times ranked | 7051 citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. Nature Genetics, 2013, 45, 371-384. | 21.4 | 493 |
| 2 | Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. Nature Genetics, 2017, 49, 680-691. | 21.4 | 356 |
| 3 | GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. Nature Genetics, 2013, 45, 362-370. | 21.4 | 326 |
| 4 | A great majority of GISTs with PDGFRA mutations represent gastric tumors of low or no malignant potential. Laboratory Investigation, 2004, 84, 874-883. | 3.7 | 292 |
| 5 | Identification of six new susceptibility loci for invasive epithelial ovarian cancer. Nature Genetics, 2015, 47, 164-171. | 21.4 | 221 |
| 6 | Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. Cancer Discovery, 2016, 6, 1052-1067. | 9.4 | 157 |
| 7 | Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. Nature Communications, 2013, 4, 1628. | 12.8 | 144 |
| 8 | Association of vitamin D levels and risk of ovarian cancer: a Mendelian randomization study. International Journal of Epidemiology, 2016, 45, 1619-1630. | 1.9 | 111 |
| 9 | Gastrointestinal Stromal Tumors with Internal Tandem Duplications in 3' End of KIT Juxtamembrane Domain Occur Predominantly in Stomach and Generally Seem to Have a Favorable Course. Modern Pathology, 2003, 16, 1257-1264. | 5.5 | 104 |
| 10 | Identification and molecular characterization of a new ovarian cancer susceptibility locus at 17q21.31. Nature Communications, 2013, 4, 1627. | 12.8 | 98 |
| 11 | A novel germline PALB2 deletion in Polish breast and ovarian cancer patients. BMC Medical Genetics, 2010, 11, 20. | 2.1 | 96 |
| 12 | Ovarian small cell carcinoma of hypercalcemic type â€“ evidence of germline origin and smarca4 gene inactivation. a pilot study. Polish Journal of Pathology, 2013, 4, 238-246. | 0.3 | 85 |
| 13 | PIK3CA amplification associates with resistance to chemotherapy in ovarian cancer patients. Cancer Biology and Therapy, 2009, 8, 21-26. | 3.4 | 81 |
| 14 | Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breastâ€“ovarian cancer susceptibility locus. Nature Communications, 2016, 7, 12675. | 12.8 | 78 |
| 15 | Adult body mass index and risk of ovarian cancer by subtype: a Mendelian randomization study. International Journal of Epidemiology, 2016, 45, 884-895. | 1.9 | 71 |
| 16 | Shared genetics underlying epidemiological association between endometriosis and ovarian cancer. Human Molecular Genetics, 2015, 24, 5955-5964. | 2.9 | 68 |
| 17 | Evaluation of NF2 and NF1 Tumor Suppressor Genes in Distinctive Gastrointestinal Nerve Sheath Tumors Traditionally Diagnosed as Benign Schwannomas: A Study of 20 Cases. Laboratory Investigation, 2003, 83, 1361-1371. | 3.7 | 65 |
| 18 | Cis-eQTL analysis and functional validation of candidate susceptibility genes for high-grade serous ovarian cancer. Nature Communications, 2015, 6, 8234. | 12.8 | 63 |

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|---|---|---|---|
| 19 | The putative oncogene, CRNDE, is a negative prognostic factor in ovarian cancer patients. Oncotarget, 2015, 6, 43897-43910. | 1.8 | 51 |
| 20 | Risk of Ovarian Cancer and the NF-ÎºB Pathway: Genetic Association with IL1A and TNFSF10. Cancer Research, 2014, 74, 852-861. | 0.9 | 48 |
| 21 | Common Genetic Variation In Cellular Transport Genes and Epithelial Ovarian Cancer (EOC) Risk. PLoS ONE, 2015, 10, e0128106. | 2.5 | 44 |
| 22 | The Novel Gene CRNDE Encodes a Nuclear Peptide (CRNDEP) Which Is Overexpressed in Highly Proliferating Tissues. PLoS ONE, 2015, 10, e0127475. | 2.5 | 40 |
| 23 | Cell-type-specific enrichment of risk-associated regulatory elements at ovarian cancer susceptibility loci. Human Molecular Genetics, 2015, 24, 3595-3607. | 2.9 | 40 |
| 24 | Evidence of a genetic link between endometriosis and ovarian cancer. Fertility and Sterility, 2016, 105, 35-43.e10. | 1.0 | 37 |
| 25 | Loss of heterozygosity on chromosome 22q in gastrointestinal stromal tumors (GISTs): a study on 50 cases. Laboratory Investigation, 2005, 85, 237-247. | 3.7 | 34 |
| 26 | Genome-wide Analysis Identifies Novel Loci Associated with Ovarian Cancer Outcomes: Findings from the Ovarian Cancer Association Consortium. Clinical Cancer Research, 2015, 21, 5264-5276. | 7.0 | 33 |
| 27 | Germline SMARCA4 mutations in patients with ovarian small cell carcinoma of hypercalcemic type. Orphanet Journal of Rare Diseases, 2015, 10, 32. | 2.7 | 31 |
| 28 | Unsupervised analysis reveals two molecular subgroups of serous ovarian cancer with distinct gene expression profiles and survival. Journal of Cancer Research and Clinical Oncology, 2016, 142, 1239-1252. | 2.5 | 30 |
| 29 | Network-Based Integration of GWAS and Gene Expression Identifies a HOX-Centric Network Associated with Serous Ovarian Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1574-1584. | 2.5 | 28 |
| 30 | Common Genetic Variation in Circadian Rhythm Genes and Risk of Epithelial Ovarian Cancer (EOC). Journal of Genetics and Genome Research, 2015, 2, . | 0.3 | 25 |
| 31 | Genome-wide association study of subtype-specific epithelial ovarian cancer risk alleles using pooled DNA. Human Genetics, 2014, 133, 481-497. | 3.8 | 23 |
| 32 | Enrichment of putative PAX8 target genes at serous epithelial ovarian cancer susceptibility loci. British Journal of Cancer, 2017, 116, 524-535. | 6.4 | 23 |
| 33 | Epithelialâ€Mesenchymal Transition (EMT) Gene Variants and Epithelial Ovarian Cancer (EOC) Risk. Genetic Epidemiology, 2015, 39, 689-697. | 1.3 | 22 |
| 34 | Large-Scale Evaluation of Common Variation in Regulatory T Cellâ€“Related Genes and Ovarian Cancer Outcome. Cancer Immunology Research, 2014, 2, 332-340. | 3.4 | 21 |
| 35 | Analysis of Over 10,000 Cases Finds No Association between Previously Reported Candidate Polymorphisms and Ovarian Cancer Outcome. Cancer Epidemiology Biomarkers and Prevention, 2013, 22, 987-992. | 2.5 | 20 |
| 36 | Clinical importance of FANCD2, BRIP1, BRCA1, BRCA2 and FANCF expression in ovarian carcinomas. Cancer Biology and Therapy, 2019, 20, 843-854. | 3.4 | 20 |

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|---|---|---|---|
| 37 | Assessing the genetic architecture of epithelial ovarian cancer histological subtypes. Human Genetics, 2016, 135, 741-756. | 3.8 | 19 |
| 38 | No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. Gynecologic Oncology, 2016, 141, 386-401. | 1.4 | 18 |
| 39 | The significance of c.690G>T polymorphism (rs34529039) and expression of the CEBPA gene in ovarian cancer outcome. Oncotarget, 2016, 7, 67412-67424. | 1.8 | 17 |
| 40 | Consortium analysis of gene and geneâ€“folate interactions in purine and pyrimidine metabolism pathways with ovarian carcinoma risk. Molecular Nutrition and Food Research, 2014, 58, 2023-2035. | 3.3 | 16 |
| 41 | Evaluating the ovarian cancer gonadotropin hypothesis: A candidate gene study. Gynecologic Oncology, 2015, 136, 542-548. | 1.4 | 15 |
| 42 | Adult height is associated with increased risk of ovarian cancer: a Mendelian randomisation study. British Journal of Cancer, 2018, 118, 1123-1129. | 6.4 | 15 |
| 43 | Assessment of moderate coffee consumption and risk of epithelial ovarian cancer: a Mendelian randomization study. International Journal of Epidemiology, 2018, 47, 450-459. | 1.9 | 15 |
| 44 | Variation in NF-Î²B Signaling Pathways and Survival in Invasive Epithelial Ovarian Cancer. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 1421-1427. | 2.5 | 13 |
| 45 | Inherited variants affecting RNA editing may contribute to ovarian cancer susceptibility: results from a large-scale collaboration. Oncotarget, 2016, 7, 72381-72394. | 1.8 | 13 |
| 46 | p19<sup>INK4d</sup>mRNA and protein expression as new prognostic factors in ovarian cancer patients. Cancer Biology and Therapy, 2013, 14, 973-981. | 3.4 | 11 |
| 47 | Variants in genes encoding small GTPases and association with epithelial ovarian cancer susceptibility. PLoS ONE, 2018, 13, e0197561. | 2.5 | 9 |
| 48 | Assessment of variation in immunosuppressive pathway genes reveals TGFBR2 to be associated with risk of clear cell ovarian cancer. Oncotarget, 2016, 7, 69097-69110. | 1.8 | 5 |
| 49 | The utility of fluorescence in situ hybridization (FISH) in determining DNA damage-inducible transcript 3 (DDIT3) amplification in dedifferentiated liposarcomas â€“ an important diagnostic pitfall. Pathology Research and Practice, 2021, 225, 153555. | 2.3 | 4 |
| 50 | Clinical importance of the EMSY gene expression and polymorphisms in ovarian cancer. Oncotarget, 2018, 9, 17735-17755. | 1.8 | 4 |
| 51 | No Evidence That Genetic Variation in the Myeloid-Derived Suppressor Cell Pathway Influences Ovarian Cancer Survival. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 420-424. | 2.5 | 3 |
| 52 | Unique gastrointestinal stromal tumor with PDGFRA D842Y mutationâ€”evaluation of in vivo sensitivity to imatinib. Memo - Magazine of European Medical Oncology, 2021, 14, 208-213. | 0.5 | 1 |