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	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
2	FGFR1 gene aberrations and FGFR1 protein expression in squamous non-small cell lung cancer (Sq-NSCLC). , 2021, , .		0
3	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
4	Molecular profiling of Sq-NSCLC with enhanced FGFR1-4 and MET gene expression â€“ NGS pilot study. , 2020, , .		0
5	Clinical importance of FANCD2, BRIP1, BRCA1, BRCA2 and FANCF expression in ovarian carcinomas. Cancer Biology and Therapy, 2019, 20, 843-854.	3.4	20
6	The EGFR mutation detection in NSCLC by Next Generation Sequencing (NGS): cons and pros. , 2019, , .		1
7	FGFR1-4 and MET expression analysis and evaluation of reliable reference genes in Sq-NSCLC. , 2019, , .		1
8	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
9	Clinical importance of the EMSY gene expression and polymorphisms in ovarian cancer. Oncotarget, 2018, 9, 17735-17755.	1.8	4
10	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. Nature Genetics, 2017, 49, 680-691.	21.4	356
11	PALB2,CHEK2andATMrare variants and cancer risk: data from COGS. Journal of Medical Genetics, 2016, 53, 800-811.	3.2	174
12	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
13	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
14	Genome-wide significant risk associations for mucinous ovarian carcinoma. Nature Genetics, 2015, 47, 888-897.	21.4	78
15	Germline SMARCA4 mutations in patients with ovarian small cell carcinoma of hypercalcemic type. Orphanet Journal of Rare Diseases, 2015, 10, 32.	2.7	31
16	Common variants at theCHEK2gene locus and risk of epithelial ovarian cancer. Carcinogenesis, 2015, 36, 1341-1353.	2.8	24
17	Analiza mutacji i polimorfizmÃ³w w genie FANCD2 u pacjentek chorych na raka jajnika z populacji polskiej. Nowotwory, 2015, 65, 7-13.	0.3	1
18	Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. Nature Communications, 2013, 4, 1628.	12.8	144

#	Article	IF	Citations
19	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