



Australian Ovarian Cancer Study

Publications List

2020 Publications		
Author	Title of Paper	PubMed Link
Gorringe KL et al	<i>Therapeutic options for mucinous ovarian carcinoma</i>	Click to view

2019 Publications		
Author	Title of Paper	PubMed Link
Cheasley D et al	<i>The molecular origin and taxonomy of mucinous ovarian carcinoma</i>	Click to view
Christie EL et al	<i>Multiple ABCB1 transcriptional fusions in drug resistant high-grade serous ovarian and breast cancer</i>	Click to view
Harris HR et al	<i>Association between genetically predicted polycystic ovary syndrome and ovarian cancer: a Mendelian randomization study</i>	Click to view
Kondrashova O et al	<i>Clinical Utility of Real-Time Targeted Molecular Profiling in the Clinical Management of Ovarian Cancer: The ALLOCATE Study</i>	Click to view
Lawrenson K et al	<i>Genome-wide association studies identify susceptibility loci for epithelial ovarian cancer in east Asian women</i>	Click to view
Mavaddat N et al	<i>Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. American journal of human genetics</i>	Click to view
Minlikeeva AN et al	<i>Joint exposure to smoking, excessive weight, and physical inactivity and survival of ovarian cancer patients, evidence from the Ovarian Cancer Association Consortium. Cancer Causes Control</i>	Click to view
Nagle CM et al	<i>The association between the inflammatory potential of diet and risk of developing, and survival following, a diagnosis of ovarian cancer</i>	Click to view
Takenaka M et al	<i>Survival Following Chemotherapy in Ovarian Clear Cell Carcinoma Is Not Associated with Pathological Misclassification of Tumor Histotype</i>	Click to view

2018 Publications		
Author	Title of Paper	PubMed Link
Aziz D et al	<i>19q12 amplified and non-amplified subsets of high grade serous ovarian cancer with overexpression of cyclin E1 differ in their molecular drivers and clinical outcomes</i>	Click to view
Babic A et al	<i>Menstrual pain and risk of epithelial ovarian cancer: Results from the Ovarian Cancer Association Consortium</i>	Click to view
Block MS et al	<i>MyD88 and TRL4 Expression in Epithelial Ovarian Cancer</i>	Click to view
Colombo M et al	<i>The BRCA2 c.68-7T > A variant is not pathogenic: A model for clinical calibration of spliceogenicity</i>	Click to view
Dixon-Suen SC et al	<i>Adult height is associated with increased risk of ovarian cancer: a Medelian randomisation study</i>	Click to view
Dobrovic A	<i>White Blood Cell BRCA1 Promoter Methylation Status and Ovarian Cancer Risk: A Perspective</i>	Click to view

Author	Title of Paper	PubMed Link
Garsed DW et al	<i>Homologous Recombination DNA Repair Pathway Disruption and Retinoblastoma Protein Loss Are Associated with Exceptional Survival in High-Grade Serous Ovarian Cancer</i>	Click to view
Harris HR et al	<i>Polycystic Ovary Syndrome, Oligomenorrhea, and Risk of Ovarian Cancer Histotypes: Evidence from the Ovarian Cancer Association Consortium</i>	Click to view
Heinzelmann-Schwarz V et al	<i>Letrozole may be a valuable maintenance treatment in high-grade serous ovarian cancer patients</i>	Click to view
Hillman RT et al	<i>Genomic Rearrangement Signatures and Clinical Outcomes in High Grade Serous Ovarian Cancer</i>	Click to view
Kelemen LE et al	<i>rs495139 in the TYMS-ENOSF1 Region and Risk of Ovarian Carcinoma of Mucinous Histology</i>	Click to view
Kondrashova O et al	<i>Methylation of all BRCA1 copies predicts response to the PARP inhibitor rucaparib in ovarian carcinoma</i>	Click to view
Lindemann K et al	<i>Response rates to second-line platinum-based therapy in ovarian cancer patients challenge the clinical definition of platinum resistance</i>	Click to view
Moujaber T et al	<i>BRAF mutations in low-grade serous ovarian cancer and response to BRAF inhibition</i>	Click to view
O'Donnell T et al	<i>Chemotherapy weakly contributes to predicted neoantigen expression in ovarian cancer</i>	Click to view
Peres LC et al	<i>Racial/ethnic differences in the epidemiology of ovarian cancer: a pooled analysis of 12 case-control studies</i>	Click to view
Rambau PF et al	<i>Association of p16 expression with prognosis varies across ovarian carcinoma histotypes: an Ovarian Tumor Tissue Analysis consortium study</i>	Click to view
Ring BZ et al	<i>Transducin-Like Enhancer of Split 3 (TLE3) expression is associated with taxane sensitivity in non-serous ovarian carcinoma in a three-cohort study</i>	Click to view
Ugai T et al	<i>Ovarian cancer risk. ALDH2 polymorphism and alcohol drinking: Asian data from the ovarian cancer Association Consortium</i>	Click to view

2017 Publications		
Author	Title of Paper	PubMed Link
Babic A et al	<i>Menstrual pain and risk of epithelial ovarian cancer: Results from the Ovarian Cancer Association Consortium</i>	Click to view
Day FR et al	<i>Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk</i>	Click to view
Dixon SC et al	<i>Use of common analgesic medications and ovarian cancer survival: results from a pooled analysis in the Ovarian Cancer Association Consortium</i>	Click to view
Etemadmoghadam D et al	<i>EIF1AX and NRAS Mutations Co-occur and Cooperate in Low-Grade Serous Ovarian Carcinomas</i>	Click to view
Fagerholm R et al	<i>TP53-based interaction analysis identifies cis-eQTL variants for TP53BP2, FBXO28, and FAM53A that associate with survival and treatment outcome in breast cancer</i>	Click to view
Glubb DM et al	<i>Analyses of germline variants associated with ovarian cancer survival identify functional candidates at the 1q22 and 19p12 outcome loci</i>	Click to view
Goode, Ellen et al	<i>Dose-Response Relationship of CD8+ Tumor Infiltrating Lymphocytes and Survival Time in High-Grade Serous Ovarian Cancer</i>	Click to view
Guo Q et al	<i>Body mass index and breast cancer survival: a Mendelian randomization analysis</i>	Click to view
Jiao X et al	<i>PHIP – a novel candidate breast cancer susceptibility locus on 6q14.1</i>	Click to view
Kondrashova O et al	<i>Secondary Somatic Mutations Restoring RAD51C and RAD51D Associated with Acquired Resistance to the PARP Inhibitor Rucaparib in High-Grade Ovarian Carcinoma</i>	Click to view
Liu G et al	<i>Robust Tests for Additive Gene-Environment Interaction in Case-Control Studies Using Gene-Environment Independence</i>	Click to view
Michailidou K et al	<i>Association analysis identifies 65 new breast cancer risk loci</i>	Click to view
Milne RL et al	<i>Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer</i>	Click to view
Minlikeeva AN et al	<i>History of Comorbidities and Survival of Ovarian Cancer Patients, Results from the Ovarian Cancer Association Consortium</i>	Click to view

Author	Title of Paper	PubMed Link
Minlikeeva AN et al	<i>History of hypertension, heart disease, and diabetes and ovarian cancer patient survival: evidence from the ovarian cancer association consortium</i>	Click to view
Minlikeeva AN et al	<i>History of thyroid disease and survival of ovarian cancer patients: results from the Ovarian Cancer Association Consortium, a brief report</i>	Click to view
Phelan CM et al	<i>Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer</i>	Click to view
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Praestegaard C et al	<i>Cigarette smoking is associated with adverse survival among women with ovarian cancer: Results from a pooled analysis of 19 studies</i>	Click to view
Rasmussen CB et al	<i>Pelvic Inflammatory Disease and the Risk of Ovarian Cancer and Borderline Ovarian Tumors: A Pooled analysis of 13 Case-Control Studies</i>	Click to view
Reid BM et al	<i>Integration of Population-Level Genotype Data with Functional Annotation Reveals Over-Representation of Long Noncoding RNAs at Ovarian Cancer Susceptibility Loci</i>	Click to view
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Sucheston-Campbell LE et al	<i>No Evidence That Genetic Variation in the Myeloid-Derived Suppressor Cell Pathway Influences Ovarian Cancer Survival</i>	Click to view
Yang WL et al	<i>Elevation of TP53 Autoantibody Before CA125 in Preclinical Invasive Epithelial Ovarian Cancer</i>	Click to view

2016 Publications		
Author	Title of Paper	PubMed Link
Alsop K et al	<i>A community-based model of rapid autopsy in end-stage cancer patients</i>	Click to view
Bolton KL et al	<i>Corrigendum: Common variants at 19p13 are associated with susceptibility to ovarian cancer</i>	Click to view
Cannioto R et al	<i>Chronic Recreational Physical Inactivity and Epithelial Ovarian Cancer Risk: Evidence from the Ovarian Cancer Association Consortium</i>	Click to view
Cheng TH et al	<i>Five endometrial cancer risk loci identified through genome-wide association analysis</i>	Click to view
Cuellar-Partida G et al	<i>Assessing the genetic architecture of epithelial ovarian cancer histological subtypes</i>	Click to view
Dickson KA et al	<i>The RING finger domain E3 ubiquitin ligases BRCA1 and the RNF20/RNF40 complex in global loss of the chromatin mark histone H2B monoubiquitination (H2Bub1) in cell line models and primary high-grade serous ovarian cancer</i>	Click to view
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Ghousaini M et al	<i>Evidence that the 5p12 Variant rs10941679 Confers Susceptibility to Estrogen-Receptor-Positive Breast Cancer through FGF10 and MRPS30 Regulation</i>	Click to view
Hamdi Y et al	<i>Association of breast cancer risk with genetic variants showing differential allelic expression: Identification of a novel breast cancer susceptibility locus at 4q21</i>	Click to view
Hamprass SS et al	<i>Assessment of variation in immunosuppressive pathway genes reveals TGFBR2 to be associated with risk of clear cell ovarian cancer</i>	Click to view
Horne HN et al	<i>Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus</i>	Click to view
Jim HS et al	<i>Common Genetic Variation in Circadian Rhythm Genes and Risk of Epithelial Ovarian Cancer (EOC)</i>	Click to view

Author	Title of Paper	PubMed Link
Kar SP et al	<i>Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types</i>	Click to view
Lawrenson K et al	<i>Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus</i>	Click to view
Lei J et al	<i>Genetic variation in the immunosuppression pathway genes and breast cancer susceptibility: a pooled analysis of 42,510 cases and 40,577 controls from the Breast Cancer Association Consortium</i>	Click to view
Mercieca-Bebber RL et al	<i>Ovarian cancer study dropouts had worse health-related quality of life and psychosocial symptoms at baseline and over time</i>	Click to view
Ong JS et al	<i>Association of vitamin D levels and risk of ovarian cancer: a Mendelian randomization study</i>	Click to view
Painter JN et al	<i>Genetic Risk Score Mendelian Randomization Shows that Obesity Measured as Body Mass Index, but not Waist:Hip Ratio, Is Causal for Endometrial Cancer</i>	Click to view
Pelttari LM et al	<i>RAD51B in Familial Breast Cancer</i>	Click to view
Permuth JB et al	<i>Exome genotyping arrays to identify rare and low frequency variants associated with epithelial ovarian cancer risk</i>	Click to view
Permuth JB et al	<i>Inherited variants affecting RNA editing may contribute to ovarian cancer susceptibility: results from a large-scale collaboration</i>	Click to view
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Price MA et al	<i>Helplessness/hopelessness, minimization and optimism predict survival in women with invasive ovarian cancer: a role for targeted support during initial treatment decision-making?</i>	Click to view
Sinha D et al	<i>Pericytes Promote Malignant Ovarian Cancer Progression in Mice and Predict Poor Prognosis in Serous Ovarian Cancer Patients</i>	Click to view
Southey MC et al	<i>PALB2, CHEK2 and ATM rare variants and cancer risk: data from COGS</i>	Click to view
Thompson DJ et al	<i>CYP19A1 fine-mapping and Mendelian randomization: estradiol is causal for endometrial cancer</i>	Click to view
Winham SJ et al	<i>Investigation of Exomic Variants Associated with Overall Survival in Ovarian Cancer</i>	Click to view
Wyszynski A et al	<i>An intergenic risk locus containing an enhancer deletion in 2q35 modulates breast cancer risk by deregulating IGFBP5 expression</i>	Click to view
Zeng C et al	<i>Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus</i>	Click to view

2015 Publications		
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Amankwah EK et al	<i>Epithelial-Mesenchymal Transition (EMT) Gene Variants and Epithelial Ovarian Cancer (EOC) Risk</i>	Click to view
Candido-dos-Reis FJ et al	<i>Germline mutation in BRCA1 or BRCA2 and ten-year survival for women diagnosed with epithelial ovarian cancer</i>	Click to view
Carvajal-Carmona LG et al	<i>Candidate locus analysis of the TERT-CLPTM1L cancer risk region on chromosome 5p15 identifies multiple independent variants associated with endometrial cancer risk</i>	Click to view
Chornokur G et al	<i>Common Genetic Variation In Cellular Transport Genes and Epithelial Ovarian Cancer (EOC) Risk</i>	Click to view
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Crook A et al	<i>Connecting patients, researchers and clinical genetics services: the experiences of participants in the Australian Ovarian Cancer Study (AOCS)</i>	Click to view
Darabi H et al	<i>Polymorphisms in a Putative Enhancer at the 10q21.2 Breast Cancer Risk Locus Regulate NRBF2 Expression</i>	Click to view
Day FR et al	<i>Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair</i>	Click to view
Hernandez L et al	<i>A dual role for Caspase8 and NF-κB interactions in regulating apoptosis and necroptosis of ovarian cancer, with correlation to patient survival</i>	Click to view
Hunter SM et al	<i>Molecular profiling of low grade serous ovarian tumours identifies novel candidate driver genes</i>	Click to view
Jim HS et al	<i>Common Genetic Variation in Circadian Rhythm Genes and Risk of Epithelial Ovarian Cancer (EOC)</i>	Click to view
Johnatty SE et al	<i>Genome-wide Analysis Identifies Novel Loci Associated with Ovarian Cancer Outcomes: Findings from the Ovarian Cancer Association Consortium</i>	Click to view
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Keleman LE et al	<i>Genome-wide significant risk associations for mucinous ovarian carcinoma</i>	Click to view
Kondrashova O et al	<i>High-Throughput Amplicon-Based Copy Number Detection of 11 Genes in Formalin-Fixed Paraffin-Embedded Ovarian Tumour Samples by MLPA-Seq</i>	Click to view
Kuchenbaecker KB et al	<i>Identification of six new susceptibility loci for invasive epithelial ovarian cancer</i>	Click to view
Lawrenson K et al	<i>Cis-eQTL analysis and functional validation of candidate susceptibility genes for high-grade serous ovarian cancer</i>	Click to view
Lawrenson K et al	<i>Common variants at the CHEK2 gene locus and risk of epithelial ovarian cancer</i>	Click to view
Lee AW et al	<i>Evaluating the ovarian cancer gonadotropin hypothesis: a candidate gene study</i>	Click to view
Leong HS et al	<i>Efficient molecular subtype classification of high-grade serous ovarian cancer</i>	Click to view
Lin WY et al	<i>Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk</i>	Click to view
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Meeks HD	<i>BRCA2 Polymorphic Stop Codon K3326X and the Risk of Breast, Prostate, and Ovarian Cancers</i>	Click to view
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Nagle CM et al	<i>Aspirin, nonaspirin nonsteroidal anti-inflammatory drugs, acetaminophen and ovarian cancer survival</i>	Click to view
Nagle CM et al	<i>Obesity and survival among women with ovarian cancer: results from the Ovarian Cancer Association Consortium</i>	Click to view
OCAC et al	<i>No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer</i>	Click to view
O'Mara TA et al	<i>Comprehensive genetic assessment of the ESR1 locus identifies a risk region for endometrial cancer</i>	Click to view

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Orr N et al	<i>Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2</i>	Click to view
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Patch AM et al	<i>Whole-genome characterization of chemoresistant ovarian cancer</i>	Click to view
Patch AM et al	<i>Corrigendum: Whole-genome characterization of chemoresistant ovarian cancer</i>	Click to view
Ramus SJ et al	<i>Germline Mutations in the BRIP1, BARD1, PALB2, and NBN Genes in Women With Ovarian Cancer</i>	Click to view
Rudolph A et al	<i>Investigation of gene-environment interactions between 47 newly identified breast cancer susceptibility loci and environmental risk factors</i>	Click to view
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Ryland GL et al	<i>Mutational landscape of mucinous ovarian carcinoma and its neoplastic precursors</i>	Click to view
Sinha D et al	<i>Pericytes promote malignant ovarian cancer progression in mice and predict poor prognosis in serous ovarian cancer patients</i>	Click to view
Sobral-Leite M et al	<i>Annexin A1 expression in a pooled breast cancer series: association with tumor subtypes and prognosis</i>	Click to view
Song H et al	<i>Contribution of Germline Mutations in the RAD51B, RAD51C, and RAD51D Genes to Ovarian Cancer in the Population</i>	Click to view
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2014 Publications		
Author	Title of Paper	PubMed Link
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Au-Yeung G et al	<i>Impact of obesity on chemotherapy dosing for women with advanced stage serous ovarian cancer in the Australian Ovarian Cancer Study (AOCS)</i>	Click to view
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Charbonneau B et al	<i>Risk of ovarian cancer and the NF-κB pathway: genetic association with IL1A and TNFSF10</i>	Click to view
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Ghousaini M et al	<i>Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation</i>	Click to view
Hedditch EL et al	<i>ABCA transporter gene expression and poor outcome in epithelial ovarian cancer</i>	Click to view
Johnson N et al	<i>Genetic variation at CYP3A is associated with age at menarche and breast cancer risk: a case-control study</i>	Click to view
Kelemen LE et al	<i>Consortium analysis of gene and gene-folate interactions in purine and pyrimidine metabolism pathways with ovarian carcinoma risk</i>	Click to view

Author	Title of Paper	PubMed Link
Khan S et al	<i>MicroRNA related polymorphisms and breast cancer risk</i>	Click to view
Köbel M et al	<i>Evidence for a time-dependent association between FOLR1 expression and survival from ovarian carcinoma: implications for clinical testing. An Ovarian Tumour Tissue Analysis consortium study</i>	Click to view
Milne RL et al	<i>A large-scale assessment of two-way SNP interactions in breast cancer susceptibility using 46,450 cases and 42,461 controls from the breast cancer association consortium.</i>	Click to view
Milne RL et al	<i>Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium.</i>	Click to view
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Trabert B et al	<i>Aspirin, nonaspirin nonsteroidal anti-inflammatory drug, and acetaminophen use and risk of invasive epithelial ovarian cancer: a pooled analysis in the Ovarian Cancer Association Consortium</i>	Click to view

2013 Publications		
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Barnes DR et al	<i>Estimating single nucleotide polymorphism associations using pedigree data: applications to breast cancer.</i>	Click to view
Baumbusch LO et al	<i>High levels of genomic aberrations in serous ovarian cancers are associated with better survival</i>	Click to view
Beesley VL et al	<i>Changes in supportive care needs after first-line treatment for ovarian cancer: identifying care priorities and risk factors for future unmet needs.</i>	Click to view
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Dixon SC et al	<i>Dietary folate and related micronutrients, folate-metabolising genes, and ovarian cancer survival</i>	Click to view
Earp MA et al	<i>Genome-wide association study of subtype-specific epithelial ovarian cancer risk alleles using pooled DNA</i>	Click to view
Etemadmoghadam D et al	<i>Synthetic lethality between CCNE1 amplification and loss of BRCA1</i>	Click to view
Faber MT et al	<i>Cigarette smoking and risk of ovarian cancer: a pooled analysis of 21 case-control studies</i>	Click to view
George J et al	<i>Non-equivalent gene expression and copy number alterations in high-grade serous ovarian cancers with BRCA1 and BRCA2 mutations</i>	Click to view
Hallowell N et al	<i>The responses of research participants and their next of kin to receiving feedback of genetic test results following participation in the Australian Ovarian Cancer Study</i>	Click to view
Johnatty SE et al	<i>ABCB1 (MDR1) polymorphisms and ovarian cancer progression and survival: a comprehensive analysis from the Ovarian Cancer Association Consortium and The Cancer Genome Atlas</i>	Click to view
Jordan S et al	<i>Patterns of chemotherapy treatment for women with invasive epithelial ovarian cancer--a population-based study</i>	Click to view
Kelemen LE et al	<i>Recent alcohol consumption and risk of incident ovarian carcinoma: a pooled analysis of 5,342 cases and 10,358 controls from the Ovarian Cancer Association Consortium</i>	Click to view
Kondalsamy-Chennakesavan S et al	<i>Differentiating stage 1 epithelial ovarian cancer from benign ovarian tumours using a combination of tumour markers HE4, CA125, and CEA and patient's age</i>	Click to view
Meyer KB et al	<i>Fine-scale mapping of the FGFR2 breast cancer risk locus: putative functional variants differentially bind FOXA1 and E2F1</i>	Click to view

Author	Title of Paper	PubMed Link
Michailidou K et al	<i>Large-scale genotyping identifies 41 new loci associated with breast cancer risk</i>	Click to view
Mikeska T et al	<i>No evidence for PALB2 methylation in high-grade serous ovarian cancer</i>	Click to view
Nagle CM et al	<i>Glycemic index, glycemic load and endometrial cancer risk: results from the Australian National Endometrial Cancer study and an updated systematic review and meta-analysis.</i>	Click to view
Nickels S et al	<i>Evidence of gene-environment interactions between common breast cancer susceptibility loci and established environmental risk factors</i>	Click to view
Olsen CM et al	<i>Obesity and risk of ovarian cancer subtypes: evidence from the Ovarian Cancer Association Consortium</i>	Click to view
Pearce CL et al	<i>Combined and interactive effects of environmental and GWAS-identified risk factors in ovarian cancer. Cancer epidemiology, biomarkers & prevention</i>	Click to view
Permeth-Wey J et al	<i>Identification and molecular characterization of a new ovarian cancer susceptibility locus at 17q21.31</i>	Click to view
Pharoah PD et al	<i>GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer</i>	Click to view
Poole EM et al	<i>Hormonal and reproductive risk factors for epithelial ovarian cancer by tumor aggressiveness</i>	Click to view
Price MA et al	<i>Physical symptoms, coping styles and quality of life in recurrent ovarian cancer: a prospective population-based study over the last year of life</i>	Click to view
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Ryland GL et al	<i>RNF43 is a tumour suppressor gene mutated in mucinous tumours of the ovary</i>	Click to view
Shen H et al	<i>Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer</i>	Click to view
Sieh W et al	<i>Tubal ligation and risk of ovarian cancer subtypes: a pooled analysis of case-control studies</i>	Click to view
Terry KL et al	<i>Genital powder use and risk of ovarian cancer: a pooled analysis of 8,525 cases and 9,859 controls</i>	Click to view
White KL et al	<i>Analysis of over 10,000 Cases finds no association between previously reported candidate polymorphisms and ovarian cancer outcome</i>	Click to view

2012 Publications		
Author	Title of Paper	PubMed Link
Alsop K et al	<i>BRCA mutation frequency and patterns of treatment response in BRCA mutation-positive women with ovarian cancer.</i>	Click to view
Beesley VL et al	<i>Changes in supportive care needs after first-line treatment for ovarian cancer: identifying care priorities and risk factors for future unmet needs.</i>	Click to view
Colagiuri B et al	<i>A comparison of the FACT-G and the Supportive Care Needs Survey (SCNS) in women with ovarian cancer: unidimensionality of constructs.</i>	Click to view
Cowin PA et al	<i>LRP1B deletion in high-grade serous ovarian cancers is associated with acquired chemotherapy resistance to liposomal doxorubicin.</i>	Click to view
Ghousaini M et al	<i>Genome-wide association analysis identifies three new breast cancer susceptibility loci.</i>	Click to view
Hein R, et al.	<i>Comparison of 6q25 Breast Cancer Hits from Asian and European Genome Wide Association Studies in the Breast Cancer Association Consortium (BCAC).</i>	Click to view
Hunter SM et al	<i>Pre-invasive ovarian mucinous tumors are characterized by CDKN2A and RAS pathway aberrations.</i>	Click to view
Ibiebele TI et al	<i>Intake of omega-3 and omega-6 fatty acids and risk of ovarian cancer.</i>	Click to view
Kirchoff T et al	<i>Breast cancer risk and 6q22.33: combined results from Breast Cancer Association Consortium and Consortium of Investigators on Modifiers of BRCA1/2</i>	Click to view
Kulbe H et al	<i>A dynamic inflammatory cytokine network in the human ovarian cancer microenvironment.</i>	Click to view

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Lahmann PH et al	<i>Estimated intake of dietary phytoestrogens in Australian women and evaluation of correlates of phyto-oestrogen intake.</i>	Click to view
Lambrechts D et al	<i>11q13 is a Susceptibility Locus for Hormone Receptor Positive Breast Cancer.</i>	Click to view
Lu Y et al	<i>Genome-wide association study for ovarian cancer susceptibility using pooled DNA.</i>	Click to view
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