

Jolanta Kupryjanczyk

List of Publications by Citations

Source: <https://exaly.com/author-pdf/392458/jolanta-kupryjanczyk-publications-by-citations.pdf>
Version: 2024-04-17

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.
The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

19 papers	1,612 citations	15 h-index	22 g-index
22 ext. papers	1,924 ext. citations	11.1 avg, IF	2.66 L-index

#	Paper	IF	Citations
19	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , 2013 , 45, 371-84, 384e1-2	36.3	422
18	GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. <i>Nature Genetics</i> , 2013 , 45, 362-70, 370e1-2	36.3	267
17	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017 , 49, 680-691	36.3	190
16	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015 , 47, 164-71	36.3	177
15	Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. <i>Nature Communications</i> , 2013 , 4, 1628	17.4	124
14	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>Cancer Discovery</i> , 2016 , 6, 1052-67	14.4	104
13	A novel germline PALB2 deletion in Polish breast and ovarian cancer patients. <i>BMC Medical Genetics</i> , 2010 , 11, 20	2.1	86
12	Gene expression analysis in ovarian cancer - faults and hints from DNA microarray study. <i>Frontiers in Oncology</i> , 2014 , 4, 6	5.3	55
11	The putative oncogene, CRNDE, is a negative prognostic factor in ovarian cancer patients. <i>Oncotarget</i> , 2015 , 6, 43897-910	3.3	46
10	Unsupervised analysis reveals two molecular subgroups of serous ovarian cancer with distinct gene expression profiles and survival. <i>Journal of Cancer Research and Clinical Oncology</i> , 2016 , 142, 1239-52	4.9	20
9	High frequency of allelic loss at the BRCA1 locus in ovarian cancers: clinicopathologic and molecular associations. <i>Cancer Genetics</i> , 2012 , 205, 94-100	2.3	18
8	TP53, BCL-2 and BAX analysis in 199 ovarian cancer patients treated with taxane-platinum regimens. <i>Gynecologic Oncology</i> , 2009 , 112, 179-84	4.9	18
7	TP53 status and taxane-platinum versus platinum-based therapy in ovarian cancer patients: a non-randomized retrospective study. <i>BMC Cancer</i> , 2008 , 8, 27	4.8	18
6	Fibronectin and Periostin as Prognostic Markers in Ovarian Cancer. <i>Cells</i> , 2020 , 9,	7.9	17
5	Nuclear survivin expression is a positive prognostic factor in taxane-platinum-treated ovarian cancer patients. <i>Journal of Ovarian Research</i> , 2011 , 4, 20	5.5	17
4	Clinical importance of FANCD2, BRIP1, BRCA1, BRCA2 and FANCF expression in ovarian carcinomas. <i>Cancer Biology and Therapy</i> , 2019 , 20, 843-854	4.6	9
3	Prognosis of patients with BRCA1-associated ovarian carcinomas depends on TP53 accumulation status in tumor cells. <i>Gynecologic Oncology</i> , 2017 , 144, 369-376	4.9	7

- 2 Cross-Cancer Genome-Wide Association Study of Endometrial Cancer and Epithelial Ovarian Cancer Identifies Genetic Risk Regions Associated with Risk of Both Cancers. *Cancer Epidemiology Biomarkers and Prevention*, **2021**, 30, 217-228 4 7
- 1 Menopausal hormone therapy prior to the diagnosis of ovarian cancer is associated with improved survival. *Gynecologic Oncology*, **2020**, 158, 702-709 4.9 5