

Mohammad Obaidul Hoque

List of Publications by Year in descending order

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128
papers

9,312
citations

28274

55
h-index

40979

93
g-index

129
all docs

129
docs citations

129
times ranked

12707
citing authors

#	ARTICLE	IF	CITATIONS
1	Dysfunctional KEAP1â€“NRF2 Interaction in Non-Small-Cell Lung Cancer. <i>PLoS Medicine</i> , 2006, 3, e420.	8.4	894
2	Pharmacologic unmasking of epigenetically silenced tumor suppressor genes in esophageal squamous cell carcinoma. <i>Cancer Cell</i> , 2002, 2, 485-495.	16.8	315
3	Quantitation of Promoter Methylation of Multiple Genes in Urine DNA and Bladder Cancer Detection. <i>Journal of the National Cancer Institute</i> , 2006, 98, 996-1004.	6.3	237
4	A Quantitative Promoter Methylation Profile of Prostate Cancer. <i>Clinical Cancer Research</i> , 2004, 10, 8472-8478.	7.0	234
5	Patient-derived xenografts effectively capture responses to oncology therapy in a heterogeneous cohort of patients with solid tumors. <i>Annals of Oncology</i> , 2017, 28, 2595-2605.	1.2	229
6	Quantitative Methylation-Specific Polymerase Chain Reaction Gene Patterns in Urine Sediment Distinguish Prostate Cancer Patients From Control Subjects. <i>Journal of Clinical Oncology</i> , 2005, 23, 6569-6575.	1.6	227
7	Detection of Aberrant Methylation of Four Genes in Plasma DNA for the Detection of Breast Cancer. <i>Journal of Clinical Oncology</i> , 2006, 24, 4262-4269.	1.6	219
8	The Human MitoChip: A High-Throughput Sequencing Microarray for Mitochondrial Mutation Detection. <i>Genome Research</i> , 2004, 14, 812-819.	5.5	218
9	Quantitative Detection of Promoter Hypermethylation of Multiple Genes in the Tumor, Urine, and Serum DNA of Patients with Renal Cancer. <i>Cancer Research</i> , 2004, 64, 5511-5517.	0.9	218
10	Involvement of aquaporins in colorectal carcinogenesis. <i>Oncogene</i> , 2003, 22, 6699-6703.	5.9	175
11	Detection of Promoter Hypermethylation of Multiple Genes in the Tumor and Bronchoalveolar Lavage of Patients with Lung Cancer. <i>Clinical Cancer Research</i> , 2004, 10, 2284-2288.	7.0	163
12	Evaluation of Promoter Hypermethylation Detection in Body Fluids as a Screening/Diagnosis Tool for Head and Neck Squamous Cell Carcinoma. <i>Clinical Cancer Research</i> , 2008, 14, 97-107.	7.0	163
13	Aquaporin 1 Is Overexpressed in Lung Cancer and Stimulates NIH-3T3 Cell Proliferation and Anchorage-Independent Growth. <i>American Journal of Pathology</i> , 2006, 168, 1345-1353.	3.8	150
14	Quantitative detection of Merkel cell virus in human tissues and possible mode of transmission. <i>International Journal of Cancer</i> , 2010, 126, 2991-2996.	5.1	146
15	The ratio of CD8 to Treg tumor-infiltrating lymphocytes is associated with response to cisplatin-based neoadjuvant chemotherapy in patients with muscle invasive urothelial carcinoma of the bladder. <i>Oncolmmunology</i> , 2016, 5, e1134412.	4.6	135
16	PD-L1 Expression Heterogeneity in Nonâ€“Small Cell Lung Cancer: Defining Criteria for Harmonization between Biopsy Specimens and Whole Sections. <i>Journal of Thoracic Oncology</i> , 2018, 13, 1113-1120.	1.1	135
17	Targeting Cancer Stem Cells: A Strategy for Effective Eradication of Cancer. <i>Cancers</i> , 2019, 11, 732.	3.7	134
18	Genome-Wide Promoter Analysis Uncovers Portions of the Cancer Methylome. <i>Cancer Research</i> , 2008, 68, 2661-2670.	0.9	131

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19	Promoter Hypermethylation as an Independent Prognostic Factor for Relapse in Patients with Prostate Cancer Following Radical Prostatectomy. <i>Clinical Cancer Research</i> , 2005, 11, 8321-8325.	7.0	129
20	Targeted sequencing reveals clonal genetic changes in the progression of early lung neoplasms and paired circulating DNA. <i>Nature Communications</i> , 2015, 6, 8258.	12.8	129
21	Inverse Correlation between Cyclin A1 Hypermethylation and p53 Mutation in Head and Neck Cancer Identified by Reversal of Epigenetic Silencing. <i>Cancer Research</i> , 2004, 64, 5982-5987.	0.9	127
22	An Epigenetic Marker Panel for Detection of Lung Cancer Using Cell-Free Serum DNA. <i>Clinical Cancer Research</i> , 2011, 17, 4494-4503.	7.0	126
23	The TGF β 2-miR200a-MIG6 Pathway Orchestrates the EMT-Associated Kinase Switch That Induces Resistance to EGFR Inhibitors. <i>Cancer Research</i> , 2014, 74, 3995-4005.	0.9	123
24	Quantitative Assessment of Promoter Methylation Profiles in Thyroid Neoplasms. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2005, 90, 4011-4018.	3.6	122
25	High Promoter Methylation Levels of <i>APC</i> Predict Poor Prognosis in Sextant Biopsies from Prostate Cancer Patients. <i>Clinical Cancer Research</i> , 2007, 13, 6122-6129.	7.0	122
26	Genome-wide genetic characterization of bladder cancer: a comparison of high-density single-nucleotide polymorphism arrays and PCR-based microsatellite analysis. <i>Cancer Research</i> , 2003, 63, 2216-22.	0.9	122
27	A time for YAP1: Tumorigenesis, immunosuppression and targeted therapy. <i>International Journal of Cancer</i> , 2018, 143, 2133-2144.	5.1	119
28	Quantitative <i>RARβ2</i> Hypermethylation. <i>Clinical Cancer Research</i> , 2004, 10, 4010-4014.	7.0	117
29	Mitochondrial Cytochrome B Gene Mutation Promotes Tumor Growth in Bladder Cancer. <i>Cancer Research</i> , 2008, 68, 700-706.	0.9	117
30	A Panel of Novel Detection and Prognostic Methylated DNA Markers in Primary Non-Small Cell Lung Cancer and Serum DNA. <i>Clinical Cancer Research</i> , 2017, 23, 7141-7152.	7.0	116
31	Molecular Analysis of Plasma DNA for the Early Detection of Lung Cancer by Quantitative Methylation-Specific PCR. <i>Clinical Cancer Research</i> , 2010, 16, 3463-3472.	7.0	105
32	Assessment of gene promoter hypermethylation for detection of cervical neoplasia. <i>International Journal of Cancer</i> , 2006, 119, 1908-1914.	5.1	97
33	N-Methyl-d-Aspartate Receptor Type 2B Is Epigenetically Inactivated and Exhibits Tumor-Suppressive Activity in Human Esophageal Cancer. <i>Cancer Research</i> , 2006, 66, 3409-3418.	0.9	97
34	Epigenetic Heterogeneity of High-Grade Prostatic Intraepithelial Neoplasia: Clues for Clonal Progression in Prostate Carcinogenesis. <i>Molecular Cancer Research</i> , 2006, 4, 1-8.	3.4	85
35	Role of HGF/c-met system in invasion and metastasis of oral squamous cell carcinoma cells in vitro and its clinical significance. <i>International Journal of Cancer</i> , 2001, 93, 489-496.	5.1	84
36	Detection of Promoter Hypermethylation in Salivary Rinses as a Biomarker for Head and Neck Squamous Cell Carcinoma Surveillance. <i>Clinical Cancer Research</i> , 2011, 17, 4782-4789.	7.0	84

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37	CDC91L1 (PIG-U) is a newly discovered oncogene in human bladder cancer. <i>Nature Medicine</i> , 2004, 10, 374-381.	30.7	79
38	YAP1 and COX2 Coordinately Regulate Urothelial Cancer Stem-like Cells. <i>Cancer Research</i> , 2018, 78, 168-181.	0.9	77
39	PD-L1 Assays 22C3 and SP263 are Not Interchangeable in Non-Small Cell Lung Cancer When Considering Clinically Relevant Cutoffs. <i>American Journal of Surgical Pathology</i> , 2018, 42, 1384-1389.	3.7	77
40	Aquaporin expression in human lymphocytes and dendritic cells. <i>American Journal of Hematology</i> , 2004, 75, 128-133.	4.1	76
41	Oxidized guanine lesions and hOgg1 activity in lung cancer. <i>Oncogene</i> , 2005, 24, 4496-4508.	5.9	76
42	KIF1A and EDNRB are differentially methylated in primary HNSCC and salivary rinses. <i>International Journal of Cancer</i> , 2010, 127, 2351-2359.	5.1	75
43	Changes in CpG Islands Promoter Methylation Patterns during Ductal Breast Carcinoma Progression. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 2694-2700.	2.5	73
44	Aberrant Promoter Methylation of Multiple Genes during Pathogenesis of Bladder Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 2786-2794.	2.5	72
45	Association between Lifestyle Factors and CpG Island Methylation in a Cancer-Free Population. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 2984-2991.	2.5	68
46	Integrated, Genome-Wide Screening for Hypomethylated Oncogenes in Salivary Gland Adenoid Cystic Carcinoma. <i>Clinical Cancer Research</i> , 2011, 17, 4320-4330.	7.0	68
47	MT1G Hypermethylation Is Associated with Higher Tumor Stage in Prostate Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2005, 14, 1274-1278.	2.5	65
48	Midkine induces epithelial-mesenchymal transition through Notch2/Jak2-Stat3 signaling in human keratinocytes. <i>Cell Cycle</i> , 2008, 7, 1613-1622.	2.6	65
49	Pharmacologic Unmasking of Epigenetically Silenced Genes in Breast Cancer. <i>Clinical Cancer Research</i> , 2009, 15, 1184-1191.	7.0	64
50	Immunohistochemical p53 Expression Patterns in Sarcomatoid Carcinomas of the Upper Respiratory Tract. <i>American Journal of Surgical Pathology</i> , 2002, 26, 1024-1031.	3.7	63
51	DNA methylation changes in prostate cancer: current developments and future clinical implementation. <i>Expert Review of Molecular Diagnostics</i> , 2009, 9, 243-257.	3.1	61
52	Clear cell papillary renal cell carcinoma: micro-RNA expression profiling and comparison with clear cell renal cell carcinoma and papillary renal cell carcinoma. <i>Human Pathology</i> , 2014, 45, 1130-1138.	2.0	61
53	Absence of V599E BRAF mutations in desmoplastic melanomas. <i>Cancer</i> , 2005, 103, 788-792.	4.1	60
54	Frequent 14-3-3 Promoter Methylation in Benign and Malignant Prostate Lesions. <i>DNA and Cell Biology</i> , 2005, 24, 264-269.	1.9	60

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55	A survey of methylated candidate tumor suppressor genes in nasopharyngeal carcinoma. <i>International Journal of Cancer</i> , 2011, 128, 1393-1403.	5.1	59
56	Performance of mitochondrial DNA mutations detecting early stage cancer. <i>BMC Cancer</i> , 2008, 8, 285.	2.6	57
57	Immune profiles in primary squamous cell carcinoma of the head and neck. <i>Oral Oncology</i> , 2019, 96, 77-88.	1.5	57
58	OGDHL Is a Modifier of AKT-Dependent Signaling and NF- κ B Function. <i>PLoS ONE</i> , 2012, 7, e48770.	2.5	56
59	Hypermethylation of Cyclin D2 is associated with loss of mRNA expression and tumor development in prostate cancer. <i>Journal of Molecular Medicine</i> , 2006, 84, 911-918.	3.9	54
60	Intravesical BCG Induces CD4+ T-Cell Expansion in an Immune Competent Model of Bladder Cancer. <i>Cancer Immunology Research</i> , 2017, 5, 594-603.	3.4	54
61	\hat{I}^m Np63 \hat{I}^\pm Confers Tumor Cell Resistance to Cisplatin through the AKT1 Transcriptional Regulation. <i>Cancer Research</i> , 2011, 71, 1167-1176.	0.9	51
62	Correlation between BRAF mutation and promoter methylation of TIMP3, RAR \hat{I}^2 and RASSF1A in thyroid cancer. <i>Epigenetics</i> , 2012, 7, 710-719.	2.7	51
63	Involvement of Epigenetics and EMT-Related miRNA in Arsenic-Induced Neoplastic Transformation and Their Potential Clinical Use. <i>Cancer Prevention Research</i> , 2015, 8, 208-221.	1.5	51
64	Patient \hat{I}^c derived xenografts as tools in pharmaceutical development. <i>Clinical Pharmacology and Therapeutics</i> , 2016, 99, 612-621.	4.7	50
65	Expression of programmed cell death ligand 1 in non \hat{I}^c small cell lung cancer: Comparison between cytologic smears, core biopsies, and whole sections using the SP263 assay. <i>Cancer Cytopathology</i> , 2019, 127, 52-61.	2.4	49
66	Tissue Inhibitor of Metalloproteinases-3 Promoter Methylation is an Independent Prognostic Factor for Bladder Cancer. <i>Journal of Urology</i> , 2008, 179, 743-747.	0.4	48
67	GSTP1 Promoter Methylation is Associated with Recurrence in Early Stage Prostate Cancer. <i>Journal of Urology</i> , 2014, 192, 1542-1548.	0.4	48
68	Quantitative hypermethylation of a small panel of genes augments the diagnostic accuracy in fine-needle aspirate washings of breast lesions. <i>Breast Cancer Research and Treatment</i> , 2008, 109, 27-34.	2.5	47
69	High-throughput molecular analysis of urine sediment for the detection of bladder cancer by high-density single-nucleotide polymorphism array. <i>Cancer Research</i> , 2003, 63, 5723-6.	0.9	44
70	LKB1/STK11 Suppresses Cyclooxygenase-2 Induction and Cellular Invasion through PEA3 in Lung Cancer. <i>Cancer Research</i> , 2006, 66, 7870-7879.	0.9	43
71	Identification and Validation of Protein Biomarkers of Response to Neoadjuvant Platinum Chemotherapy in Muscle Invasive Urothelial Carcinoma. <i>PLoS ONE</i> , 2015, 10, e0131245.	2.5	42
72	Epigenetically regulated PAX6 drives cancer cells toward a stem-like state via GLI-SOX2 signaling axis in lung adenocarcinoma. <i>Oncogene</i> , 2018, 37, 5967-5981.	5.9	42

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73	Epigenomics and ovarian carcinoma. <i>Biomarkers in Medicine</i> , 2010, 4, 543-570.	1.4	38
74	Genome-wide methylation profiling and the PI3K-AKT pathway analysis associated with smoking in urothelial cell carcinoma. <i>Cell Cycle</i> , 2013, 12, 1058-1070.	2.6	36
75	Mitochondria in chronic obstructive pulmonary disease and lung cancer: where are we now?. <i>Biomarkers in Medicine</i> , 2017, 11, 475-489.	1.4	36
76	Effect of COVID-19 on Lungs: Focusing on Prospective Malignant Phenotypes. <i>Cancers</i> , 2020, 12, 3822.	3.7	36
77	Forced cytochrome B gene mutation expression induces mitochondrial proliferation and prevents apoptosis in human uroepithelial SV60 cells. <i>International Journal of Cancer</i> , 2009, 125, 2829-2835.	5.1	35
78	Gemcitabine and cisplatin neoadjuvant chemotherapy for muscle-invasive urothelial carcinoma: Predicting response and assessing outcomes. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2015, 33, 204.e1-204.e7.	1.6	34
79	Association of Promoter Methylation of VGF and PGP9.5 with Ovarian Cancer Progression. <i>PLoS ONE</i> , 2013, 8, e70878.	2.5	34
80	PUMA in head and neck cancer. <i>Cancer Letters</i> , 2003, 199, 75-81.	7.2	33
81	Hypermethylation of Genes Detected in Urine from Ghanaian Adults with Bladder Pathology Associated with <i>Schistosoma haematobium</i> Infection. <i>PLoS ONE</i> , 2013, 8, e59089.	2.5	33
82	An integrated genome-wide approach to discover deregulated microRNAs in non-small cell lung cancer: Clinical significance of miR-23b-3p deregulation. <i>Scientific Reports</i> , 2015, 5, 13236.	3.3	32
83	Epigenetic silencing of human T (brachyury homologue) gene in non-small-cell lung cancer. <i>Biochemical and Biophysical Research Communications</i> , 2008, 365, 221-226.	2.1	31
84	Genetic and Epigenetic Analysis of erbB Signaling Pathway Genes in Lung Cancer. <i>Journal of Thoracic Oncology</i> , 2010, 5, 1887-1893.	1.1	31
85	Promoter methylation of MCAM, ER α and ER β in serum of early stage prostate cancer patients. <i>Oncotarget</i> , 2017, 8, 15431-15440.	1.8	31
86	Diabetes and Tumor Formation in Transgenic Mice Expressing Reg I. <i>Biochemical and Biophysical Research Communications</i> , 2000, 278, 368-376.	2.1	30
87	Overexpression of p27Kip1 induces growth arrest and apoptosis in an oral cancer cell line. <i>Oral Oncology</i> , 2002, 38, 730-736.	1.5	28
88	SH3GL2 is frequently deleted in non-small cell lung cancer and downregulates tumor growth by modulating EGFR signaling. <i>Journal of Molecular Medicine</i> , 2013, 91, 381-393.	3.9	28
89	Cigarette smoke induces methylation of the tumor suppressor gene <i>NISCH</i> . <i>Epigenetics</i> , 2013, 8, 383-388.	2.7	28
90	Quantitative Methylation Profiles for Multiple Tumor Suppressor Gene Promoters in Salivary Gland Tumors. <i>PLoS ONE</i> , 2010, 5, e10828.	2.5	27

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91	In silico analysis of pathways activation landscape in oral squamous cell carcinoma and oral leukoplakia. <i>Cell Death Discovery</i> , 2017, 3, 17022.	4.7	27
92	CD24 regulates cancer stem cell (CSC)-like traits and a panel of CSC-related molecules serves as a non-invasive urinary biomarker for the detection of bladder cancer. <i>British Journal of Cancer</i> , 2018, 119, 961-970.	6.4	27
93	PD-L1 expression comparison between primary and relapsed non-small cell lung carcinoma using whole sections and clone SP263. <i>Oncotarget</i> , 2018, 9, 30465-30471.	1.8	26
94	MicroRNA expression profiling of Xp11 renal cell carcinoma. <i>Human Pathology</i> , 2017, 67, 18-29.	2.0	25
95	Genome-wide analysis of genetic alterations in testicular primary seminoma using high resolution single nucleotide polymorphism arrays. <i>Genomics</i> , 2011, 97, 341-349.	2.9	24
96	Arsenic promotes the <sc>COX2/PGE2</sc> axis to increase the malignant stemness properties of urothelial cells. <i>International Journal of Cancer</i> , 2018, 143, 113-126.	5.1	21
97	AKT signaling pathway activated by H1N-1 methylation in non-small cell lung cancer. <i>Tumor Biology</i> , 2012, 33, 307-314.	1.8	19
98	GULP1 regulates the NRF2-KEAP1 signaling axis in urothelial carcinoma. <i>Science Signaling</i> , 2020, 13, .	3.6	19
99	An epigenetic marker panel for recurrence risk prediction of low grade papillary urothelial cell carcinoma (LGPUC) and its potential use for surveillance after transurethral resection using urine. <i>Oncotarget</i> , 2014, 5, 5218-5233.	1.8	19
100	Positive Correlation of Tissue Inhibitor of Metalloproteinase-3 and Death-Associated Protein Kinase Hypermethylation in Head and Neck Squamous Cell Carcinoma. <i>Laryngoscope</i> , 2007, 117, 1376-1380.	2.0	18
101	A<i>IM1</i> promoter hypermethylation as a predictor of decreased risk of recurrence following radical prostatectomy. <i>Prostate</i> , 2012, 72, 1133-1139.	2.3	18
102	Presence of 5-methylcytosine in CpNpG trinucleotides in the human genome. <i>Genomics</i> , 2010, 96, 67-72.	2.9	17
103	Epigenetic inactivation of <i>VGF</i> associated with Urothelial Cell Carcinoma and its potential as a non-invasive biomarker using urine. <i>Oncotarget</i> , 2014, 5, 3350-3361.	1.8	17
104	Integrated transcriptomic and epigenomic analysis of ovarian cancer reveals epigenetically silenced GULP1. <i>Cancer Letters</i> , 2018, 433, 242-251.	7.2	16
105	Identification of EGF as an angiogenic factor present in conditioned medium from human salivary gland adenocarcinoma cell clones with varying degrees of metastatic potential. <i>Cancer Letters</i> , 1994, 84, 189-198.	7.2	15
106	Proteolytic enzymes in salivary extravasation mucocoeles. <i>Journal of Oral Pathology and Medicine</i> , 1995, 24, 299-302.	2.7	15
107	Cyclin A1 expression predicts progression in pT1 urothelial carcinoma of bladder: a tissue microarray study of 149 patients treated by transurethral resection. <i>Histopathology</i> , 2015, 66, 262-269.	2.9	15
108	Significant correlation between matrix metalloproteinase activity and tumor necrosis factor in salivary extravasation mucocoeles. <i>Journal of Oral Pathology and Medicine</i> , 1998, 27, 30-33.	2.7	13

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109	Involvement of miR-518c-5p to Growth and Metastasis in Oral Cancer. PLoS ONE, 2014, 9, e115936.	2.5	12
110	Somatic mitochondrial mutation discovery using ultra-deep sequencing of the mitochondrial genome reveals spatial tumor heterogeneity in head and neck squamous cell carcinoma. Cancer Letters, 2020, 471, 49-60.	7.2	12
111	Increased matrix metalloproteinase-2 activity induced by TGF-beta1 in duct cells of human salivary gland is associated with the development of cyst formation in vivo. Journal of Oral Pathology and Medicine, 1996, 25, 467-473.	2.7	10
112	Therapeutic Targeting of Cancer Stem Cells in Lung, Head and Neck, and Bladder Cancers. Cancers, 2021, 13, 5098.	3.7	9
113	Concurrent Targeting of Potential Cancer Stem Cells Regulating Pathways Sensitizes Lung Adenocarcinoma to Standard Chemotherapy. Molecular Cancer Therapeutics, 2020, 19, 2175-2185.	4.1	8
114	Epigenetic silencing of S100A2 in bladder and head and neck cancers. Oncoscience, 2015, 2, 410-418.	2.2	8
115	High-performance detection of somatic D-loop mutation in urothelial cell carcinoma patients by polymorphism ratio sequencing. Journal of Molecular Medicine, 2016, 94, 1015-1024.	3.9	7
116	A single nucleotide polymorphism in the human PIGK gene associates with low PIGK expression in colorectal cancer patients. International Journal of Oncology, 2012, 41, 1405-1410.	3.3	6
117	Urothelial Carcinoma In Situ of the Bladder: Correlation of CK20 Expression With Adaptive Immune Resistance, Response to BCG Therapy, and Clinical Outcome. Applied Immunohistochemistry and Molecular Morphology, 2021, 29, 127-135.	1.2	5
118	Expression of integrin subunits in normal and malignant human salivary gland cell clones and its regulation by transforming growth factor- β 1. Cancer Letters, 1996, 109, 91-99.	7.2	4
119	Expression of GULP1 in bronchial epithelium is associated with the progression of emphysema in chronic obstructive pulmonary disease. Respiratory Medicine, 2017, 124, 72-78.	2.9	4
120	Abstract 4891: GULP1, a potential tumor suppressor gene in ovarian tumors and its utility as a biomarker. , 2010, , .		3
121	Abstract 4943: Engulfment gene GULP1 is a functional tumor suppressor through influencing TGF- β 2 pathway and is silenced by promoter methylation in urothelial carcinoma. , 2015, , .		2
122	Genetic and Epigenetic Analysis of erbB Signaling Pathway Genes in Lung Cancer: Erratum. Journal of Thoracic Oncology, 2011, 6, 409.	1.1	1
123	Correction: The TGF- β 1-miR200a-Mig6 Pathway Orchestrates the EMT-Associated Kinase Switch That Induces Resistance to EGFR Inhibitors. Cancer Research, 2014, 74, 4950-4950.	0.9	1
124	MicroRNAs, promising biomarkers in the diagnosis of Xp11 translocation RCC-reply. Human Pathology, 2017, 68, 206-207.	2.0	0
125	Differences in the Molecular Characteristics of Bladder Cancer between Smokers and Nonsmokers. European Urology Focus, 2018, 4, 98-99.	3.1	0
126	Development of biomarkers for real precision medicine. Translational Lung Cancer Research, 2018, 7, S228-S231.	2.8	0

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127	GSTP1 as a Potential Marker of Early Chemotherapy Response for Noninvasive Detection. European Urology, 2019, 76, 313-314.	1.9	0
128	Abstract 811: Induction of stem-like cells with malignant properties by chronic exposure of immortalized normal human urothelial cell line to arsenic., 2015, , .		0