

# Alfons Navarro

## List of Publications by Year in descending order

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121  
papers

4,285  
citations

117453

34  
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110170

64  
g-index

122  
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122  
docs citations

122  
times ranked

7394  
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification by Real-time PCR of 13 mature microRNAs differentially expressed in colorectal cancer and non-tumoral tissues. <i>Molecular Cancer</i> , 2006, 5, 29.	7.9	744
2	miR-34a as a prognostic marker of relapse in surgically resected non-small-cell lung cancer. <i>Carcinogenesis</i> , 2009, 30, 1903-1909.	1.3	314
3	Pint lincRNA connects the p53 pathway with epigenetic silencing by the Polycomb repressive complex 2. <i>Genome Biology</i> , 2013, 14, R104.	13.9	224
4	Overlapping expression of microRNAs in human embryonic colon and colorectal cancer. <i>Cell Research</i> , 2008, 18, 823-833.	5.7	174
5	MicroRNA expression profiling in classic Hodgkin lymphoma. <i>Blood</i> , 2008, 111, 2825-2832.	0.6	161
6	Regulation of JAK2 by miR-135a: prognostic impact in classic Hodgkin lymphoma. <i>Blood</i> , 2009, 114, 2945-2951.	0.6	157
7	miR-141 and miR-200c as Markers of Overall Survival in Early Stage Non-Small Cell Lung Cancer Adenocarcinoma. <i>PLoS ONE</i> , 2014, 9, e101899.	1.1	124
8	Epigenetic regulation mechanisms of microRNA expression. <i>Biomolecular Concepts</i> , 2017, 8, 203-212.	1.0	109
9	Mannan-binding lectin pathway deficiencies and invasive fungal infections following allogeneic stem cell transplantation. <i>Experimental Hematology</i> , 2006, 34, 1435-1441.	0.2	89
10	YKT6 expression, exosome release, and survival in non-small cell lung cancer. <i>Oncotarget</i> , 2016, 7, 51515-51524.	0.8	88
11	Hematopoiesis-related microRNA expression in myelodysplastic syndromes. <i>Leukemia and Lymphoma</i> , 2009, 50, 1854-1859.	0.6	85
12	The lincRNA<i>HOTAIRM1</i>, located in the<i>HOXA</i>genomic region, is expressed in acute myeloid leukemia, impacts prognosis in patients in the intermediate-risk cytogenetic category, and is associated with a distinctive microRNA signature. <i>Oncotarget</i> , 2015, 6, 31613-31627.	0.8	78
13	LincRNA-p21 Impacts Prognosis in Resected Non-“Small Cell Lung Cancer Patients through Angiogenesis Regulation. <i>Journal of Thoracic Oncology</i> , 2016, 11, 2173-2182.	0.5	76
14	Single-nucleotide polymorphisms in base excision repair, nucleotide excision repair, and double strand break genes as markers for response to radiotherapy in patients with Stage I to II head-and-neck cancer. <i>International Journal of Radiation Oncology Biology Physics</i> , 2006, 66, 1022-1030.	0.4	75
15	Low miR-145 and high miR-367 are associated with unfavourable prognosis in resected nonsmall cell lung cancer. <i>European Respiratory Journal</i> , 2013, 41, 1172-1178.	3.1	72
16	Tumour CD133 mRNA expression and clinical outcome in surgically resected colorectal cancer patients. <i>European Journal of Cancer</i> , 2010, 46, 642-649.	1.3	68
17	Lestaurtinib Inhibition of the JAK/STAT Signaling Pathway in Hodgkin Lymphoma Inhibits Proliferation and Induces Apoptosis. <i>PLoS ONE</i> , 2011, 6, e18856.	1.1	64
18	Genomic polymorphisms provide prognostic information in intermediate-risk acute myeloblastic leukemia. <i>Blood</i> , 2006, 107, 4871-4879.	0.6	62

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19	Role of miR-200 family members in survival of colorectal cancer patients treated with fluoropyrimidines. <i>Journal of Surgical Oncology</i> , 2014, 109, 676-683.	0.8	62
20	MicroRNA expression at diagnosis adds relevant prognostic information to molecular categorization in patients with intermediate-risk cytogenetic acute myeloid leukemia. <i>Leukemia</i> , 2014, 28, 804-812.	3.3	59
21	MicroRNAs Expressed during Lung Cancer Development Are Expressed in Human Pseudoglandular Lung Embryogenesis. <i>Oncology</i> , 2009, 76, 162-169.	0.9	57
22	Single Nucleotide Polymorphisms in Nucleotide Excision Repair Genes XPA, XPD, XPG and ERCC1 in Advanced Colorectal Cancer Patients Treated with First-Line Oxaliplatin/Fluoropyrimidine. <i>Oncology</i> , 2007, 72, 364-370.	0.9	56
23	Impact of MiRSNPs on Survival and Progression in Patients with Multiple Myeloma Undergoing Autologous Stem Cell Transplantation. <i>Clinical Cancer Research</i> , 2012, 18, 3697-3704.	3.2	54
24	Prognostic implications of miR-16 expression levels in resected non-small cell lung cancer. <i>Journal of Surgical Oncology</i> , 2011, 103, 411-415.	0.8	52
25	A Dual Role for KRT81: A miR-SNP Associated with Recurrence in Non-Small-Cell Lung Cancer and a Novel Marker of Squamous Cell Lung Carcinoma. <i>PLoS ONE</i> , 2011, 6, e22509.	1.1	50
26	Effect of NOD2/CARD15 variants in T-cell depleted allogeneic stem cell transplantation. <i>Haematologica</i> , 2006, 91, 1372-6.	1.7	48
27	Acute myeloid leukemia with translocation (8;16)(p11;p13) and MYST3-CREBBP rearrangement harbors a distinctive microRNA signature targeting RET proto-oncogene. <i>Leukemia</i> , 2013, 27, 595-603.	3.3	45
28	The significance of PIWI family expression in human lung embryogenesis and non-small cell lung cancer. <i>Oncotarget</i> , 2015, 6, 31544-31556.	0.8	45
29	A serum microRNA signature associated with complete remission and progression after autologous stem-cell transplantation in patients with multiple myeloma. <i>Oncotarget</i> , 2015, 6, 1874-1883.	0.8	42
30	MicroRNAs in Human Embryonic and Cancer Stem Cells. <i>Yonsei Medical Journal</i> , 2010, 51, 622.	0.9	41
31	miR-328 mediates a metabolic shift in colon cancer cells by targeting SLC2A1/GLUT1. <i>Clinical and Translational Oncology</i> , 2018, 20, 1161-1167.	1.2	41
32	PiwiRNA-651 as marker of treatment response and survival in classical Hodgkin lymphoma. <i>Oncotarget</i> , 2016, 7, 46002-46013.	0.8	41
33	Identifying High-Risk Stage II Colon Cancer Patients: A Three-MicroRNA-Based Score as a Prognostic Biomarker. <i>Clinical Colorectal Cancer</i> , 2016, 15, e175-e182.	1.0	36
34	Sonic hedgehog mRNA expression by real-time quantitative PCR in normal and tumor tissues from colorectal cancer patients. <i>Cancer Letters</i> , 2006, 233, 117-123.	3.2	35
35	Exosomal microRNAs isolated from plasma of mesenteric veins linked to liver metastases in resected patients with colon cancer. <i>Oncotarget</i> , 2017, 8, 30859-30869.	0.8	35
36	Common variants in NLRP2 and NLRP3 genes are strong prognostic factors for the outcome of HLA-identical sibling allogeneic stem cell transplantation. <i>Blood</i> , 2008, 112, 4337-4342.	0.6	34

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37	miR-21, miR-99b and miR-375 combination as predictive response signature for preoperative chemoradiotherapy in rectal cancer. <i>PLoS ONE</i> , 2018, 13, e0206542.	1.1	34
38	Peri-prosthetic bone cysts after total ankle replacement. A systematic review and meta-analysis. <i>Foot and Ankle Surgery</i> , 2019, 25, 96-105.	0.8	34
39	Extracellular Vesicle lincRNA-p21 Expression in Tumor-Draining Pulmonary Vein Defines Prognosis in NSCLC and Modulates Endothelial Cell Behavior. <i>Cancers</i> , 2020, 12, 734.	1.7	34
40	Prognostic Impact of miR-200 Family Members in Plasma and Exosomes from Tumor-Draining versus Peripheral Veins of Colon Cancer Patients. <i>Oncology</i> , 2018, 95, 309-318.	0.9	32
41	Epigenetic events in normal colonic mucosa surrounding colorectal cancer lesions. <i>European Journal of Cancer</i> , 2008, 44, 2689-2695.	1.3	30
42	Exosome Analysis in Tumor-Draining Pulmonary Vein Identifies NSCLC Patients with Higher Risk of Relapse after Curative Surgery. <i>Cancers</i> , 2019, 11, 249.	1.7	30
43	Impact on response and survival of DNA repair single nucleotide polymorphisms in relapsed or refractory multiple myeloma patients treated with thalidomide. <i>Leukemia Research</i> , 2011, 35, 1178-1183.	0.4	26
44	MicroRNA expression in chronic lymphocytic leukemia developing autoimmune hemolytic anemia. <i>Leukemia and Lymphoma</i> , 2013, 54, 2016-2022.	0.6	26
45	circFBXW7 attenuates malignant progression in lung adenocarcinoma by sponging miR-942-5p. <i>Translational Lung Cancer Research</i> , 2021, 10, 1457-1473.	1.3	24
46	MicroRNAs 142-3p, miR-155 and miR-203 Are Deregulated in Gastric MALT Lymphomas Compared to Chronic Gastritis. <i>Cancer Genomics and Proteomics</i> , 2017, 14, 75-82.	1.0	23
47	Clinical significance of long non-coding RNA HOTTIP in early-stage non-small-cell lung cancer. <i>BMC Pulmonary Medicine</i> , 2019, 19, 55.	0.8	22
48	MiR-SNPs as Markers of Toxicity and Clinical Outcome in Hodgkin Lymphoma Patients. <i>PLoS ONE</i> , 2013, 8, e64716.	1.1	21
49	miR-203 and miR-221 regulate SOCS1 and SOCS3 in essential thrombocythemia. <i>Blood Cancer Journal</i> , 2016, 6, e406-e406.	2.8	21
50	Proteomic Analysis of Liquid Biopsy from Tumor-Draining Vein Indicates that High Expression of Exosomal ECM1 Is Associated with Relapse in Stage I-III Colon Cancer. <i>Translational Oncology</i> , 2018, 11, 715-721.	1.7	20
51	The expression level of BAALC-associated microRNA miR-3151 is an independent prognostic factor in younger patients with cytogenetic intermediate-risk acute myeloid leukemia. <i>Blood Cancer Journal</i> , 2015, 5, e352-e352.	2.8	19
52	Immunophenotypic analysis and quantification of B-1 and B-2 B cells during human fetal hematopoietic development. <i>Leukemia</i> , 2016, 30, 1603-1606.	3.3	18
53	Impact of global and gene-specific DNA methylation pattern in relapsed multiple myeloma patients treated with bortezomib. <i>Leukemia Research</i> , 2013, 37, 641-646.	0.4	17
54	NKX2-1 expression as a prognostic marker in early-stage non-small-cell lung cancer. <i>BMC Pulmonary Medicine</i> , 2017, 17, 197.	0.8	17

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55	Dihydropyrimidine dehydrogenases and cytidineâ€¢deaminase gene polymorphisms as outcome predictors in resected gastric cancer patients treated with fluoropyrimidine adjuvant chemotherapy. <i>Journal of Surgical Oncology</i> , 2008, 98, 130-134.	0.8	16
56	High levels of global DNA methylation are an independent adverse prognostic factor in a series of 90 patients with de novo myelodysplastic syndrome. <i>Leukemia Research</i> , 2014, 38, 874-881.	0.4	16
57	A 4-gene expression prognostic signature might guide post-remission therapy in patients with intermediate-risk cytogenetic acute myeloid leukemia. <i>Leukemia and Lymphoma</i> , 2018, 59, 2394-2404.	0.6	16
58	Non-Coding RNAs in Hodgkin Lymphoma. <i>International Journal of Molecular Sciences</i> , 2017, 18, 1154.	1.8	15
59	Ultrastructural and Immunohistochemical Analysis of Intestinal Myofibroblasts During the Early Organogenesis of the Human Small Intestine. <i>Anatomical Record</i> , 2011, 294, 462-471.	0.8	14
60	Epigenetic regulation of microRNA expression in Hodgkin lymphoma. <i>Leukemia and Lymphoma</i> , 2015, 56, 2683-2689.	0.6	13
61	Differential MIR-21 Expression in Plasma From Mesenteric Versus Peripheral Veins. <i>Medicine (United Tj ETQq1 1 0.784314 rgBT /Over</i>	0.4	12
62	Insertional anatomy of peroneal brevis and longus tendon â€” A cadaveric study. <i>Foot and Ankle Surgery</i> , 2019, 25, 636-639.	0.8	11
63	Pharmacogenomics: a tool for improving cancer chemotherapy. <i>Clinical and Translational Oncology</i> , 2008, 10, 628-637.	1.2	10
64	Anti-tumor activity of the combination of bendamustine with vorinostat in diffuse large B-cell lymphoma cells. <i>Leukemia and Lymphoma</i> , 2016, 57, 692-699.	0.6	10
65	Role of the epithelial-mesenchymal transition-related circular RNA, circ-10720, in non-small-cell lung cancer. <i>Translational Lung Cancer Research</i> , 2021, 10, 1804-1818.	1.3	9
66	Regulation of a2-adrenoceptor gene expression by chronic lithium treatment in rat brain. <i>Methods and Findings in Experimental and Clinical Pharmacology</i> , 2010, 32, 721.	0.8	9
67	Characterization of the MicroRNA Cargo of Extracellular Vesicles Isolated from a Pulmonary Tumor-Draining Vein Identifies miR-203a-3p as a Relapse Biomarker for Resected Non-Small Cell Lung Cancer. <i>International Journal of Molecular Sciences</i> , 2022, 23, 7138.	1.8	7
68	Single Nucleotide Polymorphisms in Tobacco Metabolism and DNA Repair Genes and Prognosis in Resected Non-Small-Cell Lung Cancer. <i>Journal of Surgical Research</i> , 2011, 167, e5-e12.	0.8	6
69	LncRNA Quantification from Extracellular Vesicles Isolated from Blood Plasma or Conditioned Media. <i>Methods in Molecular Biology</i> , 2021, 2348, 285-304.	0.4	6
70	Lincp21-RNA as Predictive Response Marker for Preoperative Chemoradiotherapy in Rectal Cancer. <i>Journal of Personalized Medicine</i> , 2021, 11, 420.	1.1	6
71	LincRNA-p21 Levels Relates to Survival and Post-Operative Radiotherapy Benefit in Rectal Cancer Patients. <i>Life</i> , 2020, 10, 172.	1.1	5
72	Genetic Polymorphisms in the Inflammasomes Are Associated with Relapse and Survival in HLA-Identical Sibling Donor Allogeneic Stem Cell Transplantation.. <i>Blood</i> , 2007, 110, 1075-1075.	0.6	5

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73	The DNA damage repair-related gene PKMYT1 is a potential biomarker in various malignancies. <i>Translational Lung Cancer Research</i> , 2021, 10, 4600-4616.	1.3	5
74	The Os Peroneum incidence – A cadaveric study. <i>Foot and Ankle Surgery</i> , 2020, 26, 325-327.	0.8	4
75	Monitoring HOTTIP levels on extracellular vesicles for predicting recurrence in surgical non-small cell lung cancer patients. <i>Translational Oncology</i> , 2021, 14, 101144.	1.7	4
76	Prognostic Significance Of a 4-Microrna Signature Targeting JAK2 In Classical Hodgkin Lymphoma. <i>Blood</i> , 2013, 122, 629-629.	0.6	4
77	Piwirna-651 Expression Influences Treatment Response and Impacts Survival in Classical Hodgkin Lymphoma Patients through Regulation of ABCC5. <i>Blood</i> , 2014, 124, 134-134.	0.6	4
78	Screening for Prognostic microRNAs Associated with Treatment Failure in Diffuse Large B Cell Lymphoma. <i>Cancers</i> , 2022, 14, 1065.	1.7	4
79	Long Non-Coding RNA NNCI/NKX2-1 Duplex Impacts Prognosis in Stage I Non-Small-Cell Lung Cancer. <i>Archivos De Bronconeumologia</i> , 2020, 56, 630-636.	0.4	3
80	Editorial: The Role of ncRNAs in Solid Tumors Prognosis: From Laboratory to Clinical Utility. <i>Frontiers in Oncology</i> , 2020, 10, 631316.	1.3	3
81	Impact of Single Nucleotide Polymorphisms in Genes Involved in DNA Repair and Drug Metabolism On Survival After Autologous Stem Cell Transplantation in Patients with Multiple Myeloma.. <i>Blood</i> , 2012, 120, 2934-2934.	0.6	3
82	Twist1 activated circRNA-10720 is a new player in hepatocellular carcinoma metastasis. <i>Translational Cancer Research</i> , 2019, 8, S135-S140.	0.4	3
83	Midface Advancement in a Simple Approach. <i>Journal of Craniofacial Surgery</i> , 2017, 28, 108-112.	0.3	2
84	KRAS mutations by digital PCR in circulating tumor cells isolated from the mesenteric vein are associated with residual disease and overall survival in resected colorectal cancer patients. <i>International Journal of Colorectal Disease</i> , 2020, 35, 805-813.	1.0	2
85	Deciphering miRNA–Target Relationships to Understand miRNA-Mediated Carcinogenesis. <i>Cancers</i> , 2021, 13, 2415.	1.7	2
86	Exploring the Expression Profile of Long Non-Coding RNA (lncRNA) in Different Acute Myeloid Leukemia (AML) Subtypes: t(8;16)(p11;p13)/MYST3-Crebbp AML Harbors a Distinctive Lncrna Signature. <i>Blood</i> , 2015, 126, 1397-1397.	0.6	2
87	Argonaute-crosslinking and immunoprecipitation deciphers the liver miR-122 targetome. <i>Non-coding RNA Investigation</i> , 0, 1, 23-23.	0.6	2
88	A Distinctive MicroRNA Signature Characterizes Acute Myeloid Leukemia with Translocation (8;16)(p11;p13) and MYST3-CREBBP Rearrangement. <i>Blood</i> , 2010, 116, 230-230.	0.6	1
89	Salvage Therapy for Multiple Myeloma with Lenalidomide-Based Regimens: Impact of Single Nucleotide Polymorphisms Related to miRNA Pathways. <i>Blood</i> , 2014, 124, 2045-2045.	0.6	1
90	piRNA-651 as a prognostic marker in surgically resected non-small-cell lung cancer.. <i>Journal of Clinical Oncology</i> , 2015, 33, 11043-11043.	0.8	1

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91	Regulation of Jak2 Kinase Gene by the Mir-135 Family in Classical Hodgkin Lymphoma. Blood, 2008, 112, 518-518.	0.6	1
92	Octogenarians may benefit from stage-specific small cell lung cancer treatment. Translational Lung Cancer Research, 2021, 10, 3973-3982.	1.3	1
93	Long Non-Coding RNA NNCI/NKX2-1 Duplex Impacts Prognosis in Stage I Non-Small-Cell Lung Cancer. Archivos De Bronconeumologia, 2020, 56, 630-636.	0.4	1
94	Human embryonic mesenchymal lung-conditioned medium promotes differentiation to myofibroblast and loss of stemness phenotype in lung adenocarcinoma cell lines. Journal of Experimental and Clinical Cancer Research, 2022, 41, 37.	3.5	1
95	2.36 Chronic Lymphocytic Leukemia Associated with Autoimmune Hemolytic Anemia is Characterized by a Distinctive miRNA Signature. Clinical Lymphoma, Myeloma and Leukemia, 2011, 11, S181-S182.	0.2	0
96	Antitumor activity of the combination of bendamustine with vorinostat in diffuse large B-cell lymphoma cells. Clinical Lymphoma, Myeloma and Leukemia, 2015, 15, S61-S62.	0.2	0
97	Can long non-coding RNAs reveal the hidden code of lung cancer?. Non-coding RNA Investigation, 2020, 4, 6-6.	0.6	0
98	LincRNA-p21 as predictive response marker for preoperative chemoradiotherapy in rectal cancer.. Journal of Clinical Oncology, 2021, 39, e15534-e15534.	0.8	0
99	Clinic and therapeutic potential of non-coding RNAs in cancer. Translational Cancer Research, 2021, 10, 0-0.	0.4	0
100	Germline Polymorphisms Provide Strong Prognostic Information in Intermediate Risk Acute Myeloblastic Leukemia (AML).. Blood, 2004, 104, 398-398.	0.6	0
101	Polymorphisms of NOD2/CARD15 Are Associated with Clinical Outcome in T-Cell Depleted HLA-Identical Sibling Allogeneic Stem Cell Transplantation.. Blood, 2005, 106, 1408-1408.	0.6	0
102	Effect of Germline Polymorphisms on Clinical Outcome in Hodgkin's Lymphoma (HL).. Blood, 2006, 108, 2267-2267.	0.6	0
103	Analysis of microRNA Patterns in Hodgkin's Lymphoma (HL).. Blood, 2006, 108, 474-474.	0.6	0
104	Hematopoietic miRNAs Expression in Myelodysplastic Syndromes.. Blood, 2007, 110, 2422-2422.	0.6	0
105	Mir-135a Expression Is Associated with Relapse in Hodgkin Lymphoma.. Blood, 2007, 110, 2270-2270.	0.6	0
106	MicroRNA Analysis by In Situ Hybridization in Hodgkin Lymphoma.. Blood, 2007, 110, 2271-2271.	0.6	0
107	Impact on Response and Survival of DNA Repair Single Nucleotide Polymorphisms In Relapsed or Refractory Multiple Myeloma Patients Treated with Thalidomide. Blood, 2010, 116, 4041-4041.	0.6	0
108	Epigenetic Regulation of MicroRNA Expression In Hodgkin Lymphoma. Blood, 2010, 116, 3885-3885.	0.6	0

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109	Lestaurtinib (CEP701) Inhibits Proliferation and Activates Apoptosis In Hodgkin Lymphoma Cells through Inhibition of the JAK/STAT Signaling Pathway. Blood, 2010, 116, 3919-3919.	0.6	0
110	Mir-SNP Haplotype in the MicroRNA Maturation Pathway As a Marker of Clinical Outcome in Hodgkin Lymphoma (HL). Blood, 2011, 118, 1561-1561.	0.6	0
111	MicroRNAs in Intermediate Risk Cytogenetic Acute Myeloid Leukemia Add Relevant Prognostic Information to Molecular Categorization. Blood, 2011, 118, 235-235.	0.6	0
112	Impact of Global and Gene-Specific DNA Methylation Pattern in Relapsed Multiple Myeloma Patients Treated with Bortezomib. Blood, 2011, 118, 132-132.	0.6	0
113	Chronic Lymphocytic Leukemia Associated with Autoimmune Hemolytic Anemia Is Characterized by a Distinctive miRNA Signature,. Blood, 2011, 118, 3897-3897.	0.6	0
114	The Distinctive MicroRNA Signature of Acute Myeloid Leukemia with Translocation t(8;16)(p11;p13)/MYST3-CREBBP Is Responsible for RET Overexpression and Is Regulated by Epigenetic Mechanisms. Blood, 2011, 118, 2434-2434.	0.6	0
115	Impact of Single Nucleotide Polymorphisms in Genes Involved in miRNA Network on Survival and Progression After Autologous Stem Cell Transplantation in Patients with Multiple Myeloma,. Blood, 2011, 118, 3945-3945.	0.6	0
116	Overall Hydroxymethylation Levels As an Independent Prognostic Marker in Intermediate-Cytogenetic Risk Acute Myeloid Leukemia. Blood, 2012, 120, 1375-1375.	0.6	0
117	Global Methylation Provides Independent Prognostic Information in Chronic Lymphocytic Leukemia and Is Associated with a MicroRNA Signature. Blood, 2012, 120, 3873-3873.	0.6	0
118	Association of miR-141 and miR-200c with time to recurrence (TTR) and overall survival (OS) in resected non-small-cell lung cancer (NSCLC) adenocarcinoma (ADC) patients (p).. Journal of Clinical Oncology, 2013, 31, 7547-7547.	0.8	0
119	BAALC-Associated Mir-3151 Is An Independent Prognostic Factor In Younger Patients With Intermediate-Risk Cytogenetic Acute Myeloid Leukemia. Blood, 2013, 122, 2577-2577.	0.6	0
120	The LincRNA HOTAIRM1, Located in the HOXA genomic Region, impacts Prognosis in Acute Myeloid Leukemia and Is Associated with a Distinctive microRNA Signature. Blood, 2014, 124, 1003-1003.	0.6	0
121	NKX2-1 impacts prognosis in early stage NSCLC not harboring TP53 mutations. , 2016, , .		0