

Silvano Piazza

List of Publications by Year in descending order

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Version: 2024-02-01

47
papers

5,256
citations

218677

26
h-index

276875

41
g-index

48
all docs

48
docs citations

48
times ranked

11643
citing authors

#	ARTICLE	IF	CITATIONS
1	Immune dysfunction in the cerebellum of mice lacking the autism candidate gene Engrailed 2. <i>Journal of Neuroimmunology</i> , 2022, 367, 577870.	2.3	3
2	HMGA1 positively regulates the microtubule-destabilizing protein stathmin promoting motility in TNBC cells and decreasing tumour sensitivity to paclitaxel. <i>Cell Death and Disease</i> , 2022, 13, 429.	6.3	2
3	Hyperinsulinemia and insulin resistance in the obese may develop as part of a homeostatic response to elevated free fatty acids: A mechanistic case-control and a population-based cohort study. <i>EBioMedicine</i> , 2021, 65, 103264.	6.1	51
4	A selective p53 activator and anticancer agent to improve colorectal cancer therapy. <i>Cell Reports</i> , 2021, 35, 108982.	6.4	20
5	Integrative microRNAome analysis of skeletal muscle of <i>Colossoma macropomum</i> (tambaqui), <i>Piaractus mesopotamicus</i> (pacu), and the hybrid tambacu, based on next-generation sequencing data. <i>BMC Genomics</i> , 2021, 22, 237.	2.8	3
6	B-cell receptor signaling and genetic lesions in TP53 and CDKN2A/CDKN2B cooperate in Richter transformation. <i>Blood</i> , 2021, 138, 1053-1066.	1.4	33
7	Characterization of black patina from the Tiber River embankments using Next-Generation Sequencing. <i>PLoS ONE</i> , 2020, 15, e0227639.	2.5	16
8	Architecture of The Human Ape1 Interactome Defines Novel Cancers Signatures. <i>Scientific Reports</i> , 2020, 10, 28.	3.3	22
9	The altered transcriptome of pediatric myelodysplastic syndrome revealed by RNA sequencing. <i>Journal of Hematology and Oncology</i> , 2020, 13, 135.	17.0	4
10	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
11	Fasting-mimicking diet and hormone therapy induce breast cancer regression. <i>Nature</i> , 2020, 583, 620-624.	27.8	198
12	Mutant p53 induces Golgi tubulo-vesiculation driving a prometastatic secretome. <i>Nature Communications</i> , 2020, 11, 3945.	12.8	52
13	Essential Oils as Alternative Biocides for the Preservation of Waterlogged Archaeological Wood. <i>Microorganisms</i> , 2020, 8, 2015.	3.6	18
14	Changes in the Expression of Pre-Replicative Complex Genes in hTERT and ALT Pediatric Brain Tumors. <i>Cancers</i> , 2020, 12, 1028.	3.7	8
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	Characterization of black patina from the Tiber River embankments using Next-Generation Sequencing. , 2020, 15, e0227639.		0
20	Characterization of black patina from the Tiber River embankments using Next-Generation Sequencing. , 2020, 15, e0227639.		0
21	SLMP53-2 Restores Wild-Type-Like Function to Mutant p53 through Hsp70: Promising Activity in Hepatocellular Carcinoma. <i>Cancers</i> , 2019, 11, 1151.	3.7	21
22	HMGA1 promotes breast cancer angiogenesis supporting the stability, nuclear localization and transcriptional activity of FOXM1. <i>Journal of Experimental and Clinical Cancer Research</i> , 2019, 38, 313.	8.6	67
23	Introduction to Bioinformatics. <i>Methods in Molecular Biology</i> , 2019, 1986, 1-15.	0.9	12
24	MiR-181 family-specific behavior in different cancers: a meta-analysis view. <i>Cancer and Metastasis Reviews</i> , 2018, 37, 17-32.	5.9	63
25	A covalent PIN1 inhibitor selectively targets cancer cells by a dual mechanism of action. <i>Nature Communications</i> , 2017, 8, 15772.	12.8	102
26	Mammalian APE1 controls miRNA processing and its interactome is linked to cancer RNA metabolism. <i>Nature Communications</i> , 2017, 8, 797.	12.8	107
27	HMGA1 regulates the Plasminogen activation system in the secretome of breast cancer cells. <i>Scientific Reports</i> , 2017, 7, 11768.	3.3	36
28	PIN1 in breast development and cancer: a clinical perspective. <i>Cell Death and Differentiation</i> , 2017, 24, 200-211.	11.2	51
29	CTSE1: a novel TEAD4-E2F1 target gene involved in cell protrusions formation in triple-negative breast cancer cell models. <i>Oncotarget</i> , 2017, 8, 67422-67438.	1.8	17
30	Epigenetic silencing of miR-296 and miR-512 ensures hTERT dependent apoptosis protection and telomere maintenance in basal-type breast cancer cells. <i>Oncotarget</i> , 2017, 8, 95674-95691.	1.8	33
31	<scp>YAP</scp> enhances the proâ€proliferative transcriptional activity of mutant p53 proteins. <i>EMBO Reports</i> , 2016, 17, 188-201.	4.5	154
32	Proteasome machinery is instrumental in a common gain-of-function program of the p53 missense mutants in cancer. <i>Nature Cell Biology</i> , 2016, 18, 897-909.	10.3	205
33	Translating Proteomic Into Functional Data: An High Mobility Group A1 (HMGA1) Proteomic Signature Has Prognostic Value in Breast Cancer. <i>Molecular and Cellular Proteomics</i> , 2016, 15, 109-123.	3.8	41
34	A novel HMGA1-CCNE2-YAP axis regulates breast cancer aggressiveness. <i>Oncotarget</i> , 2015, 6, 19087-19101.	1.8	70
35	Prolylâ€isomerase Pin1 controls normal and cancer stem cells of the breast. <i>EMBO Molecular Medicine</i> , 2014, 6, 99-119.	6.9	130
36	A promoter-level mammalian expression atlas. <i>Nature</i> , 2014, 507, 462-470.	27.8	1,838

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37	Metabolic control of YAP and TAZ by the mevalonate pathway. <i>Nature Cell Biology</i> , 2014, 16, 357-366.	10.3	630
38	Specific Mesothelial Signature Marks the Heterogeneity of Mesenchymal Stem Cells From High-Grade Serous Ovarian Cancer. <i>Stem Cells</i> , 2014, 32, 2998-3011.	3.2	16
39	miR-155 Drives Telomere Fragility in Human Breast Cancer by Targeting TRF1. <i>Cancer Research</i> , 2014, 74, 4145-4156.	0.9	108
40	Oncogenic miR-181a/b affect the DNA damage response in aggressive breast cancer. <i>Cell Cycle</i> , 2013, 12, 1679-1687.	2.6	109
41	HMGA1 promotes metastatic processes in basal-like breast cancer regulating EMT and stemness. <i>Oncotarget</i> , 2013, 4, 1293-1308.	1.8	145
42	GTSE1 Is a Microtubule Plus-End Tracking Protein That Regulates EB1-Dependent Cell Migration. <i>PLoS ONE</i> , 2012, 7, e51259.	2.5	52
43	A Pin1/Mutant p53 Axis Promotes Aggressiveness in Breast Cancer. <i>Cancer Cell</i> , 2011, 20, 79-91.	16.8	256
44	Wiring the oncogenic circuitry: Pin1 unleashes mutant p53. <i>Oncotarget</i> , 2011, 2, 654-656.	1.8	22
45	Multipotent Progenitor Cells Are Present in Human Peripheral Blood. <i>Circulation Research</i> , 2009, 104, 1225-1234.	4.5	126
46	Multipotent cells can be generated in vitro from several adult human organs (heart, liver, and bone) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	1.4	336
47	The Transcriptional Repressor hDaxx Potentiates p53-dependent Apoptosis. <i>Journal of Biological Chemistry</i> , 2004, 279, 48013-48023.	3.4	61