Gyeong Hoon Kang

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	CpG island methylator phenotype underlies sporadic microsatellite instability and is tightly associated with BRAF mutation in colorectal cancer. Nature Genetics, 2006, 38, 787-793.	21.4	1,715
2	Open versus laparoscopic surgery for mid-rectal or low-rectal cancer after neoadjuvant chemoradiotherapy (COREAN trial): survival outcomes of an open-label, non-inferiority, randomised controlled trial. Lancet Oncology, The, 2014, 15, 767-774.	10.7	713
3	Epstein-Barr Virus-Positive Gastric Carcinoma Demonstrates Frequent Aberrant Methylation of Multiple Genes and Constitutes CpG Island Methylator Phenotype-Positive Gastric Carcinoma. American Journal of Pathology, 2002, 160, 787-794.	3.8	308
4	Aberrant CpG Island Hypermethylation Along Multistep Hepatocarcinogenesis. American Journal of Pathology, 2003, 163, 1371-1378.	3.8	301
5	Identification of MicroRNA-21 as a Biomarker for Chemoresistance and Clinical Outcome Following Adjuvant Therapy in Resectable Pancreatic Cancer. PLoS ONE, 2010, 5, e10630.	2.5	261
6	Aberrant CpG island hypermethylation of multiple genes in prostate cancer and prostatic intraepithelial neoplasia. Journal of Pathology, 2004, 202, 233-240.	4.5	222
7	Aberrant CpG Island Hypermethylation of Chronic Gastritis, in Relation to Aging, Gender, Intestinal Metaplasia, and Chronic Inflammation. American Journal of Pathology, 2003, 163, 1551-1556.	3.8	221
8	Methylation of RUNX3 in various types of human cancers and premalignant stages of gastric carcinoma. Laboratory Investigation, 2004, 84, 479-484.	3.7	199
9	Panels of Immunohistochemical Markers Help Determine Primary Sites of Metastatic Adenocarcinoma. Archives of Pathology and Laboratory Medicine, 2007, 131, 1561-1567.	2.5	189
10	Hypermethylation of CpG island loci and hypomethylation of LINE-1 and Alu repeats in prostate adenocarcinoma and their relationship to clinicopathological features. Journal of Pathology, 2007, 211, 269-277.	4.5	172
11	Treatment Guidelines for Branch Duct Type Intraductal Papillary Mucinous Neoplasms of the Pancreas: When Can We Operate or Observe?. Annals of Surgical Oncology, 2008, 15, 199-205.	1.5	165
12	DNA methylation profiles of gastric carcinoma characterized by quantitative DNA methylation analysis. Laboratory Investigation, 2008, 88, 161-170.	3.7	156
13	Cholangioscopic findings in bile duct tumors. Gastrointestinal Endoscopy, 2000, 52, 630-634.	1.0	150
14	Mesenchymal Stem Cells Transfer Mitochondria to the Cells with Virtually No Mitochondrial Function but Not with Pathogenic mtDNA Mutations. PLoS ONE, 2012, 7, e32778.	2.5	146
15	Aberrant CpG island hypermethylation of multiple genes in colorectal neoplasia. Laboratory Investigation, 2004, 84, 884-893.	3.7	143
16	Aberrant CpG Island Methylation of Multiple Genes in Intrahepatic Cholangiocarcinoma. American Journal of Pathology, 2002, 161, 1015-1022.	3.8	140
17	Promoter CpG island hypermethylation during breast cancer progression. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2011, 458, 73-84.	2.8	139
18	Distinct Clinical Features and Outcomes of Gastric Cancers with Microsatellite Instability. Modern Pathology, 2002, 15, 632-640.	5.5	132

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19	Promoter hypomethylation of a novel cancer/testis antigen gene CAGE is correlated with its aberrant expression and is seen in premalignant stage of gastric carcinoma. Biochemical and Biophysical Research Communications, 2003, 307, 52-63.	2.1	131
20	Profile of Aberrant CpG Island Methylation Along the Multistep Pathway of Gastric Carcinogenesis. Laboratory Investigation, 2003, 83, 635-641.	3.7	130
21	Clinicopathological features of CpG island methylator phenotypeâ€positive colorectal cancer and its adverse prognosis in relation to <i>KRAS</i> / <i>BRAF</i> mutation. Pathology International, 2008, 58, 104-113.	1.3	127
22	Abatacept alleviates severe autoimmune symptoms in a patient carrying a de novo variant in CTLA-4. Journal of Allergy and Clinical Immunology, 2016, 137, 327-330.	2.9	125
23	Appendiceal orifice inflammation as a skip lesion in ulcerative colitis: an analysis in relation to medical therapy and disease extent. Gastrointestinal Endoscopy, 1999, 49, 743-747.	1.0	123
24	Prognostic implications of CpG island hypermethylator phenotype in colorectal cancers. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2009, 455, 485-494.	2.8	116
25	Correlation of p16 Hypermethylation with p16 Protein Loss in Sporadic Gastric Carcinomas. Laboratory Investigation, 2000, 80, 689-695.	3.7	114
26	Recurrence and Prognostic Factors of Ampullary Carcinoma after Radical Resection: Comparison with Distal Extrahepatic Cholangiocarcinoma. Annals of Surgical Oncology, 2007, 14, 3195-3201.	1.5	109
27	Comparison of CpG island hypermethylation and repetitive DNA hypomethylation in premalignant stages of gastric cancer, stratified for <i>Helicobacter pylori</i> infection. Journal of Pathology, 2009, 219, 410-416.	4.5	109
28	Genotypic analysis at multiple loci across Kaposi's sarcoma herpesvirus (KSHV) DNA molecules: clustering patterns, novel variants and chimerism. Journal of Clinical Virology, 2002, 23, 119-148.	3.1	108
29	STAT3 inhibits the degradation of HIF-1α by pVHL-mediated ubiquitination. Experimental and Molecular Medicine, 2008, 40, 479.	7.7	103
30	Polymorphism in folate- and methionine-metabolizing enzyme and aberrant CpG island hypermethylation in uterine cervical cancer. Gynecologic Oncology, 2005, 96, 173-180.	1.4	99
31	Loss of CDX2 expression is associated with poor prognosis in colorectal cancer patients. World Journal of Gastroenterology, 2015, 21, 1457.	3.3	98
32	Tumor Mutation Burden and Prognosis in Patients with Colorectal Cancer Treated with Adjuvant Fluoropyrimidine and Oxaliplatin. Clinical Cancer Research, 2019, 25, 6141-6147.	7.0	98
33	Prognostic implication of the CpG island methylator phenotype in colorectal cancers depends on tumour location. British Journal of Cancer, 2013, 109, 1004-1012.	6.4	97
34	RNA editing in <i>RHOQ</i> promotes invasion potential in colorectal cancer. Journal of Experimental Medicine, 2014, 211, 613-621.	8.5	97
35	Prognostic implications of tumor-infiltrating FoxP3+ regulatory T cells and CD8+ cytotoxic T cells in microsatellite-unstable gastric cancers. Human Pathology, 2014, 45, 285-293.	2.0	96
36	Profile of Aberrant CpG Island Methylation along Multistep Gastric Carcinogenesis. Laboratory Investigation, 2003, 83, 519-526.	3.7	95

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37	BRAF andKRAS mutations in prostatic adenocarcinoma. International Journal of Cancer, 2006, 119, 1858-1862.	5.1	93
38	Comparison of DNA hypermethylation patterns in different types of uterine cancer: Cervical squamous cell carcinoma, cervical adenocarcinoma and endometrial adenocarcinoma. International Journal of Cancer, 2006, 118, 2168-2171.	5.1	90
39	Characterisation of PD-L1-positive subsets of microsatellite-unstable colorectal cancers. British Journal of Cancer, 2016, 115, 490-496.	6.4	88
40	Epigenetic inactivation of protein kinase D1 in gastric cancer and its role in gastric cancer cell migration and invasion. Carcinogenesis, 2007, 29, 629-637.	2.8	84
41	Prognostic Implications of and Relationship Between CpG Island Hypermethylation and Repetitive DNA Hypomethylation in Hepatocellular Carcinoma. Clinical Cancer Research, 2009, 15, 812-820.	7.0	83
42	Methylome of Fetal and Maternal Monocytes and Macrophages at the Fetoâ€Maternal Interface. American Journal of Reproductive Immunology, 2012, 68, 8-27.	1.2	82
43	Loss of CDX2/CK20 Expression Is Associated With Poorly Differentiated Carcinoma, the CpG Island Methylator Phenotype, and Adverse Prognosis in Microsatellite-unstable Colorectal Cancer. American Journal of Surgical Pathology, 2013, 37, 1532-1541.	3.7	80
44	Molecular and prognostic heterogeneity of microsatellite-unstable colorectal cancer. World Journal of Gastroenterology, 2014, 20, 4230.	3.3	79
45	Estrogen and progesterone receptor status affect genome-wide DNA methylation profile in breast cancer. Human Molecular Genetics, 2010, 19, 4273-4277.	2.9	78
46	Kaposi's sarcoma-associated herpesvirus in Kikuchi's disease. Human Pathology, 1998, 29, 1091-1096.	2.0	76
47	The CpG island methylator phenotype may confer a survival benefit in patients with stage II or III colorectal carcinomas receiving fluoropyrimidine-based adjuvant chemotherapy. BMC Cancer, 2011, 11, 344.	2.6	76
48	Adenocarcinoma of the small intestine: a multi-institutional study of 197 surgically resected cases. Human Pathology, 2010, 41, 1087-1096.	2.0	75
49	Molecular Subtypes of Colorectal Cancer and Their Clinicopathologic Features, With an Emphasis on the Serrated Neoplasia Pathway. Archives of Pathology and Laboratory Medicine, 2016, 140, 406-412.	2.5	75
50	BRCA2 Fine-Tunes the Spindle Assembly Checkpoint through Reinforcement of BubR1 Acetylation. Developmental Cell, 2012, 22, 295-308.	7.0	71
51	<i>ALU</i> and <i>LINEâ€1</i> hypomethylations in multistep gastric carcinogenesis and their prognostic implications. International Journal of Cancer, 2012, 131, 1323-1331.	5.1	71
52	Targeted Sequencing of Cancer-Related Genes in Colorectal Cancer Using Next-Generation Sequencing. PLoS ONE, 2013, 8, e64271.	2.5	71
53	Four Molecular Subtypes of Colorectal Cancer and Their Precursor Lesions. Archives of Pathology and Laboratory Medicine, 2011, 135, 698-703.	2.5	71
54	Intratumoral Fusobacterium nucleatum abundance correlates with macrophage infiltration and CDKN2A methylation in microsatellite-unstable colorectal carcinoma. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2017, 471, 329-336.	2.8	70

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55	Association Between Fusobacterium nucleatum, Pathway Mutation, and Patient Prognosis in Colorectal Cancer. Annals of Surgical Oncology, 2018, 25, 3389-3395.	1.5	69
56	Chemoradiation-Induced Alteration of Programmed Death-Ligand 1 and CD8 + Tumor-Infiltrating Lymphocytes Identified Patients With Poor Prognosis in Rectal Cancer: AÂMatched Comparison Analysis. International Journal of Radiation Oncology Biology Physics, 2017, 99, 1216-1224.	0.8	68
57	Alu and LINE-1 Hypomethylation Is Associated with HER2 Enriched Subtype of Breast Cancer. PLoS ONE, 2014, 9, e100429.	2.5	66
58	Prognostic Implication of M2 Macrophages Are Determined by the Proportional Balance of Tumor Associated Macrophages and Tumor Infiltrating Lymphocytes in Microsatellite-Unstable Gastric Carcinoma. PLoS ONE, 2015, 10, e0144192.	2.5	62
59	Role of <i>Helicobacter pylori</i> infection in aberrant DNA methylation along multistep gastric carcinogenesis. Cancer Science, 2010, 101, 1337-1346.	3.9	61
60	Improved results of LINE-1 methylation analysis in formalin-fixed, paraffin-embedded tissues with the application of a heating step during the DNA extraction process. Clinical Epigenetics, 2017, 9, 1.	4.1	61
61	Nur77 Activated by Hypoxia-Inducible Factor-1α Overproduces Proopiomelanocortin in von Hippel-Lindau-Mutated Renal Cell Carcinoma. Cancer Research, 2004, 64, 35-39.	0.9	59
62	CpG Island Methylator Phenotype in Colorectal Cancers: Comparison of the New and Classic CpG Island Methylator Phenotype Marker Panels. Archives of Pathology and Laboratory Medicine, 2008, 132, 1657-1665.	2.5	59
63	Clinical and Histological Features of Nonalcoholic Fatty Liver Disease in Children. Digestive Diseases and Sciences, 2009, 54, 2225-2230.	2.3	56
64	Evaluation of myenteric ganglion cells and interstitial cells of Cajal in patients with chronic idiopathic constipation. International Journal of Colorectal Disease, 2002, 17, 253-258.	2.2	54
65	Changes in aberrant DNA methylation after <i>Helicobacter pylori</i> eradication: A longâ€ŧerm followâ€up study. International Journal of Cancer, 2013, 133, 2034-2042.	5.1	53
66	Usefulness of cholangioscopy in patients with focal stricture of the intrahepatic duct unrelated to intrahepatic stones. Gastrointestinal Endoscopy, 1999, 49, 204-209.	1.0	52
67	Depth of Mesorectal Extension Has Prognostic Significance in Patients With T3 Rectal Cancer. Diseases of the Colon and Rectum, 2012, 55, 1220-1228.	1.3	52
68	Distinct patterns of promoter CpG island methylation of breast cancer subtypes are associated with stem cell phenotypes. Modern Pathology, 2012, 25, 185-196.	5.5	52
69	KRAS Mutation is Associated with Worse Prognosis in Stage III or High-risk Stage II Colon Cancer Patients Treated with Adjuvant FOLFOX. Annals of Surgical Oncology, 2015, 22, 187-194.	1.5	52
70	Feasibility of endoscopic papillectomy in early stage ampulla of Vater cancer. Journal of Gastroenterology and Hepatology (Australia), 2009, 24, 120-124.	2.8	51
71	Clinical Outcomes of Patients with Microsatellite-Unstable Colorectal Carcinomas Depend on L1 Methylation Level. Annals of Surgical Oncology, 2012, 19, 3441-3448.	1.5	50
72	Interleukin-10 receptor mutations in children with neonatal-onset Crohn's disease and intractable ulcerating enterocolitis. European Journal of Gastroenterology and Hepatology, 2013, 25, 1.	1.6	50

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73	Association between mutations of critical pathway genes and survival outcomes according to the tumor location in colorectal cancer. Cancer, 2017, 123, 3513-3523.	4.1	50
74	Open versus laparoscopic surgery for mid or low rectal cancer after neoadjuvant chemoradiotherapy (COREAN trial): 10-year follow-up of an open-label, non-inferiority, randomised controlled trial. The Lancet Gastroenterology and Hepatology, 2021, 6, 569-577.	8.1	50
75	Histology of Gastroesophageal Junction in Fetal and Pediatric Autopsy. Archives of Pathology and Laboratory Medicine, 2003, 127, 451-455.	2.5	50
76	Source of errors in the evaluation of early rectal cancer by endoluminal ultrasonography. Diseases of the Colon and Rectum, 2001, 44, 1302-1309.	1.3	49
77	Identification of Novel Methylation Markers in Hepatocellular Carcinoma using a Methylation Array. Journal of Korean Medical Science, 2010, 25, 1152.	2.5	49
78	DNA methylation profile during multistage progression of pulmonary adenocarcinomas. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2011, 459, 201-211.	2.8	49
79	Down-Regulation of Dual-Specificity Phosphatase 5 in Gastric Cancer by Promoter CpG Island Hypermethylation and Its Potential Role in Carcinogenesis. American Journal of Pathology, 2013, 182, 1275-1285.	3.8	49
80	Methylation and microsatellite status and recurrence following adjuvant FOLFOX in colorectal cancer. International Journal of Cancer, 2013, 132, 2209-2216.	5.1	49
81	Changes in Hepatic Gene Expression upon Oral Administration of Taurine-Conjugated Ursodeoxycholic Acid in ob/ob Mice. PLoS ONE, 2010, 5, e13858.	2.5	47
82	Nodular regenerative hyperplasia of the liver in Budd–Chiari syndrome: CT and MR features. Abdominal Imaging, 2000, 25, 255-258.	2.0	46
83	Gastric-type expression signature in serrated pathway–associated colorectal tumors. Human Pathology, 2015, 46, 643-656.	2.0	45
84	Zinc deficiency manifested by dermatitis and visual dysfunction in a patient with Crohn's disease. Journal of Gastroenterology, 1998, 33, 876-879.	5.1	44
85	Low-dose CT for the diagnosis of appendicitis in adolescents and young adults (LOCAT): a pragmatic, multicentre, randomised controlled non-inferiority trial. The Lancet Gastroenterology and Hepatology, 2017, 2, 793-804.	8.1	44
86	Genomic hypomethylation and CpG island hypermethylation in prostatic intraepithelial neoplasm. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2009, 454, 17-23.	2.8	43
87	The beneficial effect of palliative resection in metastatic colorectal cancer. British Journal of Cancer, 2013, 108, 1425-1431.	6.4	43
88	Adverse prognostic impact of the CpG island methylator phenotype in metastatic colorectal cancer. British Journal of Cancer, 2016, 115, 164-171.	6.4	43
89	CpG island hypermethylation and repetitive DNA hypomethylation in premalignant lesion of extrahepatic cholangiocarcinoma. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2009, 455, 343-351.	2.8	42
90	Differential clinicopathological features in microsatellite instability-positive colorectal cancers depending on CIMP status. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2011, 459, 55-63.	2.8	42

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91	Overexpression of POSTN in Tumor Stroma Is a Poor Prognostic Indicator of Colorectal Cancer. Journal of Pathology and Translational Medicine, 2017, 51, 306-313.	1.1	42
92	Methylation Profiles of Multiple CpG Island Loci in Extrahepatic Cholangiocarcinoma Versus Those of Intrahepatic Cholangiocarcinomas. Archives of Pathology and Laboratory Medicine, 2007, 131, 923-930.	2.5	42
93	BRAF Activation Initiates but Does Not Maintain Invasive Prostate Adenocarcinoma. PLoS ONE, 2008, 3, e3949.	2.5	41
94	The distribution of intratumoral macrophages correlates with molecular phenotypes and impacts prognosis in colorectal carcinoma. Histopathology, 2018, 73, 663-671.	2.9	41
95	Heterodimerization of Glycosylated Insulin-Like Growth Factor-1 Receptors and Insulin Receptors in Cancer Cells Sensitive to Anti-IGF1R Antibody. PLoS ONE, 2012, 7, e33322.	2.5	41
96	Genome-Wide DNA Methylation Profiles in Noncancerous Gastric Mucosae with Regard to Helicobacter pylori Infection and the Presence of Gastric Cancer. Helicobacter, 2011, 16, 179-188.	3.5	40
97	Prognostic implication of mucinous histology in colorectal cancer patients treated with adjuvant FOLFOX chemotherapy. British Journal of Cancer, 2013, 108, 1978-1984.	6.4	40
98	Distinct clinical outcomes of two CIMP-positive colorectal cancer subtypes based on a revised CIMP classification system. British Journal of Cancer, 2017, 116, 1012-1020.	6.4	40
99	Detection of KIT and PDGFRA mutations in the plasma of patients with gastrointestinal stromal tumor. Targeted Oncology, 2015, 10, 597-601.	3.6	39
100	Expression Profile of LGR5 and Its Prognostic Significance in Colorectal Cancer Progression. American Journal of Pathology, 2018, 188, 2236-2250.	3.8	39
101	Synovial Sarcoma of the Kidney With Rhabdoid Features. American Journal of Surgical Pathology, 2004, 28, 634-637.	3.7	38
102	Radiological Spectrum of Hepatic Mesenchymal Hamartoma in Children. Korean Journal of Radiology, 2007, 8, 498.	3.4	38
103	Pharmacogenetic analysis of adjuvant FOLFOX for Korean patients with colon cancer. Cancer Chemotherapy and Pharmacology, 2013, 71, 843-851.	2.3	37
104	Whole-Slide Image Analysis Reveals Quantitative Landscape of Tumor–Immune Microenvironment in Colorectal Cancers. Clinical Cancer Research, 2020, 26, 870-881.	7.0	37
105	Smooth Muscle Distribution in the Extrahepatic Bile Duct. American Journal of Surgical Pathology, 2000, 24, 660-667.	3.7	36
106	Mixed-type gastric cancer and its association with high-frequency CpG island hypermethylation. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2010, 456, 625-633.	2.8	36
107	CpG island hypermethylator phenotype in gastric carcinoma and its clinicopathological features. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2010, 457, 415-422.	2.8	36
108	Prognostic Impact of <i>Fusobacterium nucleatum</i> Depends on Combined Tumor Location and Microsatellite Instability Status in Stage II/III Colorectal Cancers Treated with Adjuvant Chemotherapy. Journal of Pathology and Translational Medicine, 2019, 53, 40-49.	1.1	36

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109	An Experimental Study of Embolic Effect According to Infusion Rate and Concentration of Suspension in Transarterial Particulate Embolization. Investigative Radiology, 1997, 32, 260-267.	6.2	35
110	Distinct features between <i>MLH1</i> -methylated and unmethylated colorectal carcinomas with the CpG island methylator phenotype: implications in the serrated neoplasia pathway. Oncotarget, 2016, 7, 14095-14111.	1.8	35
111	Standardized Pathology Report for Colorectal Cancer, 2nd Edition. Journal of Pathology and Translational Medicine, 2020, 54, 1-19.	1.1	35
112	Tufting Enteropathy with EpCAM Mutations in Two Siblings. Gut and Liver, 2010, 4, 407-410.	2.9	34
113	Phase II study of biweekly S-1 and oxaliplatin combination chemotherapy in metastatic colorectal cancer and pharmacogenetic analysis. Cancer Chemotherapy and Pharmacology, 2011, 67, 1323-1331.	2.3	34
114	Downregulation of acetyl-CoA synthetase 2 is a metabolic hallmark of tumor progression and aggressiveness in colorectal carcinoma. Modern Pathology, 2017, 30, 267-277.	5.5	34
115	CpG Island Methylator Phenotype-High Colorectal Cancers and Their Prognostic Implications and Relationships with the Serrated Neoplasia Pathway. Gut and Liver, 2017, 11, 38-46.	2.9	34
116	Endoscopic treatment of gastritis cystica polyposa found in an unoperated stomach. Gastrointestinal Endoscopy, 2001, 54, 101-103.	1.0	33
117	Bile-Based Detection of Extrahepatic Cholangiocarcinoma with Quantitative DNA Methylation Markers and Its High Sensitivity. Journal of Molecular Diagnostics, 2012, 14, 256-263.	2.8	33
118	CpG methylation in exon 1 of transcription factor 4 increases with age in normal gastric mucosa and is associated with gene silencing in intestinal-type gastric cancers. Carcinogenesis, 2008, 29, 1623-1631.	2.8	32
119	Prediction of the risk for gastric cancer using candidate methylation markers in the nonâ€neoplastic gastric mucosae. Journal of Pathology, 2012, 226, 654-665.	4.5	32
120	Aberrant GATA2 epigenetic dysregulation induces a GATA2/GATA6 switch in human gastric cancer. Oncogene, 2018, 37, 993-1004.	5.9	32
121	p53 expression status is associated with cancer-specific survival in stage III and high-risk stage II colorectal cancer patients treated with oxaliplatin-based adjuvant chemotherapy. British Journal of Cancer, 2019, 120, 797-805.	6.4	32
122	p53 Mutation and overexpression in hepatocellular carcinoma and dysplastic nodules in the liver. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 1998, 432, 27-32.	2.8	31
123	Clinical Manifestations and Treatment Outcomes of Eosinophilic Gastroenteritis in Children. Pediatric Gastroenterology, Hepatology and Nutrition, 2015, 18, 253.	1.2	30
124	Differential immune microenvironmental features of microsatellite-unstable colorectal cancers according to Fusobacterium nucleatum status. Cancer Immunology, Immunotherapy, 2021, 70, 47-59.	4.2	30
125	Clinicopathologic and Molecular Characteristics of Synchronous Colorectal Cancers. Diseases of the Colon and Rectum, 2012, 55, 181-190.	1.3	29
126	Expression status of wild-type HSP110 correlates with HSP110 T17 deletion size and patient prognosis in microsatellite-unstable colorectal cancer. Modern Pathology, 2014, 27, 443-453.	5.5	29

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127	Image Analyzer-Based Assessment of Tumor-Infiltrating T Cell Subsets and Their Prognostic Values in Colorectal Carcinomas. PLoS ONE, 2015, 10, e0122183.	2.5	29
128	Efficiency of combined colonoscopy and computed tomography for diagnosis of colonic actinomycosis: a retrospective evaluation of eight consecutive patients. International Journal of Colorectal Disease, 2000, 15, 236-242.	2.2	28
129	Isolated splenic metastasis from colorectal carcinoma: a case report. Journal of Korean Medical Science, 2000, 15, 355.	2.5	28
130	Distribution of intestinal stem cell markers in colorectal precancerous lesions. Histopathology, 2016, 68, 567-577.	2.9	28
131	Risk Factors for Metachronous Gastric Neoplasms in Patients Who Underwent Endoscopic Resection of a Gastric Neoplasm. Gut and Liver, 2016, 10, 228.	2.9	28
132	Nested PCR for diagnosis of tuberculous lymphadenitis and PCR-SSCP for identification of rifampicin resistance in fine-needle aspirates. Diagnostic Cytopathology, 2002, 26, 228-231.	1.0	27
133	Helicobacter pylori-infection-associated CpG island hypermethylation in the stomach and its possible association with Polycomb repressive marks. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2008, 452, 515-524.	2.8	27
134	Total lesion glycolysis (TLG) as an imaging biomarker in metastatic colorectal cancer patients treated with regorafenib. European Journal of Nuclear Medicine and Molecular Imaging, 2017, 44, 757-764.	6.4	27
135	Decrease of 5hmC in gastric cancers is associated withTET1silencing due to with DNA methylation and bivalent histone marks atTET1CpG island 3′-shore. Oncotarget, 2015, 6, 37647-37662.	1.8	27
136	Association of Promoter Hypermethylation of the RASSF1A Gene With Prognostic Parameters in Endometrial Cancer. Oncology Research, 2006, 16, 205-209.	1.5	26
137	Subsets of microsatellite-unstable colorectal cancers exhibit discordance between the CpG island methylator phenotype and MLH1 methylation status. Modern Pathology, 2013, 26, 1013-1022.	5.5	26
138	Evolving pathologic concepts of serrated lesions of the colorectum. Journal of Pathology and Translational Medicine, 2020, 54, 276-289.	1.1	26
139	DNA Hypomethylation ofCAGEPromotors in Squamous Cell Carcinoma of Uterine Cervix. Annals of the New York Academy of Sciences, 2006, 1091, 218-224.	3.8	25
140	Leptin, MUC2 and mTOR in Appendiceal Mucinous Neoplasms. Pathobiology, 2012, 79, 45-53.	3.8	25
141	Comparative validation of assessment criteria for Crohn-like lymphoid reaction in colorectal carcinoma. Journal of Clinical Pathology, 2015, 68, 22-28.	2.0	25
142	Prognostic significance of promoter CpG island hypermethylation and repetitive DNA hypomethylation in stage I lung adenocarcinoma. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2015, 466, 675-683.	2.8	25
143	Comparison of Clinicopathologic Characteristics and Genetic Alterations Between Microsatellite Instability-Positive and Microsatellite Instability-Negative Sporadic Colorectal Carcinomas in Patients Younger than 40 Years Old. Diseases of the Colon and Rectum, 2002, 45, 219-228.	1.3	24
144	Hypermethylation of the COX-2 gene is a potential prognostic marker for cervical cancer. Journal of Obstetrics and Gynaecology Research, 2007, 33, 236-241.	1.3	24

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145	High survivin expression as a predictor of poor response to preoperative chemoradiotherapy in locally advanced rectal cancer. International Journal of Colorectal Disease, 2011, 26, 1019-1023.	2.2	24
146	Prognostic significance of CD103+ immune cells in solid tumor: a systemic review and meta-analysis. Scientific Reports, 2019, 9, 3808.	3.3	24
147	Annexin A10 expression in colorectal cancers with emphasis on the serrated neoplasia pathway. World Journal of Gastroenterology, 2015, 21, 9749.	3.3	24
148	Methylation Profiles of CpG Island Loci in Major Types of Human Cancers. Journal of Korean Medical Science, 2007, 22, 311.	2.5	23
149	Epigenetic alterations in colorectal cancer: the CpG island methylator phenotype. Histology and Histopathology, 2013, 28, 585-95.	0.7	23
150	CpG Island Hypermethylation in Gastric Carcinoma and Its Premalignant Lesions. Korean Journal of Pathology, 2012, 46, 1.	1.3	23
151	α-Fetoprotein-producing gastric carcinoma presenting focal hepatoid differentiation in metastatic lymph nodes. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 1998, 432, 85-87.	2.8	22
152	Primary hepatic carcinoid tumor with a paranuclear clear zone: a case report. Journal of Korean Medical Science, 1998, 13, 317.	2.5	22
153	Differential clinicopathologic features in microsatellite-unstable gastric cancers with and without MLH1 methylation. Human Pathology, 2013, 44, 1055-1064.	2.0	22
154	Combined prognostic effect of PD-L1 expression and immunoscore in microsatellite-unstable advanced gastric cancers. Oncotarget, 2017, 8, 58887-58902.	1.8	22
155	5' CpG island methylation of p16 is associated with absence of p16 expression in glioblastomas. Journal of Korean Medical Science, 2000, 15, 555.	2.5	20
156	Influence of IL1B polymorphism on CpG island hypermethylation in Helicobacter pylori-infected gastric cancer. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2010, 456, 647-652.	2.8	20
157	ITF2 Prevents Activation of the β-Catenin–TCF4 Complex in Colon Cancer Cells and Levels Decrease With Tumor Progression. Gastroenterology, 2014, 147, 430-442.e8.	1.3	20
158	Clinical Implication of Anti-Angiogenic Effect of Regorafenib in Metastatic Colorectal Cancer. PLoS ONE, 2015, 10, e0145004.	2.5	20
159	Differential protein immunoexpression profiles in appendiceal mucinous neoplasms: a special reference to classification and predictive factors. Modern Pathology, 2009, 22, 1102-1112.	5.5	19
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