Joanna Moes-Sosnowska

List of Publications by Year
in descending order

19 papers	1,021 citations	933447 10 h-index	996975 15 g-index
19 all docs	19 docs citations	19 times ranked	3285 citing authors

#	Article	IF	Citations
1	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. Nature Genetics, 2017, 49, 680-691.	21.4	356
2	PALB2,CHEK2andATMrare variants and cancer risk: data from COGS. Journal of Medical Genetics, 2016, 53, 800-811.	3.2	174
3	Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. Nature Communications, 2013, 4, 1628.	12.8	144
4	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
5	Genome-wide significant risk associations for mucinous ovarian carcinoma. Nature Genetics, 2015, 47, 888-897.	21.4	78
6	BRCA2 Polymorphic Stop Codon K3326X and the Risk of Breast, Prostate, and Ovarian Cancers. Journal of the National Cancer Institute, 2016, 108, djv315.	6.3	77
7	Germline SMARCA4 mutations in patients with ovarian small cell carcinoma of hypercalcemic type. Orphanet Journal of Rare Diseases, 2015, 10, 32.	2.7	31
8	Common variants at theCHEK2gene locus and risk of epithelial ovarian cancer. Carcinogenesis, 2015, 36, 1341-1353.	2.8	24
9	Clinical importance of FANCD2, BRIP1, BRCA1, BRCA2 and FANCF expression in ovarian carcinomas. Cancer Biology and Therapy, 2019, 20, 843-854.	3.4	20
10	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
11	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
12	Detection of EGFR mutations in liquid biopsy samples using allele-specific quantitative PCR: A comparative real-world evaluation of two popular diagnostic systems. Advances in Medical Sciences, 2021, 66, 336-342.	2.1	4
13	Clinical importance of the EMSY gene expression and polymorphisms in ovarian cancer. Oncotarget, 2018, 9, 17735-17755.	1.8	4
14	1409P Comparing different methods of FGFR1 aberrations analysis in squamous cell lung cancer (SqCLC) targeted therapy. Annals of Oncology, 2020, 31, S891.	1.2	1
15	Analiza mutacji i polimorfizmÃ³w w genie FANCD2 u pacjentek chorych na raka jajnika z populacji polskiej. Nowotwory, 2015, 65, 7-13.	0.3	1
16	The EGFR mutation detection in NSCLC by Next Generation Sequencing (NGS): cons and pros. , 2019, , .		1
17	FGFR1-4 and MET expression analysis and evaluation of reliable reference genes in Sq-NSCLC. , 2019, , .		1
18	FGFR1 gene aberrations and FGFR1 protein expression in squamous non-small cell lung cancer (Sq-NSCLC). , 2021, , .		0

#	Article	IF	Citations
19	Molecular profiling of Sq-NSCLC with enhanced FGFR1-4 and MET gene expression â€“ NGS pilot study. , 2020, , .		0