

Boris J Winterhoff

List of Publications by Year in descending order

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Version: 2024-02-01

14
papers

374
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1040056

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996975

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docs citations

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times ranked

804
citing authors

#	ARTICLE	IF	CITATIONS
1	Single cell sequencing reveals heterogeneity within ovarian cancer epithelium and cancer associated stromal cells. <i>Gynecologic Oncology</i> , 2017, 144, 598-606.	1.4	82
2	Molecular classification of high grade endometrioid and clear cell ovarian cancer using TCGA gene expression signatures. <i>Gynecologic Oncology</i> , 2016, 141, 95-100.	1.4	58
3	Development and Validation of the Gene Expression Predictor of High-grade Serous Ovarian Carcinoma Molecular SubTYPE (PrOTYPE). <i>Clinical Cancer Research</i> , 2020, 26, 5411-5423.	7.0	43
4	Gene-expression signatures in ovarian cancer: Promise and challenges for patient stratification. <i>Gynecologic Oncology</i> , 2016, 141, 379-385.	1.4	37
5	USP14 is a predictor of recurrence in endometrial cancer and a molecular target for endometrial cancer treatment. <i>Oncotarget</i> , 2016, 7, 30962-30976.	1.8	35
6	Targeting fibroblast growth factor pathways in endometrial cancer. <i>Current Problems in Cancer</i> , 2017, 41, 37-47.	2.0	30
7	A patient-centered mobile health application to motivate use of genetic counseling among women with ovarian cancer: A pilot randomized controlled trial. <i>Gynecologic Oncology</i> , 2019, 153, 100-107.	1.4	17
8	The transcription factor PAX8 promotes angiogenesis in ovarian cancer through interaction with SOX17. <i>Science Signaling</i> , 2022, 15, eabm2496.	3.6	15
9	UNC-45A is preferentially expressed in epithelial cells and binds to and co-localizes with interphase MTs. <i>Cancer Biology and Therapy</i> , 2019, 20, 1304-1313.	3.4	14
10	Dilution of Molecularâ€™Pathologic Gene Signatures by Medically Associated Factors Might Prevent Prediction of Resection Status After Debulking Surgery in Patients With Advanced Ovarian Cancer. <i>Clinical Cancer Research</i> , 2020, 26, 213-219.	7.0	12
11	Expression signature distinguishing two tumour transcriptome classes associated with progression-free survival among rare histological types of epithelial ovarian cancer. <i>British Journal of Cancer</i> , 2016, 114, 1412-1420.	6.4	8
12	De novo prediction of cell-type complexity in single-cell RNA-seq and tumor microenvironments. <i>Life Science Alliance</i> , 2019, 2, e201900443.	2.8	8
13	Assessment of the Intraoperative Consultation Service Rendered by General Pathologists in a Scenario Where a Well-Defined Decision Algorithm Is Followed. <i>American Journal of Clinical Pathology</i> , 2017, 147, 322-326.	0.7	6
14	Developing a Clinico-Molecular Test for Individualized Treatment of Ovarian Cancer: The interplay of Precision Medicine Informatics with Clinical and Health Economics Dimensions. <i>AMIA ... Annual Symposium proceedings</i> , 2018, 2018, 1093-1102.	0.2	1