Mariateresa Fulciniti

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11756105/publications.pdf

Version: 2024-02-01

21 papers 3,867 citations

471371 17 h-index 752573 20 g-index

22 all docs 22 docs citations

times ranked

22

8380 citing authors

#	Article	IF	CITATIONS
1	Heterogeneity of genomic evolution and mutational profiles in multiple myeloma. Nature Communications, 2014, 5, 2997.	5.8	741
2	Blockade of XBP1 splicing by inhibition of IRE1 $\hat{l}\pm$ is a promising therapeutic option in multiple myeloma. Blood, 2012, 119, 5772-5781.	0.6	353
3	Identification of novel mutational drivers reveals oncogene dependencies in multiple myeloma. Blood, 2018, 132, 587-597.	0.6	335
4	A high-risk, Double-Hit, group of newly diagnosed myeloma identified by genomic analysis. Leukemia, 2019, 33, 159-170.	3.3	313
5	Elevated IL-17 produced by Th17 cells promotes myeloma cell growth and inhibits immune function in multiple myeloma. Blood, 2010, 115, 5385-5392.	0.6	300
6	Targeted Disruption of the BCL9 \hat{l}^2 -Catenin Complex Inhibits Oncogenic Wnt Signaling. Science Translational Medicine, 2012, 4, 148ra117.	5.8	214
7	Pharmacologic targeting of a stem/progenitor population in vivo is associated with enhanced bone regeneration in mice. Journal of Clinical Investigation, 2008, 118, 491-504.	3.9	202
8	The Monoclonal Antibody nBT062 Conjugated to Cytotoxic Maytansinoids Has Selective Cytotoxicity Against CD138-Positive Multiple Myeloma Cells <i>In vitro</i> local cancer Research, 2009, 15, 4028-4037.	3.2	200
9	Activin A promotes multiple myeloma-induced osteolysis and is a promising target for myeloma bone disease. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 5124-5129.	3.3	196
10	Genomic landscape and chronological reconstruction of driver events in multiple myeloma. Nature Communications, 2019, 10, 3835.	5.8	183
11	Drugging the IncRNA MALAT1 via LNA gapmeR ASO inhibits gene expression of proteasome subunits and triggers anti-multiple myeloma activity. Leukemia, 2018, 32, 1948-1957.	3.3	179
12	PI3K/p110δis a novel therapeutic target in multiple myeloma. Blood, 2010, 116, 1460-1468.	0.6	177
13	Bruton tyrosine kinase inhibition is a novel therapeutic strategy targeting tumor in the bone marrow microenvironment in multiple myeloma. Blood, 2012, 120, 1877-1887.	0.6	162
14	Therapeutic Targeting of miR-29b/HDAC4 Epigenetic Loop in Multiple Myeloma. Molecular Cancer Therapeutics, 2016, 15, 1364-1375.	1.9	94
15	The Cyclophilin A–CD147 complex promotes the proliferation and homing of multiple myeloma cells. Nature Medicine, 2015, 21, 572-580.	15.2	79
16	Therapeutic Targeting of miR-29b/HDAC4 Epigenetic Loop in Multiple Myeloma. Molecular Cancer Therapeutics, 2016, 15, 1364-1375.	1.9	60
17	Genome-Wide Somatic Alterations in Multiple Myeloma Reveal a Superior Outcome Group. Journal of Clinical Oncology, 2020, 38, 3107-3118.	0.8	45
18	CRISPR Interference (CRISPRi) and CRISPR Activation (CRISPRa) to Explore the Oncogenic IncRNA Network. Methods in Molecular Biology, 2021, 2348, 189-204.	0.4	12

#	Article	IF	CITATIONS
19	The effects of MicroRNA deregulation on pre-RNA processing network in multiple myeloma. Leukemia, 2020, 34, 167-179.	3.3	11
20	Deficiency of IL-17A, but not the prototypical Th17 transcription factor $ROR\hat{I}^3$ t, decreases murine spontaneous intestinal tumorigenesis. Cancer Immunology, Immunotherapy, 2016, 65, 13-24.	2.0	10
21	Role of Wnt Signaling Pathways in Multiple Myeloma Pathogenesis. , 2013, , 85-95.		1