

Francesca Guerrieri

List of Publications by Year in descending order

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46
papers

2,443
citations

331670

21
h-index

377865

34
g-index

47
all docs

47
docs citations

47
times ranked

3888
citing authors

#	ARTICLE	IF	CITATIONS
1	Host Epigenetic Alterations and Hepatitis B Virus-Associated Hepatocellular Carcinoma. <i>Journal of Clinical Medicine</i> , 2021, 10, 1715.	2.4	12
2	The lncRNAs in HBV-Related HCCs: Targeting Chromatin Dynamics and Beyond. <i>Cancers</i> , 2021, 13, 3115.	3.7	6
3	Antibiotics Treatment Modulates Microglia-Synapses Interaction. <i>Cells</i> , 2021, 10, 2648.	4.1	17
4	Characterization of black patina from the Tiber River embankments using Next-Generation Sequencing. <i>PLoS ONE</i> , 2020, 15, e0227639.	2.5	16
5	Microbiota in Waterlogged Archaeological Wood: Use of Next-Generation Sequencing to Evaluate the Risk of Biodegradation. <i>Applied Sciences (Switzerland)</i> , 2020, 10, 4636.	2.5	12
6	Structural Variations of Vaginal and Endometrial Microbiota: Hints on Female Infertility. <i>Frontiers in Cellular and Infection Microbiology</i> , 2020, 10, 350.	3.9	67
7	Essential Oils as Alternative Biocides for the Preservation of Waterlogged Archaeological Wood. <i>Microorganisms</i> , 2020, 8, 2015.	3.6	18
8	Blockade of EIF5A hypusination limits colorectal cancer growth by inhibiting MYC elongation. <i>Cell Death and Disease</i> , 2020, 11, 1045.	6.3	39
9	Nasal Microbiota in RSV Bronchiolitis. <i>Microorganisms</i> , 2020, 8, 731.	3.6	19
10	Hepatitis Delta Virus histone mimicry drives the recruitment of chromatin remodelers for viral RNA replication. <i>Nature Communications</i> , 2020, 11, 419.	12.8	19
11	Hepatitis B protein HBx binds the DLEU2 lncRNA to sustain cccDNA and host cancer-related gene transcription. <i>Gut</i> , 2020, 69, 2016-2024.	12.1	92
12	Targeting p53 and histone methyltransferases restores exhausted CD8+ T cells in HCV infection. <i>Nature Communications</i> , 2020, 11, 604.	12.8	44
13	Characterization of black patina from the Tiber River embankments using Next-Generation Sequencing. , 2020, 15, e0227639.		0
14	Characterization of black patina from the Tiber River embankments using Next-Generation Sequencing. , 2020, 15, e0227639.		0
15	Characterization of black patina from the Tiber River embankments using Next-Generation Sequencing. , 2020, 15, e0227639.		0
16	Characterization of black patina from the Tiber River embankments using Next-Generation Sequencing. , 2020, 15, e0227639.		0
17	Characterization of black patina from the Tiber River embankments using Next-Generation Sequencing. , 2020, 15, e0227639.		0
18	Characterization of black patina from the Tiber River embankments using Next-Generation Sequencing. , 2020, 15, e0227639.		0

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19	SAT-165-Identification of chromatin-accessible domains on the host genome and hepatitis B virus mini-chromosome in infected primary human hepatocytes. <i>Journal of Hepatology</i> , 2019, 70, e702.	3.7	1
20	Microglia-Derived Microvesicles Affect Microglia Phenotype in Glioma. <i>Frontiers in Cellular Neuroscience</i> , 2019, 13, 41.	3.7	52
21	The Food Additive Maltodextrin Promotes Endoplasmic Reticulum Stress-Driven Mucus Depletion and Exacerbates Intestinal Inflammation. <i>Cellular and Molecular Gastroenterology and Hepatology</i> , 2019, 7, 457-473.	4.5	84
22	Enteric Delivery of Regenerating Family Member 3 alpha Alters the Intestinal Microbiota and Controls Inflammation in Mice With Colitis. <i>Gastroenterology</i> , 2018, 154, 1009-1023.e14.	1.3	107
23	Targeting a phospho-STAT3-miRNAs pathway improves vesicular hepatic steatosis in an in vitro and in vivo model. <i>Scientific Reports</i> , 2018, 8, 13638.	3.3	14
24	Combining amplicon sequencing and metabolomics in cirrhotic patients highlights distinctive microbiota features involved in bacterial translocation, systemic inflammation and hepatic encephalopathy. <i>Scientific Reports</i> , 2018, 8, 8210.	3.3	63
25	Genome-wide identification of direct HBx genomic targets. <i>BMC Genomics</i> , 2017, 18, 184.	2.8	52
26	Targeting mitochondrial dysfunction can restore antiviral activity of exhausted HBV-specific CD8 T cells in chronic hepatitis B. <i>Nature Medicine</i> , 2017, 23, 327-336.	30.7	251
27	Mo1927 Mucosa-Associated Microbiota and Promoter Methylation Status of Genes Involved in Immune Response in Crohn's Disease Patients. <i>Gastroenterology</i> , 2016, 150, S818.	1.3	0
28	Pathobiology of Hepatitis B Virus-Induced Carcinogenesis. <i>Molecular and Translational Medicine</i> , 2016, , 95-121.	0.4	2
29	IL6 Inhibits HBV Transcription by Targeting the Epigenetic Control of the Nuclear cccDNA Minichromosome. <i>PLoS ONE</i> , 2015, 10, e0142599.	2.5	66
30	Clinical relevance of next generation sequencing on baseline detection of minority resistance associated variants in HCV-1 patients treated with protease inhibitors. <i>Digestive and Liver Disease</i> , 2015, 47, e48.	0.9	0
31	HBx-DLEU2 lncRNA complex affects transcription of new target promoters. <i>Digestive and Liver Disease</i> , 2015, 47, e30.	0.9	0
32	Long non-coding RNAs era in liver cancer. <i>World Journal of Hepatology</i> , 2015, 7, 1971.	2.0	32
33	p53 and TP73 in cancer, an unresolved "family" puzzle of complexity, redundancy and hierarchy. <i>FEBS Letters</i> , 2014, 588, 2590-2599.	2.8	39
34	The HMGA1 protooncogene frequently deregulated in cancer is a transcriptional target of E2F1. <i>Molecular Carcinogenesis</i> , 2013, 52, 526-534.	2.7	22
35	Molecular Mechanisms of HBV-Associated Hepatocarcinogenesis. <i>Seminars in Liver Disease</i> , 2013, 33, 147-156.	3.6	96
36	The sodium/iodide symporter NIS is a transcriptional target of the p53-family members in liver cancer cells. <i>Cell Death and Disease</i> , 2013, 4, e807-e807.	6.3	18

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37	Transcriptional regulation of miR-224 upregulated in human HCCs by NF κ B inflammatory pathways. <i>Journal of Hepatology</i> , 2012, 56, 855-861.	3.7	134
38	IFN α inhibits HBV transcription and replication in cell culture and in humanized mice by targeting the epigenetic regulation of the nuclear cccDNA minichromosome. <i>Journal of Clinical Investigation</i> , 2012, 122, 529-537.	8.2	492
39	p53-paralog DNp73 oncogene is repressed by IFN α /STAT2 through the recruitment of the Ezh2 polycomb group transcriptional repressor. <i>Oncogene</i> , 2011, 30, 2670-2678.	5.9	26
40	Chromatin Dynamics of Gene Activation and Repression in Response to Interferon α (IFN α) Reveal New Roles for Phosphorylated and Unphosphorylated Forms of the Transcription Factor STAT2. <i>Journal of Biological Chemistry</i> , 2011, 286, 20217-20227.	3.4	51
41	hSirT1-Dependent Regulation of the PCAF-E2F1-p73 Apoptotic Pathway in Response to DNA Damage. <i>Molecular and Cellular Biology</i> , 2009, 29, 1989-1998.	2.3	66
42	Nuclear HBx binds the HBV minichromosome and modifies the epigenetic regulation of cccDNA function. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 19975-19979.	7.1	403
43	Differential regulation of sodium/iodide symporter NIS gene promoter by p53, p63 and p73 in human cholangiocarcinoma (CCA) and hepatocellular (HCC) cell lines. <i>Digestive and Liver Disease</i> , 2009, 41, A18-A19.	0.9	0
44	<i>Molecular Pathogenesis.</i> , 2009, , 9-25.		1
45	[71] THE SIRT1 HYSTONE DEACETYLASE REGULATES TAP TRANSCRIPTION AND APOPTOSIS IN RESPONSE TO DNA DAMAGE. <i>Journal of Hepatology</i> , 2007, 46, S32.	3.7	0
46	Gut Microbiota Structure and Metabolites, Before and After Treatment in Early Rheumatoid Arthritis Patients: A Pilot Study. <i>Frontiers in Medicine</i> , 0, 9, .	2.6	7