

Kalliopi I Pappa

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

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citations

1163117

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412
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#	ARTICLE	IF	CITATIONS
1	p16/Ki-67 Dual Staining Is a Reliable Biomarker for Risk Stratification for Patients With Borderline/Mild Cytology in Cervical Cancer Screening. <i>Anticancer Research</i> , 2022, 42, 2599-2606.	1.1	3
2	Mimiviruses: Giant viruses with novel and intriguing features (Review). <i>Molecular Medicine Reports</i> , 2022, 25, .	2.4	1
3	The p16/ki-67 assay is a safe, effective and rapid approach to triage women with mild cervical lesions. <i>PLoS ONE</i> , 2021, 16, e0253045.	2.5	8
4	Effects of Hormone Therapy and Flavonoids Capable on Reversal of Menopausal Immune Senescence. <i>Nutrients</i> , 2021, 13, 2363.	4.1	5
5	Metabolic rewiring is associated with HPV-specific profiles in cervical cancer cell lines. <i>Scientific Reports</i> , 2021, 11, 17718.	3.3	9
6	Treatment of Cervical Pregnancy with Ultrasound-Guided Intra-gestational Injection of Methotrexate: A Case Report. <i>Case Reports in Obstetrics and Gynecology</i> , 2021, 2021, 1-5.	0.3	0
7	Membrane proteomics of cervical cancer cell lines reveal insights on the process of cervical carcinogenesis. <i>International Journal of Oncology</i> , 2018, 53, 2111-2122.	3.3	6
8	Proteomic Analysis of Normal and Cancer Cervical Cell Lines Reveals Deregulation of Cytoskeleton-associated Proteins. <i>Cancer Genomics and Proteomics</i> , 2017, 14, 253-266.	2.0	30
9	Cervical Cancer Cell Line Secretome Highlights the Roles of Transforming Growth Factor-Beta-Induced Protein ig-h3, Peroxiredoxin-2, and NRF2 on Cervical Carcinogenesis. <i>BioMed Research International</i> , 2017, 2017, 1-15.	1.9	39
10	High Resolution Proteomic Analysis of the Cervical Cancer Cell Lines Secretome Documents Deregulation of Multiple Proteases. <i>Cancer Genomics and Proteomics</i> , 2017, 14, 507-521.	2.0	17
11	Proteomics approaches in cervical cancer: focus on the discovery of biomarkers for diagnosis and drug treatment monitoring. <i>Expert Review of Proteomics</i> , 2016, 13, 731-745.	3.0	27
12	Profiling of Discrete Gynecological Cancers Reveals Novel Transcriptional Modules and Common Features Shared by Other Cancer Types and Embryonic Stem Cells. <i>PLoS ONE</i> , 2015, 10, e0142229.	2.5	52
13	Low Mutational Burden of Eight Genes Involved in the MAPK/ERK, PI3K/AKT, and GNAQ/11 Pathways in Female Genital Tract Primary Melanomas. <i>BioMed Research International</i> , 2015, 2015, 1-10.	1.9	16
14	Comparative Assessment of Lymph Node Micrometastasis in Cervical, Endometrial and Vulvar Cancer: Insights on the Real Time qRT-PCR Approach versus Immunohistochemistry, Employing Dual Molecular Markers. <i>BioMed Research International</i> , 2014, 2014, 1-9.	1.9	5
15	Expression Profiling of Vulvar Carcinoma: Clues for Deranged Extracellular Matrix Remodeling and Effects on Multiple Signaling Pathways Combined with Discrete Patient Subsets. <i>Translational Oncology</i> , 2011, 4, 301-IN6.	3.7	9