Elena A Alexandrova

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5372903/publications.pdf

Version: 2024-02-01

22 papers

764

16 h-index 22 g-index

22 all docs 22 docs citations

times ranked

22

1093 citing authors

#	Article	IF	CITATIONS
1	Nasopharyngeal virome analysis of COVIDâ€19 patients during three different waves in Campania region of Italy. Journal of Medical Virology, 2022, , .	2.5	9
2	Rapid and sensitive detection of SARS-CoV-2 variants in nasopharyngeal swabs and wastewaters. Diagnostic Microbiology and Infectious Disease, 2022, 102, 115632.	0.8	6
3	NGS analysis of nasopharyngeal microbiota in SARS-CoV-2 positive patients during the first year of the pandemic in the Campania Region of Italy. Microbial Pathogenesis, 2022, 165, 105506.	1.3	12
4	Histone Methyltransferase DOT1L as a Promising Epigenetic Target for Treatment of Solid Tumors. Frontiers in Genetics, 2022, 13, 864612.	1.1	22
5	Interaction Proteomics Identifies ERbeta Association with Chromatin Repressive Complexes to Inhibit Cholesterol Biosynthesis and Exert An Oncosuppressive Role in Triple-negative Breast Cancer. Molecular and Cellular Proteomics, 2020, 19, 245-260.	2.5	18
6	An Overview of Candidate Therapeutic Target Genes in Ovarian Cancer. Cancers, 2020, 12, 1470.	1.7	20
7	Insights into the Role of Estrogen Receptor \hat{I}^2 in Triple-Negative Breast Cancer. Cancers, 2020, 12, 1477.	1.7	33
8	Small Non-Coding RNA Profiling Identifies miR-181a-5p as a Mediator of Estrogen Receptor Beta-Induced Inhibition of Cholesterol Biosynthesis in Triple-Negative Breast Cancer. Cells, 2020, 9, 874.	1.8	25
9	Molecular and Functional Characterization of the Somatic PIWIL1/piRNA Pathway in Colorectal Cancer Cells. Cells, 2019, 8, 1390.	1.8	16
10	The Histone Methyltransferase DOT1L Is a Functional Component of Estrogen Receptor Alpha Signaling in Ovarian Cancer Cells. Cancers, 2019, 11, 1720.	1.7	24
11	Inhibition of histone methyltransferase DOT1L silences $\mathrm{ER}\hat{\mathrm{l}}\pm$ gene and blocks proliferation of antiestrogen-resistant breast cancer cells. Science Advances, 2019, 5, eaav5590.	4.7	70
12	Quantitative mapping of RNA-mediated nuclear estrogen receptor \hat{l}^2 interactome in human breast cancer cells. Scientific Data, 2018, 5, 180031.	2.4	22
13	Large-scale profiling of signalling pathways reveals an asthma specific signature in bronchial smooth muscle cells. Oncotarget, 2016, 7, 25150-25161.	0.8	32
14	Small RNA profiling reveals deregulated phosphatase and tensin homolog (PTEN)/phosphoinositide 3-kinase (PI3K)/Akt pathway in bronchial smooth muscle cells from asthmatic patients. Journal of Allergy and Clinical Immunology, 2016, 137, 58-67.	1.5	30
15	Small non-coding RNA deregulation in endometrial carcinogenesis. Oncotarget, 2015, 6, 4677-4691.	0.8	49
16	Identification of cytoplasmic proteins interacting with unliganded estrogen receptor \hat{l}^{\pm} and \hat{l}^{2} in human breast cancer cells. Proteomics, 2015, 15, 1801-1807.	1.3	17
17	Sense transcripts originated from an internal part of the human retrotransposon LINE-1 5′ UTR. Gene, 2012, 511, 46-53.	1.0	21
18	Functional human endogenous retroviral LTR transcription start sites are located between the R and U5 regions. Virology, 2006, 346, 373-378.	1.1	51

#	Article	IF	CITATIONS
19	GREM, a technique for genome-wide isolation and quantitative analysis of promoter active repeats. Nucleic Acids Research, 2006, 34, e67-e67.	6.5	36
20	At Least 50% of Human-Specific HERV-K (HML-2) Long Terminal Repeats Serve In Vivo as Active Promoters for Host Nonrepetitive DNA Transcription. Journal of Virology, 2006, 80, 10752-10762.	1.5	108
21	Improving specificity of DNA hybridization-based methods. Nucleic Acids Research, 2004, 32, e130-e130.	6.5	50
22	The human genome contains many types of chimeric retrogenes generated through in vivo RNA recombination. Nucleic Acids Research, 2003, 31, 4385-4390.	6.5	93