

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	m6A demethylase ALKBH5 promotes tumor cell proliferation by destabilizing IGF2BPs target genes and worsens the prognosis of patients with non-small-cell lung cancer. <i>Cancer Gene Therapy</i> , 2022, 29, 1355-1372.	4.6	13
2	YTHDF1 and YTHDF2 are associated with better patient survival and an inflamed tumor-immune microenvironment in non-small-cell lung cancer. <i>Oncolmmunology</i> , 2021, 10, 1962656.	4.6	51
3	A Case of Encephalocele in the Temporal Bone Repeating Intracranial Infection. <i>Practica Otologica</i> , 2021, 114, 347-352.	0.0	0
4	Identification and characterization of primary cilia-positive salivary gland tumours exhibiting basaloid/myoepithelial differentiation. <i>Journal of Pathology</i> , 2021, 254, 519-530.	4.5	8
5	Diagnosis of Ion-Exchange Resin Depositions in Paraffin Sections Using Corrective Light and Electron Microscopy-NanoSuit Method. <i>Diagnostics</i> , 2021, 11, 1193.	2.6	7
6	Abstract 2131: m6A demethylase ALKBH5 promote tumor growth through IGF2BPs' recognition of m6A modified CDKN1A in non-small lung cancer. , 2021, , .		0
7	Mismatch repair proteins immunohistochemical null phenotype in colon medullary carcinoma. <i>Clinical Journal of Gastroenterology</i> , 2021, 14, 1448-1452.	0.8	4
8	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. <i>Lung Cancer</i> , 2020, 141, 21-31.	2.0	20
9	Utility of Scanning Electron Microscopy Elemental Analysis Using the NanoSuit™ Correlative Light and Electron Microscopy Method in the Diagnosis of Lanthanum Phosphate Deposition in the Esophagogastroduodenal Mucosa. <i>Diagnostics</i> , 2020, 10, 1.	2.6	57
10	CD200 and CD200R1 are differentially expressed and have differential prognostic roles in non-small cell lung cancer. <i>Oncolmmunology</i> , 2020, 9, 1746554.	4.6	19
11	Tuberous sclerosis patient with neuroendocrine carcinoma of the esophagogastric junction: A case report. <i>World Journal of Gastroenterology</i> , 2020, 26, 7263-7271.	3.3	4
12	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. <i>Lung Cancer</i> , 2019, 134, 202-209.	2.0	35
13	POLQ Overexpression Is Associated with an Increased Somatic Mutation Load and PLK4 Overexpression in Lung Adenocarcinoma. <i>Cancers</i> , 2019, 11, 722.	3.7	18
14	Genes Located on 18q23 Are Epigenetic Markers and Have Prognostic Significance for Patients with Head and Neck Cancer. <i>Cancers</i> , 2019, 11, 401.	3.7	6
15	Clinical analysis of false-negative fine needle aspiration cytology of head and neck cancers. <i>Postgraduate Medicine</i> , 2019, 131, 151-155.	2.0	5
16	Defective repair capacity of variant proteins of the DNA glycosylase NTHL1 for 5-hydroxyuracil, an oxidation product of cytosine. <i>Free Radical Biology and Medicine</i> , 2019, 131, 264-273.	2.9	11
17	Promoter methylation of galanin receptors as epigenetic biomarkers for head and neck squamous cell carcinomas. <i>Expert Review of Molecular Diagnostics</i> , 2019, 19, 137-148.	3.1	9
18	Mesothelial/monocytic incidental cardiac excrescence with bone marrow. <i>Pathology International</i> , 2018, 68, 436-438.	1.3	1

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19	BSND is a Novel Immunohistochemical Marker for Oncocytic Salivary Gland Tumors. <i>Pathology and Oncology Research</i> , 2018, 24, 439-444.	1.9	6
20	Anaplastic changes of diffuse leptomeningeal glioneuronal tumor with polar spongioblastoma pattern. <i>Brain Tumor Pathology</i> , 2018, 35, 209-216.	1.7	15
21	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. <i>Molecular Carcinogenesis</i> , 2017, 56, 781-788.	2.7	9
22	WDR62 overexpression is associated with a poor prognosis in patients with lung adenocarcinoma. <i>Molecular Carcinogenesis</i> , 2017, 56, 1984-1991.	2.7	17
23	Distinct prognostic roles and heterogeneity of <i>TTF1</i> copy number and <i>TTF1</i> protein expression in non-small cell lung cancer. <i>Genes Chromosomes and Cancer</i> , 2017, 56, 570-581.	2.8	12
24	Characterization of <i>Veset</i> and immunoglobulin domain containing 1 exerting a tumor suppressor function in gastric, lung, and esophageal cancer cells. <i>Cancer Science</i> , 2017, 108, 1701-1714.	3.9	13
25	Site-specific methylation patterns of the <i>GAL</i> and <i>GALR1/2</i> genes in head and neck cancer: Potential utility as biomarkers for prognosis. <i>Molecular Carcinogenesis</i> , 2017, 56, 1107-1116.	2.7	23
26	Mutation Spectrum Induced by 8-Bromoguanine, a Base Damaged by Reactive Brominating Species, in Human Cells. <i>Oxidative Medicine and Cellular Longevity</i> , 2017, 2017, 1-11.	4.0	4
27	Prognostic impact of CD73 and A2A adenosine receptor expression in non-small-cell lung cancer. <i>Oncotarget</i> , 2017, 8, 8738-8751.	1.8	129
28	Genes encoding neuropeptide receptors are epigenetic markers in patients with head and neck cancer: a site-specific analysis. <i>Oncotarget</i> , 2017, 8, 76318-76328.	1.8	18
29	Clinical significance of <i>PD-L1</i> and <i>PD-L2</i> copy number gains in non-small-cell lung cancer. <i>Oncotarget</i> , 2016, 7, 32113-32128.	1.8	100
30	Non-Mucinous Lepidic Predominant Adenocarcinoma Presenting with Extensive Aerogenous Spread. <i>Rare Tumors</i> , 2016, 8, 169-172.	0.6	2
31	Abnormal Expressions of DNA Glycosylase Genes NEIL1, NEIL2, and NEIL3 Are Associated with Somatic Mutation Loads in Human Cancer. <i>Oxidative Medicine and Cellular Longevity</i> , 2016, 2016, 1-10.	4.0	39
32	Functional Evaluation of Nine Missense-Type Variants of the Human DNA Glycosylase Enzyme MUTYH in the Japanese Population. <i>Human Mutation</i> , 2016, 37, 350-353.	2.5	3
33	Frequent <i>PTPRK-RSPO3</i> fusions and <i>RNF43</i> mutations in colorectal traditional serrated adenoma. <i>Journal of Pathology</i> , 2016, 239, 133-138.	4.5	99
34	Neoadjuvant imatinib treatment and laparoscopic anus-preserving surgery for a large gastrointestinal stromal tumor of the rectum. <i>World Journal of Surgical Oncology</i> , 2016, 14, 68.	1.9	14
35	Clinicopathological and Survival Analysis of Japanese Patients with Resected Non-Small-Cell Lung Cancer Harboring <i>NKX2-1</i> , <i>SETDB1</i> , <i>MET</i> , <i>HER2</i> , <i>SOX2</i> , <i>FGFR1</i> , or <i>PIK3CA</i> Gene Amplification. <i>Journal of Thoracic Oncology</i> , 2015, 10, 1590-1600.	1.1	40
36	Mediastinal mature teratoma with complete gastrointestinal and bronchial walls. <i>Respirology Case Reports</i> , 2015, 3, 89-91.	0.6	5

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37	Novel roles for LIX1L in promoting cancer cell proliferation through ROS1-mediated LIX1L phosphorylation. <i>Scientific Reports</i> , 2015, 5, 13474.	3.3	10
38	A novel APC mosaicism in a patient with familial adenomatous polyposis. <i>Human Genome Variation</i> , 2015, 2, 15057.	0.7	7
39	BSND and ATP6V1G3. <i>Medicine (United States)</i> , 2015, 94, e989.	1.0	13
40	SASS6 overexpression is associated with mitotic chromosomal abnormalities and a poor prognosis in patients with colorectal cancer. <i>Oncology Reports</i> , 2015, 34, 727-738.	2.6	13
41	CD44-SLC1A2 Fusion Transcripts in Primary Colorectal Cancer. <i>Pathology and Oncology Research</i> , 2015, 21, 759-764.	1.9	9
42	Robust quantitative assessments of cytosine modifications and changes in the expressions of related enzymes in gastric cancer. <i>Gastric Cancer</i> , 2015, 18, 516-525.	5.3	36
43	NEIL1 p.Gln282Stop variant is predominantly localized in the cytoplasm and exhibits reduced activity in suppressing mutations. <i>Gene</i> , 2015, 571, 33-42.	2.2	7
44	Effect of boron neutron capture therapy for recurrent anaplastic meningioma: an autopsy case report. <i>Brain Tumor Pathology</i> , 2015, 32, 61-65.	1.7	6
45	Impaired 8-Hydroxyguanine Repair Activity of MUTYH Variant p.Arg109Trp Found in a Japanese Patient with Early-Onset Colorectal Cancer. <i>Oxidative Medicine and Cellular Longevity</i> , 2014, 2014, 1-12.	4.0	10
46	PLK4 overexpression and its effect on centrosome regulation and chromosome stability in human gastric cancer. <i>Molecular Biology Reports</i> , 2014, 41, 6635-6644.	2.3	51
47	CLCA2 as a Novel Immunohistochemical Marker for Differential Diagnosis of Squamous Cell Carcinoma from Adenocarcinoma of the Lung. <i>Disease Markers</i> , 2014, 2014, 1-11.	1.3	22
48	<i>miR-21</i> gene amplification is a novel predictor of a poor prognosis in patients with gastric cancer. <i>Journal of Surgical Oncology</i> , 2014, 109, 189-197.	1.7	28
49	Human DNA glycosylase enzyme TDG repairs thymine mispaired with exocyclic etheno-DNA adducts. <i>Free Radical Biology and Medicine</i> , 2014, 76, 136-146.	2.9	11
50	RSPO fusion transcripts in colorectal cancer in Japanese population. <i>Molecular Biology Reports</i> , 2014, 41, 5375-5384.	2.3	53
51	A novel somatic FGFR3 mutation in primary lung cancer. <i>Oncology Reports</i> , 2014, 31, 1219-1224.	2.6	12
52	Abstract 1601: Human colorectal cancer marker PC(16:0/16:1) induces cell growth by activating Akt and Erk pathways. <i>Cancer Research</i> , 2014, 74, 1601-1601.	0.9	1
53	Accumulated phosphatidylcholine (16:0/16:1) in human colorectal cancer; possible involvement of <i>LPCAT4</i> . <i>Cancer Science</i> , 2013, 104, 1295-1302.	3.9	96
54	Identification and association study with lung cancer for novel insertion polymorphisms of human endogenous retrovirus. <i>Carcinogenesis</i> , 2013, 34, 2531-2538.	2.8	27

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55	Lipid peroxidation-induced DNA adducts in human gastric mucosa. <i>Carcinogenesis</i> , 2013, 34, 121-127.	2.8	56
56	Visualization of phosphatidylcholine (16:0/16:0) in type II alveolar epithelial cells in the human lung using imaging mass spectrometry. <i>Pathology International</i> , 2013, 63, 195-200.	1.3	13
57	SGOL1 variant B induces abnormal mitosis and resistance to taxane in non-small cell lung cancers. <i>Scientific Reports</i> , 2013, 3, 3012.	3.3	26
58	CD74-ROS1 fusion transcripts in resected non-small cell lung carcinoma. <i>Oncology Reports</i> , 2013, 30, 1675-1680.	2.6	19
59	Role of Base Excision Repair Enzyme MUTYH in the Repair of 8-Hydroxyguanine and MUTYH-Associated Polyposis (MAP). <i>Hereditary Genetics: Current Research</i> , 2012, 01, .	0.1	3
60	Chromogenic <i>in situ</i> hybridization (CISH) to detect <i>HER2</i> gene amplification in breast and gastric cancer: Comparison with immunohistochemistry (IHC) and fluorescence <i>in situ</i> hybridization (FISH). <i>Pathology International</i> , 2012, 62, 728-734.	1.3	33
61	The CRKL gene encoding an adaptor protein is amplified, overexpressed, and a possible therapeutic target in gastric cancer. <i>Journal of Translational Medicine</i> , 2012, 10, 97.	4.4	20
62	Association between dopamine beta hydroxylase rs5320 polymorphism and smoking behaviour in elderly Japanese. <i>Journal of Human Genetics</i> , 2012, 57, 385-390.	2.3	16
63	Detection of kinase amplifications in gastric cancer archives using fluorescence <i>in situ</i> hybridization. <i>Pathology International</i> , 2012, 62, 477-484.	1.3	35
64	Impaired suppressive activities of human MUTYH variant proteins against oxidative mutagenesis. <i>World Journal of Gastroenterology</i> , 2012, 18, 6935.	3.3	4
65	Suppression of hydroxyurea-induced centrosome amplification by NORE1A and down-regulation of NORE1A mRNA expression in non-small cell lung carcinoma. <i>Lung Cancer</i> , 2011, 71, 19-27.	2.0	17
66	Germline alterations in the <i>CDH1</i> gene in familial gastric cancer in the Japanese population. <i>Cancer Science</i> , 2011, 102, 1782-1788.	3.9	42
67	A novel tumor-derived SGOL1 variant causes abnormal mitosis and unstable chromatid cohesion. <i>Oncogene</i> , 2011, 30, 4453-4463.	5.9	32
68	Aberrant Expression and Mutation-Inducing Activity of AID in Human Lung Cancer. <i>Annals of Surgical Oncology</i> , 2011, 18, 2084-2092.	1.5	36
69	Reduced expression of MUTYH with suppressive activity against mutations caused by 8-hydroxyguanine is a novel predictor of a poor prognosis in human gastric cancer. <i>Journal of Pathology</i> , 2011, 225, 414-423.	4.5	45
70	Genetic susceptibility to lung cancer. <i>Frontiers in Bioscience - Scholar</i> , 2011, S3, 1463.	2.1	14
71	Glioblastoma: Germline Mutation of TP53. , 2011, , 31-38.		0
72	Secreted form of EphA7 in lung cancer. <i>International Journal of Oncology</i> , 2010, 36, .	3.3	7

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73	EML4-ALK fusion transcripts in immunohistochemically ALK-positive non-small cell lung carcinomas. <i>Experimental and Therapeutic Medicine</i> , 2010, 1, 271-275.	1.8	11
74	Fluorescence in situ hybridization analysis with a tissue microarray: <i>in situ</i> FISH and chips TM analysis of pathology archives. <i>Pathology International</i> , 2010, 60, 543-550.	1.3	29
75	Identification of 5 novel germline APC mutations and characterization of clinical phenotypes in Japanese patients with classical and attenuated familial adenomatous polyposis. <i>BMC Research Notes</i> , 2010, 3, 305.	1.4	16
76	Adenine DNA glycosylase activity of 14 Human MutY homolog (MUTYH) variant proteins found in patients with colorectal polyposis and cancer. <i>Human Mutation</i> , 2010, 31, E1861-E1874.	2.5	37
77	Association between neuropeptide Y receptor 2 polymorphism and the smoking behavior of elderly Japanese. <i>Journal of Human Genetics</i> , 2010, 55, 755-760.	2.3	15
78	Detection of Lipid Peroxidation-Induced DNA Adducts Caused by 4-Oxo-2(<i>E</i>)-nonenal and 4-Oxo-2(<i>E</i>)-hexenal in Human Autopsy Tissues. <i>Chemical Research in Toxicology</i> , 2010, 23, 1442-1448.	3.3	67
79	Association between neurexin 1 (NRXN1) polymorphisms and the smoking behavior of elderly Japanese. <i>Psychiatric Genetics</i> , 2010, 20, 135-136.	1.1	12
80	EPH-EPHRIN in human gastrointestinal cancers. <i>World Journal of Gastrointestinal Oncology</i> , 2010, 2, 421.	2.0	6
81	Three novel <i>NEIL1</i> promoter polymorphisms in gastric cancer patients. <i>World Journal of Gastrointestinal Oncology</i> , 2010, 2, 117.	2.0	13
82	Secreted form of EphA7 in lung cancer. <i>International Journal of Oncology</i> , 2010, 36, 635-40.	3.9	12
83	Altered expression of the human base excision repair gene <i>NTH1</i> in gastric cancer. <i>Carcinogenesis</i> , 2009, 30, 1345-1352.	2.8	31
84	Human Sgo1 downregulation leads to chromosomal instability in colorectal cancer. <i>Gut</i> , 2009, 58, 249-260.	12.1	72
85	Identification and characterization of a novel germline <i>p53</i> mutation in a patient with glioblastoma and colon cancer. <i>International Journal of Cancer</i> , 2009, 125, 973-976.	5.1	18
86	Absence of germline mono-allelic promoter hypermethylation of the <i>CDH1</i> gene in gastric cancer patients. <i>Molecular Cancer</i> , 2009, 8, 63.	19.2	16
87	<i>OGG1</i> , <i>MYH</i> and <i>MTH1</i> gene variants identified in gastric cancer patients exhibiting both 8-hydroxy-2-deoxyguanosine accumulation and low inflammatory cell infiltration in their gastric mucosa. <i>Journal of Genetics</i> , 2008, 87, 181-186.	0.7	20
88	Induction of centrosome amplification and chromosome instability in <i>p53</i> -deficient lung cancer cells exposed to benzo[a]pyrene diol epoxide (B[a]PDE). <i>Journal of Pathology</i> , 2008, 216, 365-374.	4.5	37
89	Inverse relationship between the length of the <i>EGFR</i> CA repeat polymorphism in lung carcinoma and protein expression of <i>EGFR</i> in the carcinoma. <i>Journal of Surgical Oncology</i> , 2008, 98, 457-461.	1.7	14
90	Association between genetic polymorphisms of the base excision repair gene <i>MUTYH</i> and increased colorectal cancer risk in a Japanese population. <i>Cancer Science</i> , 2008, 99, 355-360.	3.9	42

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91	Characterization of adenocarcinoma of the lung in a familial adenomatous polyposis patient. <i>Pathology International</i> , 2008, 58, 706-712.	1.3	1
92	EML4-ALK fusion transcripts, but no NPM-, TPM3-, CLTC-, ATIC-, or TFG-ALK fusion transcripts, in non-small cell lung carcinomas. <i>Lung Cancer</i> , 2008, 61, 163-169.	2.0	159
93	Fluorescence-labeled Methylation-sensitive Amplified Fragment Length Polymorphism (FL-MS-AFLP) Analysis for Quantitative Determination of DNA Methylation and Demethylation Status. <i>Japanese Journal of Clinical Oncology</i> , 2008, 38, 317-322.	1.3	8
94	Identification and characterization of a novel germ line p53 mutation in familial gastric cancer in the Japanese population. <i>Carcinogenesis</i> , 2007, 28, 2013-2018.	2.8	33
95	Association between CDH1 haplotypes and gastric cancer risk in a Japanese population. <i>Scandinavian Journal of Gastroenterology</i> , 2007, 42, 1479-1485.	1.5	23
96	Sumoylation of nucleophosmin/B23 regulates its subcellular localization, mediating cell proliferation and survival. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 9679-9684.	7.1	77
97	Direct evidence for the role of centrosomally localized p53 in the regulation of centrosome duplication. <i>Oncogene</i> , 2007, 26, 2939-2944.	5.9	86
98	Thr199 phosphorylation targets nucleophosmin to nuclear speckles and represses pre-mRNA processing. <i>FEBS Letters</i> , 2006, 580, 399-409.	2.8	52
99	<i>EPHA2</i> / <i>EFNA1</i> expression in human gastric cancer. <i>Cancer Science</i> , 2005, 96, 42-47.	3.9	118
100	Effect of splice-site polymorphisms of the <i>TMPRSS4</i> , <i>NPHP4</i> and <i>ORCTL4</i> genes on their mRNA expression. <i>Journal of Genetics</i> , 2005, 84, 131-136.	0.7	8
101	Downregulation of <i>EphA7</i> by hypermethylation in colorectal cancer. <i>Oncogene</i> , 2005, 24, 5637-5647.	5.9	100
102	Characterization of centrosomal association of nucleophosmin/B23 linked to Crm1 activity. <i>FEBS Letters</i> , 2005, 579, 6621-6634.	2.8	51
103	Inactivating mutations of the human base excision repair gene <i>NEIL1</i> in gastric cancer. <i>Carcinogenesis</i> , 2004, 25, 2311-2317.	2.8	85
104	A novel splice-site variant of the base excision repair gene <i>MYH</i> is associated with production of an aberrant mRNA transcript encoding a truncated <i>MYH</i> protein not localized in the nucleus. <i>Carcinogenesis</i> , 2004, 25, 1859-1866.	2.8	53
105	Effect of exogenous <i>MSH6</i> and <i>POLD1</i> expression on the mutation rate of the <i>HPRT</i> locus in a human colon cancer cell line with mutator phenotype, DLD-1. <i>International Journal of Oncology</i> , 2004, 24, 697.	3.3	0
106	<i>Tiam1</i> mediates neurite outgrowth induced by ephrin-B1 and <i>EphA2</i> . <i>EMBO Journal</i> , 2004, 23, 1075-1088.	7.8	87
107	Splice-site genetic polymorphism of the human kallikrein 12 (<i>KLK12</i>) gene correlates with no substantial expression of <i>KLK12</i> protein having serine protease activity. <i>Human Mutation</i> , 2004, 24, 273-274.	2.5	22
108	<i>EphA2</i> Up-regulation induced by deoxycholic acid in human colon carcinoma cells, an involvement of extracellular signal-regulated kinase and p53-independence. <i>Journal of Cancer Research and Clinical Oncology</i> , 2003, 129, 703-708.	2.5	61

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109	Causation of Borrmann type 4 gastric cancer: heritable factors or environmental factors?. <i>Gastric Cancer</i> , 2003, 6, 17-23.	5.3	12
110	Ile-Leu Substitution (I415L) in Germline E-cadherin Gene (CDH1) in Japanese Familial Gastric Cancer. <i>Japanese Journal of Clinical Oncology</i> , 2003, 33, 17-20.	1.3	44
111	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. <i>Carcinogenesis</i> , 2003, 24, 1031-1037.	2.8	33
112	E-cadherin gene variants in gastric cancer families whose probands are diagnosed with diffuse gastric cancer. <i>International Journal of Cancer</i> , 2002, 101, 434-441.	5.1	91
113	A single nucleotide polymorphism at the splice donor site of the human MYH base excision repair gene results in reduced translation efficiency of its transcripts. <i>Genes To Cells</i> , 2002, 7, 461-474.	1.2	39
114	The OGG1 Gene Encodes a Repair Enzyme for Oxidatively Damaged DNA and Is Involved in Human Carcinogenesis. <i>Antioxidants and Redox Signaling</i> , 2001, 3, 597-609.	5.4	127
115	Somatic mutations and single nucleotide polymorphisms of base excision repair genes involved in the repair of 8-hydroxyguanine in damaged DNA. <i>Cancer Letters</i> , 2001, 166, 65-69.	7.2	38
116	Enhancement of OGG1 protein AP lyase activity by increase of APEX protein. <i>Mutation Research DNA Repair</i> , 2001, 486, 31-40.	3.7	56
117	Segregation Analysis of Gastric Cancer in a Japanese Population. <i>International Journal of Human Genetics</i> , 2001, 1, 263-270.	0.1	2
118	OGG1 protein suppresses G:C->T:A mutation in a shuttle vector containing 8-hydroxyguanine in human cells. <i>Carcinogenesis</i> , 2001, 22, 1355-1362.	2.8	38
119	Germline p53 Mutation in a Patient with Multiple Primary Cancers. <i>Japanese Journal of Clinical Oncology</i> , 2001, 31, 349-351.	1.3	11
120	Absence of Germline CHK2 Mutations in Familial Gastric Cancer. <i>Japanese Journal of Cancer Research</i> , 2000, 91, 875-879.	1.7	15
121	Adenine excisional repair function of MYH protein on the adenine:8-hydroxyguanine base pair in double-stranded DNA. <i>Nucleic Acids Research</i> , 2000, 28, 4912-4918.	14.5	79
122	Familial gastric cancer: clinicopathological characteristics, RER phenotype and germline p53 and E-cadherin mutations. <i>Carcinogenesis</i> , 1999, 20, 1127-1131.	2.8	115
123	Mxi1 Mutations in Human Neurofibrosarcomas. <i>Japanese Journal of Cancer Research</i> , 1999, 90, 740-746.	1.7	16
124	Microsatellite Instability and K-ras Mutations in Gastric Adenomas, with Reference to Associated Gastric Cancers. <i>Cancer Detection and Prevention</i> , 1999, 23, 204-214.	2.1	36
125	Mutational Analyses of Multiple Target Genes in Histologically Heterogeneous Gastric Cancer with Microsatellite Instability. <i>Japanese Journal of Cancer Research</i> , 1998, 89, 1284-1291.	1.7	14
126	Infrequent Mutations of the OGG1 Gene, That Is Involved in the Excision of 8-Hydroxyguanine in Damaged DNA, in Human Gastric Cancer. <i>Japanese Journal of Cancer Research</i> , 1998, 89, 825-828.	1.7	96

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127	Genetic polymorphisms and alternative splicing of the hOGG1 gene, that is involved in the repair of 8-hydroxyguanine in damaged DNA. <i>Oncogene</i> , 1998, 16, 3219-3225.	5.9	408
128	Genomic structure and chromosomal localization of the mouse Ogg1 gene that is involved in the repair of 8-hydroxyguanine in DNA damage. <i>Mammalian Genome</i> , 1998, 9, 32-37.	2.2	48
129	RER phenotype and its associated mutations in familial gastric cancer. <i>Carcinogenesis</i> , 1998, 19, 247-251.	2.8	34
130	8-hydroxyguanine (7,8-dihydro-8-oxoguanine) DNA glycosylase and AP lyase activities of hOGG1 protein and their substrate specificity. <i>Mutation Research DNA Repair</i> , 1997, 385, 75-82.	3.7	76
131	Cloning of a human homolog of the yeast OGG1 gene that is involved in the repair of oxidative DNA damage. <i>Oncogene</i> , 1997, 14, 2857-2861.	5.9	249
132	Bilateral breast tumors, malignant phyllodes tumor and invasive lobular carcinoma in a 46, XX/46, XY mosaic female with family history of breast cancer*. <i>Pathology International</i> , 1997, 47, 147-154.	1.3	6
133	Localization of Menkes gene expression in the mouse brain; its association with neurological manifestations in Menkes model mice. <i>Acta Neuropathologica</i> , 1996, 91, 482-488.	7.7	45
134	Frequent co-occurrence of mutator phenotype in synchronous, independent multiple cancers of the stomach. <i>Carcinogenesis</i> , 1995, 16, 2989-2993.	2.8	31
135	Centrosome Abnormality and Human Lung Cancer. , 0, , .		2