

Ji-Fan Hu

List of Publications by Year in descending order

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120
papers

5,562
citations

71102

41
h-index

88630

70
g-index

122
all docs

122
docs citations

122
times ranked

7370
citing authors

#	ARTICLE	IF	CITATIONS
1	CTCF Mediates Interchromosomal Colocalization Between Igf2/H19 and Wsb1/Nf1. <i>Science</i> , 2006, 312, 269-272.	12.6	413
2	A novel FLI1 exonic circular RNA promotes metastasis in breast cancer by coordinately regulating TET1 and DNMT1. <i>Genome Biology</i> , 2018, 19, 218.	8.8	292
3	Manipulation of nuclear architecture through CRISPR-mediated chromosomal looping. <i>Nature Communications</i> , 2017, 8, 15993.	12.8	224
4	CTCF Regulates Allelic Expression of <i>Igf2</i> by Orchestrating a Promoter-Polycomb Repressive Complex 2 Intrachromosomal Loop. <i>Molecular and Cellular Biology</i> , 2008, 28, 6473-6482.	2.3	193
5	Regulation of telomerase by alternate splicing of human telomerase reverse transcriptase (hTERT) in normal and neoplastic ovary, endometrium and myometrium. <i>International Journal of Cancer</i> , 2000, 85, 330-335.	5.1	174
6	IVF results in de novo DNA methylation and histone methylation at an Igf2-H19 imprinting epigenetic switch. <i>Molecular Human Reproduction</i> , 2005, 11, 631-640.	2.8	164
7	Putative tumor suppressor miR-145 inhibits colon cancer cell growth by targeting oncogene friend leukemia virus integration 1 gene. <i>Cancer</i> , 2011, 117, 86-95.	4.1	156
8	Pro-inflammatory miR-223 mediates the cross-talk between the IL23 pathway and the intestinal barrier in inflammatory bowel disease. <i>Genome Biology</i> , 2016, 17, 58.	8.8	137
9	Loss of imprinting of IGF2 and H19 in osteosarcoma is accompanied by reciprocal methylation changes of a CTCF-binding site. <i>Human Molecular Genetics</i> , 2003, 12, 535-549.	2.9	132
10	Tissue-specific alternate splicing of human telomerase reverse transcriptase (hTERT) influences telomere lengths during human development. <i>International Journal of Cancer</i> , 2001, 91, 644-649.	5.1	131
11	A novel antisense long noncoding RNA within the IGF1R gene locus is imprinted in hematopoietic malignancies. <i>Nucleic Acids Research</i> , 2014, 42, 9588-9601.	14.5	130
12	Intrachromosomal Looping Is Required for Activation of Endogenous Pluripotency Genes during Reprogramming. <i>Cell Stem Cell</i> , 2013, 13, 30-35.	11.1	120
13	Long noncoding RNA-mediated intrachromosomal interactions promote imprinting at the <i>Kcnq1</i> locus. <i>Journal of Cell Biology</i> , 2014, 204, 61-75.	5.2	118
14	<i>FLI1</i> Exonic Circular RNAs as a Novel Oncogenic Driver to Promote Tumor Metastasis in Small Cell Lung Cancer. <i>Clinical Cancer Research</i> , 2019, 25, 1302-1317.	7.0	113
15	Mitochondrial peptides modulate mitochondrial function during cellular senescence. <i>Aging</i> , 2018, 10, 1239-1256.	3.1	98
16	IGFBP-2 Enhances VEGF Gene Promoter Activity and Consequent Promotion of Angiogenesis by Neuroblastoma Cells. <i>Endocrinology</i> , 2011, 152, 3332-3342.	2.8	97
17	Long noncoding RNAs coordinate functions between mitochondria and the nucleus. <i>Epigenetics and Chromatin</i> , 2017, 10, 41.	3.9	86
18	An intragenic long noncoding RNA interacts epigenetically with the <i>RUNX1</i> promoter and enhancer chromatin DNA in hematopoietic malignancies. <i>International Journal of Cancer</i> , 2014, 135, 2783-2794.	5.1	82

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19	Symmetric and Asymmetric DNA Methylation in the Human IGF2/H19 Imprinted Region. <i>Genomics</i> , 2000, 64, 132-143.	2.9	78
20	A methylated oligonucleotide inhibits IGF2 expression and enhances survival in a model of hepatocellular carcinoma. <i>Journal of Clinical Investigation</i> , 2003, 111, 265-273.	8.2	77
21	Interruption of intrachromosomal looping by CCCTC binding factor decoy proteins abrogates genomic imprinting of human insulin-like growth factor II. <i>Journal of Cell Biology</i> , 2011, 193, 475-487.	5.2	68
22	Promoter-specific Modulation of Insulin-like Growth Factor II Genomic Imprinting by Inhibitors of DNA Methylation. <i>Journal of Biological Chemistry</i> , 1996, 271, 18253-18262.	3.4	66
23	Dissociation of IGF2 and H19 imprinting in human brain. <i>Brain Research</i> , 1998, 810, 1-8.	2.2	66
24	The Histone Code Regulating Expression of the Imprinted Mouse <i>Igf2r</i> Gene. <i>Endocrinology</i> , 2003, 144, 5658-5670.	2.8	66
25	LncRNA Functions as a New Emerging Epigenetic Factor in Determining the Fate of Stem Cells. <i>Frontiers in Genetics</i> , 2020, 11, 277.	2.3	65
26	CTCF Binding at the Insulin-Like Growth Factor-II (IGF2)/H19 Imprinting Control Region Is Insufficient to Regulate IGF2/H19 Expression in Human Tissues. <i>Endocrinology</i> , 2003, 144, 4420-4426.	2.8	63
27	Nuclear-Encoded lncRNA MALAT1 Epigenetically Controls Metabolic Reprogramming in HCC Cells through the Mitophagy Pathway. <i>Molecular Therapy - Nucleic Acids</i> , 2021, 23, 264-276.	5.1	61
28	Tissue-Specific Imprinting of the Mouse Insulin-Like Growth Factor II Receptor Gene Correlates with Differential Allele-Specific DNA Methylation. <i>Molecular Endocrinology</i> , 1998, 12, 220-232.	3.7	59
29	Allele-Specific Histone Acetylation Accompanies Genomic Imprinting of the Insulin-Like Growth Factor II Receptor Gene**Supported by NIH Grant DK-36054 and by the Research Service of the Department of Veterans Affairs.. <i>Endocrinology</i> , 2000, 141, 4428-4435.	2.8	58
30	Long Noncoding RNA HOTAIR as an Independent Prognostic Marker in Cancer: A Meta-Analysis. <i>PLoS ONE</i> , 2014, 9, e105538.	2.5	58
31	A novel orthotopic tumor model to study growth factors and oncogenes in hepatocarcinogenesis. <i>Clinical Cancer Research</i> , 2003, 9, 2719-26.	7.0	57
32	The Role of Histone Acetylation in the Allelic Expression of the Imprinted Human Insulin-like Growth Factor II Gene. <i>Biochemical and Biophysical Research Communications</i> , 1998, 251, 403-408.	2.1	54
33	Human umbilical cord mesenchymal stromal cells rescue mice from acetaminophen-induced acute liver failure. <i>Cytotherapy</i> , 2014, 16, 1207-1219.	0.7	54
34	Epigenetic Targeting of Granulin in Hepatoma Cells by Synthetic CRISPR dCas9 Epi-suppressors. <i>Molecular Therapy - Nucleic Acids</i> , 2018, 11, 23-33.	5.1	52
35	The effects of mitochondria-associated long noncoding RNAs in cancer mitochondria: New players in an old arena. <i>Critical Reviews in Oncology/Hematology</i> , 2018, 131, 76-82.	4.4	51
36	Lack of Reciprocal Genomic Imprinting of Sense and Antisense RNA of Mouse Insulin-like Growth Factor II Receptor in the Central Nervous System1. <i>Biochemical and Biophysical Research Communications</i> , 1999, 257, 604-608.	2.1	46

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37	Modulation of Igf2 Genomic Imprinting in Mice Induced by 5-Azacytidine, an Inhibitor of DNA Methylation. <i>Molecular Endocrinology</i> , 1997, 11, 1891-1898.	3.7	45
38	Epigenetic reprogramming reverses the malignant epigenotype of the MMP/TIMP axis genes in tumor cells. <i>International Journal of Cancer</i> , 2014, 134, 1583-1594.	5.1	45
39	An Imprinted PEG1/MEST Antisense Expressed Predominantly in Human Testis and in Mature Spermatozoa. <i>Journal of Biological Chemistry</i> , 2002, 277, 13518-13527.	3.4	44
40	A Complex Deoxyribonucleic Acid Looping Configuration Associated with the Silencing of the Maternal Igf2 Allele. <i>Molecular Endocrinology</i> , 2008, 22, 1476-1488.	3.7	44
41	Combination of Circulating Tumor Cells with Serum Carcinoembryonic Antigen Enhances Clinical Prediction of Non-Small Cell Lung Cancer. <i>PLoS ONE</i> , 2015, 10, e0126276.	2.5	43
42	A methylated oligonucleotide inhibits IGF2 expression and enhances survival in a model of hepatocellular carcinoma. <i>Journal of Clinical Investigation</i> , 2003, 111, 265-273.	8.2	43
43	In vitro analysis of the proliferative capacity and cytotoxic effects of ex vivo induced natural killer cells, cytokine-induced killer cells, and gamma-delta T cells. <i>BMC Immunology</i> , 2015, 16, 61.	2.2	41
44	Genomic Deletion of an Imprint Maintenance Element Abolishes Imprinting of Both Insulin-like Growth Factor II and H19. <i>Journal of Biological Chemistry</i> , 1997, 272, 20715-20720.	3.4	40
45	Enhanced Therapeutic Efficacy by Simultaneously Targeting Two Genetic Defects in Tumors. <i>Molecular Therapy</i> , 2009, 17, 57-64.	8.2	40
46	Promoter histone H3K27 methylation in the control of IGF2 imprinting in human tumor cell lines. <i>Human Molecular Genetics</i> , 2014, 23, 117-128.	2.9	40
47	Histone deacetylase inhibitor valproic acid promotes the induction of pluripotency in mouse fibroblasts by suppressing reprogramming-induced senescence stress. <i>Experimental Cell Research</i> , 2015, 337, 61-67.	2.6	39
48	Correction of aberrant imprinting of IGF2 in human tumors by nuclear transfer-induced epigenetic reprogramming. <i>EMBO Journal</i> , 2006, 25, 5329-5338.	7.8	37
49	Promotion of the induction of cell pluripotency through metabolic remodeling by thyroid hormone triiodothyronine-activated PI3K/AKT signal pathway. <i>Biomaterials</i> , 2012, 33, 5514-5523.	11.4	37
50	Epigenetic regulation of IGF2/H19 imprinting at CTCF insulator binding sites. <i>Journal of Cellular Biochemistry</i> , 2003, 90, 1038-1055.	2.6	36
51	Aberrant allele-switch imprinting of a novel IGF1R intragenic antisense non-coding RNA in breast cancers. <i>European Journal of Cancer</i> , 2015, 51, 260-270.	2.8	35
52	Targeting the IGF1R Pathway in Breast Cancer Using Antisense lncRNA-Mediated Promoter cis Competition. <i>Molecular Therapy - Nucleic Acids</i> , 2018, 12, 105-117.	5.1	33
53	Mitochondrial DNA Hypomethylation Is a Biomarker Associated with Induced Senescence in Human Fetal Heart Mesenchymal Stem Cells. <i>Stem Cells International</i> , 2017, 2017, 1-12.	2.5	32
54	Transient in vitro epigenetic reprogramming of skin fibroblasts into multipotent cells. <i>Biomaterials</i> , 2010, 31, 2779-2787.	11.4	31

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55	Copper induces cellular senescence in human glioblastoma multiforme cells through downregulation of Bmi-1. <i>Oncology Reports</i> , 2013, 29, 1805-1810.	2.6	31
56	Epigenetic regulation of the taxol resistance-associated gene TRAG-3 in human tumors. <i>Cancer Genetics and Cytogenetics</i> , 2004, 151, 1-13.	1.0	30
57	Atorvastatin exerts its anti-atherosclerotic effects by targeting the receptor for advanced glycation end products. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2011, 1812, 1130-1137.	3.8	28
58	Therapeutic Potential of HGF-Expressing Human Umbilical Cord Mesenchymal Stem Cells in Mice with Acute Liver Failure. <i>International Journal of Hepatology</i> , 2016, 2016, 1-13.	1.1	28
59	Valproic Acid Enhances iPSC Induction From Human Bone Marrow-Derived Cells Through the Suppression of Reprogramming-Induced Senescence. <i>Journal of Cellular Physiology</i> , 2016, 231, 1719-1727.	4.1	28
60	Profiling the long noncoding RNA interaction network in the regulatory elements of target genes by chromatin in situ reverse transcription sequencing. <i>Genome Research</i> , 2019, 29, 1521-1532.	5.5	27
61	Repression of hepatitis B virus (HBV) transgene and HBV-induced liver injury by low protein diet. <i>Oncogene</i> , 1997, 15, 2795-2801.	5.9	26
62	Inhibition of Retinoblastoma In Vitro and In Vivo with Conditionally Replicating Oncolytic Adenovirus H101. , 2010, 51, 2626.		24
63	Loss of insulin-like growth factor II imprinting is a hallmark associated with enhanced chemo/radiotherapy resistance in cancer stem cells. <i>Oncotarget</i> , 2016, 7, 51349-51364.	1.8	24
64	JMJD3 acts in tandem with KLF4 to facilitate reprogramming to pluripotency. <i>Nature Communications</i> , 2020, 11, 5061.	12.8	24
65	CRISPR Cas9-guided chromatin immunoprecipitation identifies miR483 as an epigenetic modulator of IGF2 imprinting in tumors. <i>Oncotarget</i> , 2017, 8, 34177-34190.	1.8	23
66	Genome-wide interaction target profiling reveals a novel Peblr20-eRNA activation pathway to control stem cell pluripotency. <i>Theranostics</i> , 2020, 10, 353-370.	10.0	23
67	Potential of tumor radiotherapy by a radiation-inducible oncolytic and oncoapoptotic adenovirus in cervical cancer xenografts. <i>International Journal of Cancer</i> , 2012, 130, 443-453.	5.1	22
68	Chromatin looping is needed for iPSC induction. <i>Cell Cycle</i> , 2014, 13, 1-2.	2.6	22
69	Hematopoietic recovery of acute radiation syndrome by human superoxide dismutase-expressing umbilical cord mesenchymal stromal cells. <i>Cytotherapy</i> , 2015, 17, 403-417.	0.7	21
70	Friend leukemia virus integration 1 promotes tumorigenesis of small cell lung cancer cells by activating the miR-17-92 pathway. <i>Oncotarget</i> , 2017, 8, 41975-41987.	1.8	21
71	Oplr16 serves as a novel chromatin factor to control stem cell fate by modulating pluripotency-specific chromosomal looping and TET2-mediated DNA demethylation. <i>Nucleic Acids Research</i> , 2020, 48, 3935-3948.	14.5	20
72	Friend leukemia virus integration 1 activates the Rho GTPase pathway and is associated with metastasis in breast cancer. <i>Oncotarget</i> , 2015, 6, 23764-23775.	1.8	20

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73	Allele-Specific Histone Acetylation Accompanies Genomic Imprinting of the Insulin-Like Growth Factor II Receptor Gene. <i>Endocrinology</i> , 2000, 141, 4428-4435.	2.8	19
74	Genome-wide target interactome profiling reveals a novel epigenetic pathway for oncogenic lncRNA in breast cancer. <i>American Journal of Cancer Research</i> , 2019, 9, 714-729.	1.4	19
75	Therapeutic Efficacy by Targeting Correction of Notch1-Induced Aberrants in Uveal Tumors. <i>PLoS ONE</i> , 2012, 7, e44301.	2.5	17
76	Targeted gene suppression by inducing de novo DNA methylation in the gene promoter. <i>Epigenetics and Chromatin</i> , 2014, 7, 20.	3.9	17
77	Restoration of IGF2 imprinting by polycomb repressive complex 2 docking factor SUZ12 in colon cancer cells. <i>Experimental Cell Research</i> , 2015, 338, 214-221.	2.6	17
78	Systematic Correlation Analyses of Circulating Tumor Cells with Clinical Variables and Tumor Markers in Lung Cancer Patients. <i>Journal of Cancer</i> , 2017, 8, 3099-3104.	2.5	17
79	Combined RNA-seq and RAT-seq mapping of long noncoding RNAs in pluripotent reprogramming. <i>Scientific Data</i> , 2018, 5, 180255.	5.3	17
80	Aberrant shuttling of long noncoding RNAs during the mitochondria-nuclear crosstalk in hepatocellular carcinoma cells. <i>American Journal of Cancer Research</i> , 2019, 9, 999-1008.	1.4	17
81	Induced epigenetic modifications of the promoter chromatin silence survivin and inhibit tumor growth. <i>Biochemical and Biophysical Research Communications</i> , 2010, 393, 592-597.	2.1	16
82	Cytokine IL9 Triggers the Pathogenesis of Inflammatory Bowel Disease Through the miR21-CLDN8 Pathway. <i>Inflammatory Bowel Diseases</i> , 2018, 24, 2211-2223.	1.9	16
83	Regulation of telomerase by alternate splicing of human telomerase reverse transcriptase (hTERT) in normal and neoplastic ovary, endometrium and myometrium. <i>International Journal of Cancer</i> , 2000, 85, 330.	5.1	15
84	Gene therapy for colorectal cancer by an oncolytic adenovirus that targets loss of the insulin-like growth factor 2 imprinting system. <i>Molecular Cancer</i> , 2012, 11, 86.	19.2	14
85	Combining Telomerase Reverse Transcriptase Genetic Variant rs2736100 with Epidemiologic Factors in the Prediction of Lung Cancer Susceptibility. <i>Journal of Cancer</i> , 2016, 7, 846-853.	2.5	14
86	The combination of polyalanine expansion mutation and a novel missense substitution in transcription factor FOXL2 leads to different ovarian phenotypes in blepharophimosis-ptosis-epicanthus inversus syndrome (BPES) patients. <i>Human Reproduction</i> , 2012, 27, 3347-3357.	0.9	12
87	Antitumor potential of a synthetic interferon-alpha/PLGF-2 positive charge peptide hybrid molecule in pancreatic cancer cells. <i>Scientific Reports</i> , 2015, 5, 16975.	3.3	12
88	Bioinformatics analyses of differentially expressed genes associated with bisphosphonate-related osteonecrosis of the jaw in patients with multiple myeloma. <i>OncoTargets and Therapy</i> , 2015, 8, 2681.	2.0	12
89	Long noncoding RNA: A resident staff of genomic instability regulation in tumorigenesis. <i>Cancer Letters</i> , 2021, 503, 103-109.	7.2	12
90	Chromatin lncRNA Platr10 controls stem cell pluripotency by coordinating an intrachromosomal regulatory network. <i>Genome Biology</i> , 2021, 22, 233.	8.8	12

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91	Targeted tumor gene therapy based on loss of IGF2 imprinting. <i>Cancer Biology and Therapy</i> , 2010, 10, 290-298.	3.4	11
92	Serum peptidomic profiling identifies a minimal residual disease detection and prognostic biomarker for patients with acute leukemia. <i>Oncology Letters</i> , 2013, 6, 1453-1460.	1.8	11
93	Long noncoding RNAs and their epigenetic function in hematological diseases. <i>Hematological Oncology</i> , 2019, 37, 15-21.	1.7	11
94	Epigenetic suppression of the antitumor cytotoxicity of NK cells by histone deacetylase inhibitor valproic acid. <i>American Journal of Cancer Research</i> , 2016, 6, 600-14.	1.4	11
95	Gene therapy for cancer through adenovirus vector-mediated expression of the Ad5 early region gene 1A based on loss of IGF2 imprinting. <i>Oncology Reports</i> , 2013, 30, 1814-1822.	2.6	10
96	miRâ€³338â€³p inhibits A549 lung cancer cell proliferation and invasion by targeting AKT and Î²â€³catenin signaling pathways. <i>Molecular Medicine Reports</i> , 2019, 20, 33-40.	2.4	10
97	Targeted knockdown of Bcl2 in tumor cells using a synthetic TRAIL 3â€³â€³UTR microRNA. <i>International Journal of Cancer</i> , 2010, 126, 2229-2239.	5.1	9
98	Evaluation of Circulating Tumor Cells in Predicting Therapeutic Response in Small Cell Lung Cancer Patients. <i>Archives of Medical Research</i> , 2016, 47, 454-459.	3.3	9
99	Knockdown of <i>COPS3</i> Inhibits Lung Cancer Tumor Growth in Nude Mice by Blocking Cell Cycle Progression. <i>Journal of Cancer</i> , 2017, 8, 1129-1136.	2.5	9
100	Targeting Jurkat T Lymphocyte Leukemia Cells by an Engineered Interferon-Alpha Hybrid Molecule. <i>Cellular Physiology and Biochemistry</i> , 2017, 42, 519-529.	1.6	8
101	Profiling the epigenetic interplay of lncRNA RUNXOR and oncogenic RUNX1 in breast cancer cells by gene in situ cis-activation. <i>American Journal of Cancer Research</i> , 2019, 9, 1635-1649.	1.4	8
102	Dual Effects of Cellular Immunotherapy in Inhibition of Virus Replication and Prolongation of Survival in HCV-Positive Hepatocellular Carcinoma Patients. <i>Journal of Immunology Research</i> , 2016, 2016, 1-8.	2.2	7
103	Targeted breast cancer therapy by harnessing the inherent blood group antigen immune system. <i>Oncotarget</i> , 2017, 8, 15034-15046.	1.8	7
104	The Nucleus/Mitochondria-Shuttling lncRNAs Function as New Epigenetic Regulators of Mitophagy in Cancer. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 699621.	3.7	7
105	Cancer Stem Cell Marker Musashi-1 rs2522137 Genotype Is Associated with an Increased Risk of Lung Cancer. <i>PLoS ONE</i> , 2014, 9, e95915.	2.5	7
106	<i>Osblr8</i> orchestrates intrachromosomal loop structure required for maintaining stem cell pluripotency. <i>International Journal of Biological Sciences</i> , 2020, 16, 1861-1875.	6.4	6
107	Tissueâ€³specific alternate splicing of human telomerase reverse transcriptase (hTERT) influences telomere lengths during human development. <i>International Journal of Cancer</i> , 2001, 91, 644-649.	5.1	6
108	A Novel Anticancer Therapy That Simultaneously Targets Aberrant p53 and Notch Activities in Tumors. <i>PLoS ONE</i> , 2012, 7, e46627.	2.5	6

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109	A Novel Inherited Mutation in PRKAR1A Abrogates PreRNA Splicing in a Carney Complex Family. Canadian Journal of Cardiology, 2015, 31, 1393-1401.	1.7	5
110	Converting Skin Fibroblasts into Hepatic-like Cells by Transient Programming. Journal of Cellular Biochemistry, 2016, 117, 589-598.	2.6	5
111	Examining Histone Acetylation at Specific Genomic Regions. , 2002, 181, 285-296.		4
112	Inhibition of HIV-1 Viral Infection by an Engineered CRISPR Csy4 RNA Endoribonuclease. PLoS ONE, 2015, 10, e0141335.	2.5	4
113	Pluripotency exit is guided by the <i>PeIn1</i> -mediated disruption of intrachromosomal architecture. Journal of Cell Biology, 2022, 221, .	5.2	3
114	Effect of MALAT1 in the crosstalk between nucleus and mitochondria on mitochondrial reprogramming in hepatocellular carcinoma cells.. Journal of Clinical Oncology, 2019, 37, e14711-e14711.	1.6	2
115	A placental growth factor-positively charged peptide potentiates the antitumor activity of interferon-gamma in human brain glioblastoma U87 cells. American Journal of Cancer Research, 2016, 6, 214-25.	1.4	2
116	Tissue-specific alternate splicing of human telomerase reverse transcriptase (hTERT) influences telomere lengths during human development. , 2001, 91, 644.		1
117	Effect of long noncoding RNA RUNXOR on the epigenetic regulation of RUNX1 in acute myelocytic leukemia.. Journal of Clinical Oncology, 2015, 33, 7018-7018.	1.6	0
118	The antitumor activity of synthetic interferon-alpha in pancreatic cancer cells.. Journal of Clinical Oncology, 2015, 33, e15272-e15272.	1.6	0
119	Effect of friend leukemia virus integration 1 on tumorigenesis of small cell lung cancer cells and activation of the miR-17-92 pathway.. Journal of Clinical Oncology, 2017, 35, e20015-e20015.	1.6	0
120	FLI1 circular RNA as biomarkers for tracking disease progression and as potential therapeutic targets in small cell lung cancer.. Journal of Clinical Oncology, 2018, 36, e20573-e20573.	1.6	0