Kazuya Shinmura

List of Publications by Year
in descending order

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135 papers	4,899 citations	87888 38 h-index	110387 64 g-index
139 all docs	139 docs citations	139 times ranked	6395 citing authors

#	Article	IF	Citations
1	Genetic polymorphisms and alternative splicing of the hOGG1 gene, that is involved in the repair of 8-hydroxyguanine in damaged DNA. Oncogene, 1998, 16, 3219-3225.	5.9	408
2	Cloning of a human homolog of the yeast OGG1 gene that is involved in the repair of oxidative DNA damage. Oncogene, 1997, 14, 2857-2861.	5.9	249
3	EML4-ALK fusion transcripts, but no NPM-, TPM3-, CLTC-, ATIC-, or TFG-ALK fusion transcripts, in non-small cell lung carcinomas. Lung Cancer, 2008, 61, 163-169.	2.0	159
4	Prognostic impact of CD73 and A2A adenosine receptor expression in non-small-cell lung cancer. Oncotarget, 2017, 8, 8738-8751.	1.8	129
5	TheOGG1Gene Encodes a Repair Enzyme for Oxidatively Damaged DNA and Is Involved in Human Carcinogenesis. Antioxidants and Redox Signaling, 2001, 3, 597-609.	5.4	127
6	EPHA2/EFNA1 expression in human gastric cancer. Cancer Science, 2005, 96, 42-47.	3.9	118
7	Familial gastric cancer: clinicopathological characteristics, RER phenotype and germline p53 and E-cadherin mutations. Carcinogenesis, 1999, 20, 1127-1131.	2.8	115
8	Downregulation of EphA7 by hypermethylation in colorectal cancer. Oncogene, 2005, 24, 5637-5647.	5.9	100
9	Clinical significance of PD-L1 and PD-L2 copy number gains in non-small-cell lung cancer. Oncotarget, 2016, 7, 32113-32128.	1.8	100
10	FrequentPTPRK-RSPO3fusions andRNF43mutations in colorectal traditional serrated adenoma. Journal of Pathology, 2016, 239, 133-138.	4.5	99
11	Infrequent Mutations of thehOGG1Gene, That Is Involved in the Excision of 8-Hydroxyguanine in Damaged DNA, in Human Gastric Cancer. Japanese Journal of Cancer Research, 1998, 89, 825-828.	1.7	96
12	Accumulated phosphatidylcholine (16:0/16:1) in human colorectal cancer; possible involvement of <scp>LPCAT</scp>4. Cancer Science, 2013, 104, 1295-1302.	3.9	96
13	E-cadherin gene variants in gastric cancer families whose probands are diagnosed with diffuse gastric cancer. International Journal of Cancer, 2002, 101, 434-441.	5.1	91
14	Tiam1 mediates neurite outgrowth induced by ephrin-B1 and EphA2. EMBO Journal, 2004, 23, 1075-1088.	7.8	87
15	Direct evidence for the role of centrosomally localized p53 in the regulation of centrosome duplication. Oncogene, 2007, 26, 2939-2944.	5.9	86
16	Inactivating mutations of the human base excision repair gene NEIL1 in gastric cancer. Carcinogenesis, 2004, 25, 2311-2317.	2.8	85
17	Adenine excisional repair function of MYH protein on the adenine:8-hydroxyguanine base pair in double-stranded DNA. Nucleic Acids Research, 2000, 28, 4912-4918.	14.5	79
18	Sumoylation of nucleophosmin/B23 regulates its subcellular localization, mediating cell proliferation and survival. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 9679-9684.	7.1	77

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19	8-hydroxyguanine (7,8-dihydro-8-oxoguanine) DNA glycosylase and AP lyase activities of hOGG1 protein and their substrate specificity. Mutation Research DNA Repair, 1997, 385, 75-82.	3.7	76
20	Human Sgo1 downregulation leads to chromosomal instability in colorectal cancer. Gut, 2009, 58, 249-260.	12.1	72
21	Detection of Lipid Peroxidation-Induced DNA Adducts Caused by 4-Oxo-2(E)-nonenal and 4-Oxo-2(E)-hexenal in Human Autopsy Tissues. Chemical Research in Toxicology, 2010, 23, 1442-1448.	3.3	67
22	EphA2 Up-regulation induced by deoxycholic acid in human colon carcinoma cells, an involvement of extracellular signal-regulated kinase and p53-independence. Journal of Cancer Research and Clinical Oncology, 2003, 129, 703-708.	2.5	61
23	Utility of Scanning Electron Microscopy Elemental Analysis Using the â€˜NanoSuitâ€™ Correlative Light and Electron Microscopy Method in the Diagnosis of Lanthanum Phosphate Deposition in the Esophagogastroduodenal Mucosa. Diagnostics, 2020, 10, 1.	2.6	57
24	Enhancement of OGG1 protein AP lyase activity by increase of APEX protein. Mutation Research DNA Repair, 2001, 486, 31-40.	3.7	56
25	Lipid peroxidation-induced DNA adducts in human gastric mucosa. Carcinogenesis, 2013, 34, 121-127.	2.8	56
26	A novel splice-site variant of the base excision repair gene MYH is associated with production of an aberrant mRNA transcript encoding a truncated MYH protein not localized in the nucleus. Carcinogenesis, 2004, 25, 1859-1866.	2.8	53
27	RSPO fusion transcripts in colorectal cancer in Japanese population. Molecular Biology Reports, 2014, 41, 5375-5384.	2.3	53
28	Thr199phosphorylation targets nucleophosmin to nuclear speckles and represses pre-mRNA processing. FEBS Letters, 2006, 580, 399-409.	2.8	52
29	Characterization of centrosomal association of nucleophosmin/B23 linked to Crm1 activity. FEBS Letters, 2005, 579, 6621-6634.	2.8	51
30	PLK4 overexpression and its effect on centrosome regulation and chromosome stability in human gastric cancer. Molecular Biology Reports, 2014, 41, 6635-6644.	2.3	51
31	YTHDF1 and YTHDF2 are associated with better patient survival and an inflamed tumor-immune microenvironment in nonâ€“small-cell lung cancer. OncoImmunology, 2021, 10, 1962656.	4.6	51
32	Genomic structure and chromosomal localization of the mouse Ogg1 gene that is involved in the repair of 8-hydroxyguanine in DNA damage. Mammalian Genome, 1998, 9, 32-37.	2.2	48
33	Localization of Menkes gene expression in the mouse brain; its association with neurological manifestations in Menkes model mice. Acta Neuropathologica, 1996, 91, 482-488.	7.7	45
34	Reduced expression of MUTYH with suppressive activity against mutations caused by 8â€hydroxyguanine is a novel predictor of a poor prognosis in human gastric cancer. Journal of Pathology, 2011, 225, 414-423.	4.5	45
35	Ile-Leu Substitution (I415L) in Germline E-cadherin Gene (CDH1) in Japanese Familial Gastric Cancer. Japanese Journal of Clinical Oncology, 2003, 33, 17-20.	1.3	44
36	Association between genetic polymorphisms of the base excision repair gene MUTYH and increased colorectal cancer risk in a Japanese population. Cancer Science, 2008, 99, 355-360.	3.9	42

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37	Germline alterations in the CDH1 gene in familial gastric cancer in the Japanese population. Cancer Science, 2011, 102, 1782-1788.	3.9	42
38	Clinicopathological and Survival Analysis of Japanese Patients with Resected Non-Small-Cell Lung Cancer Harboring NKX2-1, SETDB1, MET, HER2, SOX2, FGFR1, or PIK3CA Gene Amplification. Journal of Thoracic Oncology, 2015, 10, 1590-1600.	1.1	40
39	A single nucleotide polymorphism at the splice donor site of the humanMYHbase excision repair gene results in reduced translation efficiency of its transcripts. Genes To Cells, 2002, 7, 461-474.	1.2	39
40	Abnormal Expressions of DNA Glycosylase Genes NEIL1, NEIL2, and NEIL3 Are Associated with Somatic Mutation Loads in Human Cancer. Oxidative Medicine and Cellular Longevity, 2016, 2016, 1-10.	4.0	39
41	Somatic mutations and single nucleotide polymorphisms of base excision repair genes involved in the repair of 8-hydroxyguanine in damaged DNA. Cancer Letters, 2001, 166, 65-69.	7.2	38
42	OGG1 protein suppresses G:C->T:A mutation in a shuttle vector containing 8-hydroxyguanine in human cells. Carcinogenesis, 2001, 22, 1355-1362.	2.8	38
43	Induction of centrosome amplification and chromosome instability in p53â€deficient lung cancer cells exposed to benzo[a]pyrene diol epoxide (B[a]PDE). Journal of Pathology, 2008, 216, 365-374.	4.5	37
44	Adenine DNA glycosylase activity of 14 Human MutY homolog (MUTYH) variant proteins found in patients with colorectal polyposis and cancer. Human Mutation, 2010, 31, E1861-E1874.	2.5	37
45	Aberrant Expression and Mutation-Inducing Activity of AID in Human Lung Cancer. Annals of Surgical Oncology, 2011, 18, 2084-2092.	1.5	36
46	Robust quantitative assessments of cytosine modifications and changes in the expressions of related enzymes in gastric cancer. Gastric Cancer, 2015, 18, 516-525.	5.3	36
47	Microsatellite Instability and K-ras Mutations in Gastric Adenomas, with Reference to Associated Gastric Cancers. Cancer Detection and Prevention, 1999, 23, 204-214.	2.1	36
48	Detection of kinase amplifications in gastric cancer archives using fluorescence in situ hybridization. Pathology International, 2012, 62, 477-484.	1.3	35
49	Heterogeneity analysis of PD-L1 expression and copy number status in EBUS-TBNA biopsy specimens of non-small cell lung cancer: Comparative assessment of primary and metastatic sites. Lung Cancer, 2019, 134, 202-209.	2.0	35
50	RER phenotype and its associated mutations in familial gastric cancer. Carcinogenesis, 1998, 19, 247-251.	2.8	34
51	Suppressive activities of OGG1 and MYH proteins against G:C to T:A mutations caused by 8-hydroxyguanine but not by benzo[a]pyrene diol epoxide in human cells in vivo. Carcinogenesis, 2003, 24, 1031-1037.	2.8	33
52	Identification and characterization of a novel germ line p53 mutation in familial gastric cancer in the Japanese population. Carcinogenesis, 2007, 28, 2013-2018.	2.8	33
53	Chromogenic in situ hybridization (CISH) to detect HER2 gene amplification in breast and gastric cancer: Comparison with immunohistochemistry (IHC) and fluorescence in situ hybridization (FISH). Pathology International, 2012, 62, 728-734.	1.3	33
54	A novel tumor-derived SGOL1 variant causes abnormal mitosis and unstable chromatid cohesion. Oncogene, 2011, 30, 4453-4463.	5.9	32

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55	Frequent co-occurrence of mutator phenotype in synchronous, independent multiple cancers of the stomach. Carcinogenesis, 1995, 16, 2989-2993.	2.8	31
56	Altered expression of the human base excision repair gene NTH1 in gastric cancer. Carcinogenesis, 2009, 30, 1345-1352.	2.8	31
57	Fluorescence in situ hybridization analysis with a tissue microarray: â€˜FISH and chipsâ€™ analysis of pathology archives. Pathology International, 2010, 60, 543-550.	1.3	29
58	TNK2 gene amplification is a novel predictor of a poor prognosis in patients with gastric cancer. Journal of Surgical Oncology, 2014, 109, 189-197.	1.7	28
59	Identification and association study with lung cancer for novel insertion polymorphisms of human endogenous retrovirus. Carcinogenesis, 2013, 34, 2531-2538.	2.8	27
60	SGOL1 variant B induces abnormal mitosis and resistance to taxane in non-small cell lung cancers. Scientific Reports, 2013, 3, 3012.	3.3	26
61	Association betweenCDH1haplotypes and gastric cancer risk in a Japanese population. Scandinavian Journal of Gastroenterology, 2007, 42, 1479-1485.	1.5	23
62	Siteâ€specific methylation patterns of the GAL and GALR1/2 genes in head and neck cancer: Potential utility as biomarkers for prognosis. Molecular Carcinogenesis, 2017, 56, 1107-1116.	2.7	23
63	Splice-site genetic polymorphism of the human kallikrein 12 (KLK12) gene correlates with no substantial expression of KLK12 protein having serine protease activity. Human Mutation, 2004, 24, 273-274.	2.5	22
64	CLCA2 as a Novel Immunohistochemical Marker for Differential Diagnosis of Squamous Cell Carcinoma from Adenocarcinoma of the Lung. Disease Markers, 2014, 2014, 1-11.	1.3	22
65	OGG1, MYH and MTH1 gene variants identified in gastric cancer patients exhibiting both 8-hydroxy-2â€²-deoxyguanosine accumulation and low inflammatory cell infiltration in their gastric mucosa. Journal of Genetics, 2008, 87, 181-186.	0.7	20
66	The CRKL gene encoding an adaptor protein is amplified, overexpressed, and a possible therapeutic target in gastric cancer. Journal of Translational Medicine, 2012, 10, 97.	4.4	20
67	Elucidation of the relationships of MET protein expression and gene copy number status with PD-L1 expression and the immune microenvironment in non-small cell lung cancer. Lung Cancer, 2020, 141, 21-31.	2.0	20
68	CD74-ROS1 fusion transcripts in resected non-small cell lung carcinoma. Oncology Reports, 2013, 30, 1675-1680.	2.6	19
69	CD200 and CD200R1 are differentially expressed and have differential prognostic roles in non-small cell lung cancer. OncoImmunology, 2020, 9, 1746554.	4.6	19
70	Identification and characterization of a novel germline p53 mutation in a patient with glioblastoma and colon cancer. International Journal of Cancer, 2009, 125, 973-976.	5.1	18
71	POLQ Overexpression Is Associated with an Increased Somatic Mutation Load and PLK4 Overexpression in Lung Adenocarcinoma. Cancers, 2019, 11, 722.	3.7	18
72	Genes encoding neuropeptide receptors are epigenetic markers in patients with head and neck cancer: a site-specific analysis. Oncotarget, 2017, 8, 76318-76328.	1.8	18

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73	Suppression of hydroxyurea-induced centrosome amplification by NORE1A and down-regulation of NORE1A mRNA expression in non-small cell lung carcinoma. Lung Cancer, 2011, 71, 19-27.	2.0	17
74	WDR62 overexpression is associated with a poor prognosis in patients with lung adenocarcinoma. Molecular Carcinogenesis, 2017, 56, 1984-1991.	2.7	17
75	Mxi1Mutations in Human Neurofibrosarcomas. Japanese Journal of Cancer Research, 1999, 90, 740-746.	1.7	16
76	Absence of germline mono-allelic promoter hypermethylation of the CDH1 gene in gastric cancer patients. Molecular Cancer, 2009, 8, 63.	19.2	16
77	Identification of 5 novel germline APC mutations and characterization of clinical phenotypes in Japanese patients with classical and attenuated familial adenomatous polyposis. BMC Research Notes, 2010, 3, 305.	1.4	16
78	Association between dopamine beta hydroxylase rs5320 polymorphism and smoking behaviour in elderly Japanese. Journal of Human Genetics, 2012, 57, 385-390.	2.3	16
79	Absence of GermlineCHK2Mutations in Familial Gastric Cancer. Japanese Journal of Cancer Research, 2000, 91, 875-879.	1.7	15
80	Association between neuropeptide Y receptor 2 polymorphism and the smoking behavior of elderly Japanese. Journal of Human Genetics, 2010, 55, 755-760.	2.3	15
81	Anaplastic changes of diffuse leptomeningeal glioneuronal tumor with polar spongioblastoma pattern. Brain Tumor Pathology, 2018, 35, 209-216.	1.7	15
82	Mutational Analyses of Multiple Target Genes in Histologically Heterogeneous Gastric Cancer with Microsatellite Instability. Japanese Journal of Cancer Research, 1998, 89, 1284-1291.	1.7	14
83	Inverse relationship between the length of the EGFR CA repeat polymorphism in lung carcinoma and protein expression of EGFR in the carcinoma. Journal of Surgical Oncology, 2008, 98, 457-461.	1.7	14
84	Neoadjuvant imatinib treatment and laparoscopic anus-preserving surgery for a large gastrointestinal stromal tumor of the rectum. World Journal of Surgical Oncology, 2016, 14, 68.	1.9	14
85	Genetic susceptibility to lung cancer. Frontiers in Bioscience - Scholar, 2011, S3, 1463.	2.1	14
86	Visualization of phosphatidylcholine (16:0/16:0) in type <scp>II</scp> alveolar epithelial cells in the human lung using imaging mass spectrometry. Pathology International, 2013, 63, 195-200.	1.3	13
87	BSND and ATP6V1G3. Medicine (United States), 2015, 94, e989.	1.0	13
88	SASS6 overexpression is associated with mitotic chromosomal abnormalities and a poor prognosis in patients with colorectal cancer. Oncology Reports, 2015, 34, 727-738.	2.6	13
89	Characterization of Vâ€set and immunoglobulin domain containing 1 exerting a tumor suppressor function in gastric, lung, and esophageal cancer cells. Cancer Science, 2017, 108, 1701-1714.	3.9	13
90	Three novelNEIL1promoter polymorphisms in gastric cancer patients. World Journal of Gastrointestinal Oncology, 2010, 2, 117.	2.0	13

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91	m6A demethylase ALKBH5 promotes tumor cell proliferation by destabilizing IGF2BPs target genes and worsens the prognosis of patients with non-small-cell lung cancer. Cancer Gene Therapy, 2022, 29, 1355-1372.	4.6	13
92	Causation of Borrmann type 4 gastric cancer: heritable factors or environmental factors?. Gastric Cancer, 2003, 6, 17-23.	5.3	12
93	A novel somatic FGFR3 mutation in primary lung cancer. Oncology Reports, 2014, 31, 1219-1224.	2.6	12
94	Distinct prognostic roles and heterogeneity of TTF1 copy number and <scp>TTF</scp>1 protein expression in nonâ€small cell lung cancer. Genes Chromosomes and Cancer, 2017, 56, 570-581.	2.8	12
95	Association between neurexin 1 (NRXN1) polymorphisms and the smoking behavior of elderly Japanese. Psychiatric Genetics, 2010, 20, 135-136.	1.1	12
96	Secreted form of EphA7 in lung cancer. International Journal of Oncology, 2010, 36, 635-40.	3.9	12
97	Germline p53 Mutation in a Patient with Multiple Primary Cancers. Japanese Journal of Clinical Oncology, 2001, 31, 349-351.	1.3	11
98	EML4-ALK fusion transcripts in immunohistochemically ALK-positive non-small cell lung carcinomas. Experimental and Therapeutic Medicine, 2010, 1, 271-275.	1.8	11
99	Human DNA glycosylase enzyme TDG repairs thymine mispaired with exocyclic etheno-DNA adducts. Free Radical Biology and Medicine, 2014, 76, 136-146.	2.9	11
100	Defective repair capacity of variant proteins of the DNA glycosylase NTHL1 for 5-hydroxyuracil, an oxidation product of cytosine. Free Radical Biology and Medicine, 2019, 131, 264-273.	2.9	11
101	Impaired 8-Hydroxyguanine Repair Activity of MUTYH Variant p.Arg109Trp Found in a Japanese Patient with Early-Onset Colorectal Cancer. Oxidative Medicine and Cellular Longevity, 2014, 2014, 1-12.	4.0	10
102	Novel roles for LIX1L in promoting cancer cell proliferation through ROS1-mediated LIX1L phosphorylation. Scientific Reports, 2015, 5, 13474.	3.3	10
103	CD44-SLC1A2 Fusion Transcripts in Primary Colorectal Cancer. Pathology and Oncology Research, 2015, 21, 759-764.	1.9	9
104	Reduced expression of the DNA glycosylase gene MUTYH is associated with an increased number of somatic mutations via a reduction in the DNA repair capacity in prostate adenocarcinoma. Molecular Carcinogenesis, 2017, 56, 781-788.	2.7	9
105	Promoter methylation of galanin receptors as epigenetic biomarkers for head and neck squamous cell carcinomas. Expert Review of Molecular Diagnostics, 2019, 19, 137-148.	3.1	9
106	Effect of splice-site polymorphisms of theTMPRSS4, NPHP4 andORCTL4 genes on their mRNA expression. Journal of Genetics, 2005, 84, 131-136.	0.7	8
107	Fluorescence-labeled Methylation-sensitive Amplified Fragment Length Polymorphism (FL-MS-AFLP) Analysis for Quantitative Determination of DNA Methylation and Demethylation Status. Japanese Journal of Clinical Oncology, 2008, 38, 317-322.	1.3	8
108	Identification and characterization of primary ciliaâ€positive salivary gland tumours exhibiting basaloid/myoepithelial differentiation. Journal of Pathology, 2021, 254, 519-530.	4.5	8

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109	Secreted form of EphA7 in lung cancer. International Journal of Oncology, 2010, 36, .	3.3	7
110	A novel APC mosaicism in a patient with familial adenomatous polyposis. Human Genome Variation, 2015, 2, 15057.	0.7	7
111	NEIL1 p.Gln282Stop variant is predominantly localized in the cytoplasm and exhibits reduced activity in suppressing mutations. Gene, 2015, 571, 33-42.	2.2	7
112	Diagnosis of Ion-Exchange Resin Depositions in Paraffin Sections Using Corrective Light and Electron Microscopy-NanoSuit Method. Diagnostics, 2021, 11, 1193.	2.6	7
113	Bilateral breast tumors, malignant phyllodes tumor and invasive lobular carcinoma in a 46, XX/46, XY mosaic female with family history of breast cancer*. Pathology International, 1997, 47, 147-154.	1.3	6
114	Effect of boron neutron capture therapy for recurrent anaplastic meningioma: an autopsy case report. Brain Tumor Pathology, 2015, 32, 61-65.	1.7	6
115	BSND is a Novel Immunohistochemical Marker for Oncocytic Salivary Gland Tumors. Pathology and Oncology Research, 2018, 24, 439-444.	1.9	6
116	Genes Located on 18q23 Are Epigenetic Markers and Have Prognostic Significance for Patients with Head and Neck Cancer. Cancers, 2019, 11, 401.	3.7	6
117	EPH-EPHRIN in human gastrointestinal cancers. World Journal of Gastrointestinal Oncology, 2010, 2, 421.	2.0	6
118	Mediastinal mature teratoma with complete gastrointestinal and bronchial walls. Respirology Case Reports, 2015, 3, 89-91.	0.6	5
119	Clinical analysis of false-negative fine needle aspiration cytology of head and neck cancers. Postgraduate Medicine, 2019, 131, 151-155.	2.0	5
120	Mutation Spectrum Induced by 8-Bromoguanine, a Base Damaged by Reactive Brominating Species, in Human Cells. Oxidative Medicine and Cellular Longevity, 2017, 2017, 1-11.	4.0	4
121	Mismatch repair proteins immunohistochemical null phenotype in colon medullary carcinoma. Clinical Journal of Gastroenterology, 2021, 14, 1448-1452.	0.8	4
122	Impaired suppressive activities of human MUTYH variant proteins against oxidative mutagenesis. World Journal of Gastroenterology, 2012, 18, 6935.	3.3	4
123	Tuberous sclerosis patient with neuroendocrine carcinoma of the esophagogastric junction: A case report. World Journal of Gastroenterology, 2020, 26, 7263-7271.	3.3	4
124	Role of Base Excision Repair Enzyme MUTYH in the Repair of 8-Hydroxyguanine and MUTYH-Associated Polyposis (MAP). Hereditary Genetics: Current Research, 2012, 01, .	0.1	3
125	Functional Evaluation of Nine Missense-Type Variants of the Human DNA Glycosylase Enzyme MUTYH in the Japanese Population. Human Mutation, 2016, 37, 350-353.	2.5	3
126	Segregation Analysis of Gastric Cancer in a Japanese Population. International Journal of Human Genetics, 2001, 1, 263-270.	0.1	2

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127	Non-Mucinous Lepidic Predominant Adenocarcinoma Presenting with Extensive Aerogenous Spread. Rare Tumors, 2016, 8, 169-172.	0.6	2
128	Centrosome Abnormality and Human Lung Cancer. , 0, , .		2
129	Characterization of adenocarcinoma of the lung in a familial adenomatous polyposis patient. Pathology International, 2008, 58, 706-712.	1.3	1
130	Mesothelial/monocytic incidental cardiac excrescence with bone marrow. Pathology International, 2018, 68, 436-438.	1.3	1
131	Abstract 1601: Human colorectal cancer marker PC(16:0/16:1) induces cell growth by activating Akt and Erk pathways. Cancer Research, 2014, 74, 1601-1601.	0.9	1
132	Effect of exogenous MSH6 and POLD1 expression on the mutation rate of the HPRT locus in a human colon cancer cell line with mutator phenotype, DLD-1. International Journal of Oncology, 2004, 24, 697.	3.3	0
133	A Case of Encephalocele in the Temporal Bone Repeating Intracranial Infection. Practica Otologica, 2021, 114, 347-352.	0.0	0
134	Abstract 2131: m6A demethylase ALKBH5 promote tumor growth through IGF2BPs' recognition of m6A modified CDKN1A in non-small lung cancer. , 2021, , .		0
135	Glioblastoma: Germline Mutation of TP53. , 2011, , 31-38.		0