Eva Hernando

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

3,251 23 54 57 h-index g-index citations papers 60 4.62 3,859 10.4 avg, IF L-index ext. citations ext. papers

#	Paper	IF	Citations
54	Mad2 overexpression promotes aneuploidy and tumorigenesis in mice. <i>Cancer Cell</i> , 2007 , 11, 9-23	24.3	488
53	Aberrant miR-182 expression promotes melanoma metastasis by repressing FOXO3 and microphthalmia-associated transcription factor. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009 , 106, 1814-9	11.5	461
52	The histone variant macroH2A suppresses melanoma progression through regulation of CDK8. <i>Nature</i> , 2010 , 468, 1105-9	50.4	291
51	miR-30b/30d regulation of GalNAc transferases enhances invasion and immunosuppression during metastasis. <i>Cancer Cell</i> , 2011 , 20, 104-18	24.3	278
50	Melanoma MicroRNA signature predicts post-recurrence survival. Clinical Cancer Research, 2010, 16, 157	77 <u>-</u> 86	180
49	BRD4 sustains melanoma proliferation and represents a new target for epigenetic therapy. <i>Cancer Research</i> , 2013 , 73, 6264-76	10.1	161
48	A Systems Biology Approach Identifies FUT8 as a Driver of Melanoma Metastasis. <i>Cancer Cell</i> , 2017 , 31, 804-819.e7	24.3	146
47	Control of embryonic stem cell identity by BRD4-dependent transcriptional elongation of super-enhancer-associated pluripotency genes. <i>Cell Reports</i> , 2014 , 9, 234-247	10.6	144
46	Histone Variant H2A.Z.2 Mediates Proliferation and Drug Sensitivity of Malignant Melanoma. <i>Molecular Cell</i> , 2015 , 59, 75-88	17.6	128
45	Epigenetic Silencing of CDR1as Drives IGF2BP3-Mediated Melanoma Invasion and Metastasis. <i>Cancer Cell</i> , 2020 , 37, 55-70.e15	24.3	113
44	FBXW7 modulates cellular stress response and metastatic potential through HSF1 post-translational modification. <i>Nature Cell Biology</i> , 2015 , 17, 322-332	23.4	89
43	miR-204-5p and miR-211-5p Contribute to BRAF Inhibitor Resistance in Melanoma. <i>Cancer Research</i> , 2018 , 78, 1017-1030	10.1	88
42	MicroRNA and cutaneous melanoma: from discovery to prognosis and therapy. <i>Carcinogenesis</i> , 2012 , 33, 1823-32	4.6	73
41	Limited Environmental Serine and Glycine Confer Brain Metastasis Sensitivity to PHGDH Inhibition. <i>Cancer Discovery</i> , 2020 , 10, 1352-1373	24.4	62
40	Harnessing BET Inhibitor Sensitivity Reveals AMIGO2 as a Melanoma Survival Gene. <i>Molecular Cell</i> , 2017 , 68, 731-744.e9	17.6	53
39	A miRNA-Based Signature Detected in Primary Melanoma Tissue Predicts Development of Brain Metastasis. <i>Clinical Cancer Research</i> , 2015 , 21, 4903-12	12.9	49
38	Anti-miR182 reduces ovarian cancer burden, invasion, and metastasis: an in vivo study in orthotopic xenografts of nude mice. <i>Molecular Cancer Therapeutics</i> , 2014 , 13, 1729-39	6.1	47

(2013-2011)

37	Integrative genomics identifies molecular alterations that challenge the linear model of melanoma progression. <i>Cancer Research</i> , 2011 , 71, 2561-71	10.1	45	
36	Revisiting determinants of prognosis in cutaneous melanoma. <i>Cancer</i> , 2015 , 121, 4108-23	6.4	40	
35	Identification of metastasis-suppressive microRNAs in primary melanoma. <i>Journal of the National Cancer Institute</i> , 2015 , 107,	9.7	38	
34	BET and BRAF inhibitors act synergistically against BRAF-mutant melanoma. <i>Cancer Medicine</i> , 2016 , 5, 1183-93	4.8	38	
33	circSamd4 represses myogenic transcriptional activity of PUR proteins. <i>Nucleic Acids Research</i> , 2020 , 48, 3789-3805	20.1	34	
32	Lysyl oxidase-like 3 is required for melanoma cell survival by maintaining genomic stability. <i>Cell Death and Differentiation</i> , 2018 , 25, 935-950	12.7	29	
31	MicroRNA-125a promotes resistance to BRAF inhibitors through suppression of the intrinsic apoptotic pathway. <i>Pigment Cell and Melanoma Research</i> , 2017 , 30, 328-338	4.5	23	
30	A Leukocyte Infiltration Score Defined by a Gene Signature Predicts Melanoma Patient Prognosis. <i>Molecular Cancer Research</i> , 2019 , 17, 109-119	6.6	20	
29	Limited miR-17-92 overexpression drives hematologic malignancies. <i>Leukemia Research</i> , 2015 , 39, 335-	41 .7	17	
28	Cancer. Aneuploidy advantages?. <i>Science</i> , 2008 , 322, 692-3	33.3	14	
27	A TGFEmiR-182-BRCA1 axis controls the mammary differentiation hierarchy. <i>Science Signaling</i> , 2016 , 9, ra118	8.8	12	
26	Identification of gene expression levels in primary melanoma associated with clinically meaningful characteristics. <i>Melanoma Research</i> , 2018 , 28, 380-389	3.3	12	
25	The State of Melanoma: Emergent Challenges and Opportunities. <i>Clinical Cancer Research</i> , 2021 , 27, 2678-2697	12.9	11	
24	Mutation burden as a potential prognostic marker of melanoma progression and survival <i>Journal of Clinical Oncology</i> , 2017 , 35, 9567-9567	2.2	10	
23	TYRP1 mRNA goes fishing for miRNAs in melanoma. <i>Nature Cell Biology</i> , 2017 , 19, 1311-1312	23.4	8	
22	Krppel-like factor 4 (KLF4) regulates the miR-183~96~182 cluster under physiologic and pathologic conditions. <i>Oncotarget</i> , 2017 , 8, 26298-26311	3.3	8	
21	Functional analysis of RPS27 mutations and expression in melanoma. <i>Pigment Cell and Melanoma Research</i> , 2020 , 33, 466-479	4.5	8	
20	In vivo Modeling and Molecular Characterization: A Path Toward Targeted Therapy of Melanoma Brain Metastasis. <i>Frontiers in Oncology</i> , 2013 , 3, 127	5.3	6	

19	HNRNPM controls circRNA biogenesis and splicing fidelity to sustain cancer cell fitness. <i>ELife</i> , 2021 , 10,	8.9	6
18	Network models of primary melanoma microenvironments identify key melanoma regulators underlying prognosis. <i>Nature Communications</i> , 2021 , 12, 1214	17.4	6
17	Expression of miR-16 is not a suitable reference for analysis of serum microRNAs in melanoma patients. <i>Journal of Biomedical Science and Engineering</i> , 2012 , 05, 647-651	0.7	4
16	Treatment with therapeutic anticoagulation is not associated with immunotherapy response in advanced cancer patients. <i>Journal of Translational Medicine</i> , 2021 , 19, 47	8.5	4
15	Regulates the Proliferation Capacity of Bone-Marrow Derived Mesenchymal Stem Cells. <i>Cells</i> , 2020 , 9,	7.9	3
14	Melanoma-secreted Amyloid Beta Suppresses Neuroinflammation and Promotes Brain Metastasis <i>Cancer Discovery</i> , 2022 ,	24.4	2
13	Characterization of MicroRNAs Regulating FOXO Expression. <i>Methods in Molecular Biology</i> , 2019 , 1890, 13-28	1.4	1
12	The histone demethylase PHF8 regulates TGFIsignaling and promotes melanoma metastasis <i>Science Advances</i> , 2022 , 8, eabi7127	14.3	1
11	miRNA Decoy Screen Reveals miR-124a as a Suppressor of Melanoma Metastasis <i>Frontiers in Oncology</i> , 2022 , 12, 852952	5.3	O
10	Human genes differ by their UV sensitivity estimated through analysis of UV-induced silent mutations in melanoma. <i>Human Mutation</i> , 2020 , 41, 1751-1760	4.7	
9	Preclinical testing supports combined BET and BRAF inhibition as a promising therapeutic strategy for melanoma <i>Journal of Clinical Oncology</i> , 2014 , 32, 9072-9072	2.2	
8	Targeted next-generation sequencing of melanoma patient samples to reveal mutations in non-protein coding regions of targetable oncogenes <i>Journal of Clinical Oncology</i> , 2016 , 34, 9559-9559	2.2	
7	Genomic characterization of acral lentiginous melanoma: Identification of altered metabolism as a potential therapeutic target <i>Journal of Clinical Oncology</i> , 2016 , 34, 9524-9524	2.2	
6	Newmouse models of melanoma metastasis and differences in brain tropism and metastatic growth pattern <i>Journal of Clinical Oncology</i> , 2012 , 30, e19015-e19015	2.2	
5	MicroRNA alterations associated with BRAF status in melanoma <i>Journal of Clinical Oncology</i> , 2012 , 30, 8565-8565	2.2	
4	Early alterations of microRNA expression to predict and modulate melanoma metastasis <i>Journal of Clinical Oncology</i> , 2012 , 30, 8550-8550	2.2	
3	Identification of melanoma-specific alterations in cell surface glycosylation <i>Journal of Clinical Oncology</i> , 2012 , 30, e19018-e19018	2.2	
2	Melanoma recurrence risk stratification using Bayesian systems biology modeling <i>Journal of Clinical Oncology</i> , 2013 , 31, 9089-9089	2.2	

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