

Andrea Grilli

List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

1,257
citations

623574

14
h-index

752573

20
g-index

21
all docs

21
docs citations

21
times ranked

3128
citing authors

#	ARTICLE	IF	CITATIONS
1	Jagged1 is the pathological link between Wnt and Notch pathways in colorectal cancer. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 6315-6320.	3.3	338
2	High-dimensional single cell analysis identifies stem-like cytotoxic CD8+ T cells infiltrating human tumors. Journal of Experimental Medicine, 2018, 215, 2520-2535.	4.2	250
3	miR-34a predicts survival of Ewing's sarcoma patients and directly influences cell chemosensitivity and malignancy. Journal of Pathology, 2012, 226, 796-805.	2.1	128
4	MicroRNAs involvement in fludarabine refractory chronic lymphocytic leukemia. Molecular Cancer, 2010, 9, 123.	7.9	107
5	Single-cell analyses reveal YAP/TAZ as regulators of stemness and cell plasticity in glioblastoma. Nature Cancer, 2021, 2, 174-188.	5.7	83
6	Bimodal CD40/Fas-Dependent Crosstalk between iNKT Cells and Tumor-Associated Macrophages Impairs Prostate Cancer Progression. Cell Reports, 2018, 22, 3006-3020.	2.9	62
7	Identification of Common and Distinctive Mechanisms of Resistance to Different Anti-IGF-IR Agents in Ewing's Sarcoma. Molecular Endocrinology, 2012, 26, 1603-1616.	3.7	53
8	Transcriptional profiling of human bronchial epithelial cell BEAS-2B exposed to diesel and biomass ultrafine particles. BMC Genomics, 2018, 19, 302.	1.2	43
9	CD99 triggering induces methuosis of Ewing sarcoma cells through IGF-1R/RAS/Rac1 signaling. Oncotarget, 2016, 7, 79925-79942.	0.8	40
10	Insulin-Like Growth Factor 2 mRNA-Binding Protein 3 is a Novel Post-Transcriptional Regulator of Ewing Sarcoma Malignancy. Clinical Cancer Research, 2018, 24, 3704-3716.	3.2	33
11	High NRF2 Levels Correlate with Poor Prognosis in Colorectal Cancer Patients and with Sensitivity to the Kinase Inhibitor AT9283 In Vitro. Biomolecules, 2020, 10, 1365.	1.8	22
12	A gene expression signature of Retinoblastoma loss-of-function predicts resistance to neoadjuvant chemotherapy in ER-positive/HER2-positive breast cancer patients. Breast Cancer Research and Treatment, 2018, 170, 329-341.	1.1	17
13	Exome sequencing and bioinformatic approaches reveals rare sequence variants involved in cell signalling and elastic fibre homeostasis: new evidence in the development of ectopic calcification. Cellular Signalling, 2019, 59, 131-140.	1.7	15
14	Slurry surfacing: a review of definitions, descriptions and current practices. RILEM Technical Letters, 0, 4, 103-109.	0.0	15
15	Fatal cytokine release syndrome by an aberrant FLIP/STAT3 axis. Cell Death and Differentiation, 2022, 29, 420-438.	5.0	14
16	Insulin-like growth factor 1 receptor affects the survival of primary prostate cancer patients depending on TMPRSS2-ERG status. BMC Cancer, 2017, 17, 367.	1.1	10
17	Gene expression profile correlates with molecular and clinical features in patients with myelofibrosis. Blood Advances, 2021, 5, 1452-1462.	2.5	8
18	RNA-seq in DMD urinary stem cells recognized muscle-related transcription signatures and addressed the identification of atypical mutations by whole-genome sequencing. Human Genetics and Genomics Advances, 2022, 3, 100054.	1.0	6

#	ARTICLE	IF	CITATIONS
19	RNA aptamers specific for transmembrane p24 trafficking protein 6 and Clusterin for the targeted delivery of imaging reagents and RNA therapeutics to human \hat{i}^2 cells. Nature Communications, 2022, 13, 1815.	5.8	6
20	Urine-Derived Stem Cells Express 571 Neuromuscular Disorders Causing Genes, Making Them a Potential in vitro Model for Rare Genetic Diseases. Frontiers in Physiology, 2021, 12, 716471.	1.3	4
21	Human T cells engineered with a leukemia lipid-specific TCR enables donor-unrestricted recognition of CD1c-expressing leukemia. Nature Communications, 2021, 12, 4844.	5.8	3