Daniel Onofre Vidal

List of Publications by Year
in descending order

30	1,796	567281 15	501196 28
papers	citations	h-index	g-index

30	30	30	3911
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	lncRNA Epigenetic Landscape Analysis Identifies EPIC1 as an Oncogenic lncRNA that Interacts with MYC and Promotes Cell-Cycle Progression in Cancer. Cancer Cell, 2018, 33, 706-720.e9.	16.8	400
2	Somatic Mutational Landscape of Splicing Factor Genes and Their Functional Consequences across 33 Cancer Types. Cell Reports, 2018, 23, 282-296.e4.	6.4	333
3	Integrated Molecular Characterization of Testicular Germ Cell Tumors. Cell Reports, 2018, 23, 3392-3406.	6.4	324
4	A Pan-Cancer Analysis of Enhancer Expression in Nearly 9000 Patient Samples. Cell, 2018, 173, 386-399.e12.	28.9	228
5	Sense-antisense pairs in mammals: functional and evolutionary considerations. Genome Biology, 2007, 8, R40.	9.6	55
6	High Expression of HULC Is Associated with Poor Prognosis in Osteosarcoma Patients. PLoS ONE, 2016, 11, e0156774.	2.5	54
7	MGMT and CALCA promoter methylation are associated with poor prognosis in testicular germ cell tumor patients. Oncotarget, 2017, 8, 50608-50617.	1.8	52
8	miR-450a Acts as a Tumor Suppressor in Ovarian Cancer by Regulating Energy Metabolism. Cancer Research, 2019, 79, 3294-3305.	0.9	51
9	CYP1A2*1C, CYP2E1*5B, and GSTM1 polymorphisms are predictors of risk and poor outcome in head and neck squamous cell carcinoma patients. Oral Oncology, 2009, 45, e73-e79.	1.5	48
10	Overexpression of mir-183 and mir-494 promotes proliferation and migration in human breast cancer cell lines. Oncology Letters, 2017, 14, 1054-1060.	1.8	40
11	Aberrant methylation in pediatric myelodysplastic syndrome. Leukemia Research, 2007, 31, 175-181.	0.8	39
12	Placenta-Enriched LincRNAs MIR503HG and LINC00629 Decrease Migration and Invasion Potential of JEG-3 Cell Line. PLoS ONE, 2016, 11, e0151560.	2.5	36
13	Hypermethylation of CpG island in the promoter region of CALCA in acute lymphoblastic leukemia with central nervous system (CNS) infiltration correlates with poorer prognosis. Leukemia Research, 2006, 30, 891-894.	0.8	18
14	Absence of microsatellite instability andBRAF(V600E) mutation in testicular germ cell tumors. Andrology, 2016, 4, 866-872.	3.5	18
15	The Role of microRNAs in Medulloblastoma. Pediatric Hematology and Oncology, 2013, 30, 367-378.	0.8	15
16	Characteristics of the phenotypic abnormalities of bone marrow cells in childhood myelodysplastic syndromes and juvenile myelomonocytic leukemia. Pediatric Blood and Cancer, 2017, 64, e26285.	1.5	14
17	Hotspot TERT promoter mutations are rare events in testicular germ cell tumors. Tumor Biology, 2016, 37, 4901-4907.	1.8	13
18	Brachyury oncogene is a prognostic factor in highâ€risk testicular germ cell tumors. Andrology, 2018, 6, 597-604.	3.5	11

#	Article	IF	Citations
19	Highly expressed placental miRNAs control key biological processes in human cancer cell lines. Oncotarget, 2018, 9, 23554-23563.	1.8	10
20	Insights in Osteosarcoma by Proton Nuclear Magnetic Resonance Serum Metabonomics. Frontiers in Oncology, 2020, 10, 506959.	2.8	9
21	The Brazilian TP53 mutation (R337H) and sarcomas. PLoS ONE, 2020, 15, e0227260.	2.5	6
22	Analysis of allelic differential expression in the human genome using allele-specific serial analysis of gene expression tags. Genome, 2011, 54, 120-127.	2.0	5
23	Immunophenotypic characteristics of juvenile myelomonocytic leukaemia and their relation with the molecular subgroups of the disease. British Journal of Haematology, 2021, 192, 129-136.	2.5	5
24	Drug Resistance and Methylation in Myelodysplastic Syndrome. Current Pharmaceutical Biotechnology, 2007, 8, 77-81.	1.6	3
25	Array-CGH as an adjuvant tool in cytogenetic diagnosis of pediatric MDS and JMML. Medical Oncology, 2013, 30, 734.	2.5	3
26	Dysregulation of interferon regulatory genes reinforces the concept of chronic immune response in myelodysplastic syndrome pathogenesis. Hematological Oncology, 2019, 37, 523-526.	1.7	3
27	Heteroduplex formation and S1 digestion for mapping alternative splicing sites. Genetics and Molecular Research, 2008, 7, 958-969.	0.2	2
28	1031-1034delTAAC (Leu125Stop): a novel familial UBE3A mutation causing Angelman syndrome in two siblings showing distinct phenotypes. BMC Medical Genetics, 2012, 13, 124.	2.1	1
29	Prognosis value of HER2 in osteosarcomas: A systematic review with meta-analysis.. Journal of Clinical Oncology, 2015, 33, e21504-e21504.	1.6	0
30	Brachyury, a driver of epithelial mesenchymal transition, as an independent prognostic factor in high-grade testicular germ cell tumors.. Journal of Clinical Oncology, 2017, 35, e16039-e16039.	1.6	0