## Tomomitsu Tahara

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	MicroRNA Profile of Human Small Intestinal Tumors Compared to Colorectal Tumors. Journal of Clinical Medicine, 2022, 11, 2604.	1.0	1
2	A giant Brunner gland hamartoma successfully treated by endoscopic excision followed by transanal retrieval. Medicine (United States), 2021, 100, e25048.	0.4	3
3	<i>MAFK</i> Polymorphisms Located in 3′-UTR are Associated with Severity of Atrophy and <i>CDKN2A</i> Methylation Status in the Gastric Mucosa. Genetic Testing and Molecular Biomarkers, 2021, 25, 255-262.	0.3	1
4	H. pylori negative gastric MALT lymphoma with API2-MALT1 translocation treated by endoscopic submucosal dissection. Medicine (United States), 2021, 100, e24371.	0.4	1
5	Effect of DNMT3A polymorphisms on CpG island hypermethylation in gastric mucosa. BMC Medical Genetics, 2020, 21, 205.	2.1	3
6	Influence of MIF polymorphisms on CpG island hyper-methylation of CDKN2A in the patients with ulcerative colitis. BMC Medical Genetics, 2020, 21, 201.	2.1	3
7	A Comparative Study of White Light Endoscopy, Chromoendoscopy and Magnifying Endoscopy with Narrow Band Imaging in the Diagnosis of Early Gastric Cancer after Helicobacter pylori Eradication. Journal of Gastrointestinal and Liver Diseases, 2020, 26, 357-362.	0.5	19
8	DNA methylation accumulation in gastric mucosa adjacent to cancer after <i>Helicobacter pylori</i> eradication. International Journal of Cancer, 2019, 144, 80-88.	2.3	25
9	Induced miRâ€31 by 5â€fluorouracil exposure contributes to the resistance in colorectal tumors. Cancer Science, 2019, 110, 2540-2548.	1.7	14
10	Lower LINE-1 methylation is associated with promoter hypermethylation and distinct molecular features in gastric cancer. Epigenomics, 2019, 11, 1651-1659.	1.0	4
11	Gastric Mucosal Microarchitectures Associated with Irreversibility with <i>Helicobacter pylori</i> Eradication and Downregulation of Micro RNA (miR)-124a. Cancer Investigation, 2019, 37, 417-426.	0.6	4
12	Prostate Stem Cell Antigen Gene Polymorphism Is Associated with <i>H. pylori</i> –related Promoter DNA Methylation in Nonneoplastic Gastric Epithelium. Cancer Prevention Research, 2019, 12, 579-584.	0.7	5
13	Eradication of <b><i>Helicobacter pylori</i></b> Induces Immediate Regressive Changes in Early Gastric Adenocarcinomas. Pathobiology, 2019, 86, 135-144.	1.9	6
14	Association of genetic polymorphisms in DNMT3A with the progression of gastric mucosal atrophy and susceptibility to gastric cancer in Japan. Oncology Letters, 2019, 17, 3482-3488.	0.8	1
15	Comparative study of magnifying narrow-band imaging and conventional white light endoscopy in the diagnosis of Helicobacter pylori status after eradication therapy. Medicine (United States), 2019, 98, e17697.	0.4	6
16	Molecular subtyping of gastric cancer combining genetic and epigenetic anomalies provides distinct clinicopathological features and prognostic impacts. Human Mutation, 2019, 40, 347-354.	1.1	10
17	Association between receptor interacting serine/threonine kinase 2 polymorphisms and gastric cancer susceptibility. Oncology Letters, 2018, 15, 3772-3778.	0.8	10
18	Development and endoscopic appearance of colorectal tumors are characterized by the expression profiles of miRNAs. Medical Molecular Morphology, 2018, 51, 82-88.	0.4	4

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19	Telomere length in the gastric mucosa after Helicobacter pylori eradication and its potential role in the gastric carcinogenesis. Clinical and Experimental Medicine, 2018, 18, 21-26.	1.9	6
20	Methylation status of IGF2 DMR and LINE1 in leukocyte DNA provides distinct clinicopathological features of gastric cancer patients. Clinical and Experimental Medicine, 2018, 18, 215-220.	1.9	11
21	Polymorphism rs7521584 in miR‑429 is associated with the severity of atrophic gastritis in patients with Helicobacter�pylori infection. Molecular Medicine Reports, 2018, 18, 2381-2386.	1.1	1
22	Unusual growth of an Epstein-Barr virus-associated differentiated early-stage gastric carcinoma: A case report. Molecular and Clinical Oncology, 2018, 8, 657-660.	0.4	2
23	Magnifying narrow-band imaging of gastric mucosal morphology predicts the H. pylori-related epigenetic field defect. Scientific Reports, 2017, 7, 3090.	1.6	7
24	Morphologic characterization of residual DNA methylation in the gastric mucosa after <i>Helicobacter pylori</i> eradication. Cancer Medicine, 2017, 6, 1730-1737.	1.3	9
25	Comprehensive DNA Methylation Profiling of Inflammatory Mucosa in Ulcerative Colitis. Inflammatory Bowel Diseases, 2017, 23, 165-173.	0.9	15
26	A comparative study of magnifying blue laser imaging and magnifying narrow-band imaging system for endoscopic diagnosis of Helicobacter pylori infection. Biomedical Reports, 2017, 7, 236-240.	0.9	4
27	Magnifying NBI Patterns of Gastric Mucosa After Helicobacter pylori Eradication and Its Potential Link to the Gastric Cancer Risk. Digestive Diseases and Sciences, 2017, 62, 2421-2427.	1.1	8
28	Evaluations of primary lesions by endoscopy clearly distinguishes prognosis in patients with gastric cancer who receive chemotherapy. PLoS ONE, 2017, 12, e0173663.	1.1	7
29	Potential link between Fusobacterium enrichment and DNA methylation accumulation in the inflammatory colonic mucosa in ulcerative colitis. Oncotarget, 2017, 8, 61917-61926.	0.8	29
30	Telomere Length in Leukocyte DNA in Gastric Cancer Patients and its Association with Clinicopathological Features and Prognosis. Anticancer Research, 2017, 37, 1997-2001.	0.5	4
31	Genetic polymorphisms of MAFK, encoding a small Maf protein, are associated with susceptibility to ulcerative colitis in Japan. World Journal of Gastroenterology, 2017, 23, 5364.	1.4	5
32	Endoscopic features of lymphoid follicles using blue laser imaging (BLI) endoscopy in the colorectum and its association with chronic bowel symptoms. PLoS ONE, 2017, 12, e0182224.	1.1	3
33	Telomere length shortening in gastric mucosa is a field effect associated with increased risk of gastric cancer. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2016, 469, 19-24.	1.4	20
34	Telomere length in non-neoplastic gastric mucosa and its relationship to H. pylori infection, degree of gastritis, and NSAID use. Clinical and Experimental Medicine, 2016, 16, 65-71.	1.9	14
35	A Possible Link between Gastric Mucosal Atrophy and Gastric Cancer after Helicobacter pylori Eradication. PLoS ONE, 2016, 11, e0163700.	1.1	14
36	Demonstration of potential link between Helicobacter pylori related promoter CpG island methylation and telomere shortening in human gastric mucosa. Oncotarget, 2016, 7, 43989-43996.	0.8	15

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37	Association between interleukin-1β and tumor necrosis factor-α polymorphisms and symptoms of dyspepsia. Molecular Medicine Reports, 2015, 11, 3888-3893.	1.1	3
38	Association between individual response to food taste and gastroesophageal symptoms. Journal of Digestive Diseases, 2015, 16, 337-341.	0.7	3
39	Fusobacterium Detected in Colonic Biopsy and Clinicopathological Features of Ulcerative Colitis in Japan. Digestive Diseases and Sciences, 2015, 60, 205-210.	1.1	54
40	DNA methylation as a molecular biomarker in gastric cancer. Epigenomics, 2015, 7, 475-486.	1.0	142
41	Relationship between Expression of Onco-Related miRNAs and the Endoscopic Appearance of Colorectal Tumors. International Journal of Molecular Sciences, 2015, 16, 1526-1543.	1.8	17
42	Telomere length in non-neoplastic colonic mucosa in ulcerative colitis (UC) and its relationship to the severe clinical phenotypes. Clinical and Experimental Medicine, 2015, 15, 327-332.	1.9	18
43	Endoscopic submucosal dissection of an esophageal tumor using a transnasal endoscope without sedation. Endoscopy, 2014, 46, E115-E116.	1.0	6
44	A new method using a polyglycolic acid monolayer patch to shield the mucosal defect after endoscopic submucosal dissection. Endoscopy, 2014, 46, E176-E177.	1.0	2
45	Colorectal Carcinomas With CpG Island Methylator Phenotype 1 Frequently Contain Mutations in Chromatin Regulators. Gastroenterology, 2014, 146, 530-538.e5.	0.6	76
46	<i>Fusobacterium</i> in Colonic Flora and Molecular Features of Colorectal Carcinoma. Cancer Research, 2014, 74, 1311-1318.	0.4	389
47	Sa1931 Association Between Common Genetic Variants in Pre-MicroRNAs and Prognosis of Advanced Gastric Cancer Treated Chemotherapy. Gastroenterology, 2014, 146, S-332.	0.6	8
48	DNA Methylation Status of Epithelial-Mesenchymal Transition (EMT) - Related Genes Is Associated with Severe Clinical Phenotypes in Ulcerative Colitis (UC). PLoS ONE, 2014, 9, e107947.	1.1	26
49	Change in DNA Methylation Patterns of SLC6A4 Gene in the Gastric Mucosa in Functional Dyspepsia. PLoS ONE, 2014, 9, e105565.	1.1	6
50	Influence of HRH2 promoter polymorphism on aberrant DNA methylation of DAPK and CDH1in the gastric epithelium. BMC Gastroenterology, 2013, 13, 1.	0.8	47
51	Influence of IL17A polymorphisms (rs2275913 and rs3748067) on the susceptibility to ulcerative colitis. Clinical and Experimental Medicine, 2013, 13, 239-244.	1.9	41
52	Genetic polymorphisms of SCN10A are associated with functional dyspepsia in Japanese subjects. Journal of Gastroenterology, 2013, 48, 73-80.	2.3	26
53	Examination of Whole Blood DNA Methylation as a Potential Risk Marker for Gastric Cancer. Cancer Prevention Research, 2013, 6, 1093-1100.	0.7	35
54	Association between common genetic variant of HRH2 and gastric cancer risk. International Journal of Oncology, 2012, 41, 497-503.	1.4	20

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55	NFKB1 polymorphism is associated with age-related gene methylation in Helicobacter pylori-infected subjects. International Journal of Molecular Medicine, 2012, 30, 255-262.	1.8	8
56	Genetic polymorphism of pri-microRNA 325, targeting SLC6A4 3′-UTR, is closely associated with the risk of functional dyspepsia in Japan. Journal of Gastroenterology, 2012, 47, 1091-1098.	2.3	20
57	Influence of IL17A polymorphisms on the aberrant methylation of DAPK and CDH1 in non-cancerous gastric mucosa. BMC Medical Genetics, 2012, 13, 59.	2.1	6
58	-449 C>G polymorphism ofNFKB1gene, coding nuclear factor-kappa-B, is associated with the susceptibility to ulcerative colitis. World Journal of Gastroenterology, 2012, 18, 6981.	1.4	2
59	Potential usefulness of DNA methylation as a risk marker for digestive cancer associated with inflammation. Expert Review of Molecular Diagnostics, 2012, 12, 489-497.	1.5	10
60	Presence of Minimal Change Esophagitis Closely Correlates with Pathological Conditions in the Stomach. Digestive Diseases and Sciences, 2012, 57, 958-966.	1.1	5
61	Correlation between magnifying narrow band imaging and histopathology in gastric protruding/or polypoid lesions: a pilot feasibility trial. BMC Gastroenterology, 2012, 12, 17.	0.8	29
62	Synergistic effect of IL-1β and TNF-α polymorphisms on the H. pylori-related gastric pre-malignant condition. Hepato-Gastroenterology, 2012, 59, 2416-20.	0.5	5
63	Combination Chemotherapy with S-1 and Docetaxel in Advanced Gastric Cancer patient with Peritoneal Dissemination and Malignant Ascites. Annals of Cancer Research and Therapy, 2012, 20, 07-10.	0.1	Ο
64	Role of heat-shock protein (HSP) 70-2 genotype in peptic ulcer in Japanese population. Hepato-Gastroenterology, 2012, 59, 426-9.	0.5	3
65	Examination of serum pepsinogen in functional dyspepsia. Hepato-Gastroenterology, 2012, 59, 2516-22.	0.5	4
66	Usefulness of Magnifying Narrow-Band Imaging Endoscopy in the Helicobacter pylori-Related Chronic Gastritis. Digestion, 2011, 83, 161-166.	1.2	22
67	Polymorphisms of DNA Repair and Xenobiotic Genes Predispose to CpG Island Methylation in Nonâ€Neoplastic Gastric Mucosa. Helicobacter, 2011, 16, 99-106.	1.6	8
68	Influence of MDR1 Polymorphism on H. pylori-Related Chronic Gastritis. Digestive Diseases and Sciences, 2011, 56, 103-108.	1.1	9
69	Association Study of Common Genetic Variants in Pre-microRNAs in Patients with Ulcerative Colitis. Journal of Clinical Immunology, 2011, 31, 69-73.	2.0	75
70	Changes in gastric mucosal patterns seen by magnifying NBI during H. pylori eradication. Journal of Gastroenterology, 2011, 46, 175-182.	2.3	42
71	Association between polymorphisms in the XRCC1 and GST genes, and CpG island methylation status in colonic mucosa in ulcerative colitis. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2011, 458, 205-211.	1.4	10
72	Effect of IL-1β and TNF-α polymorphisms on the prognosis and survival of gastric cancer patients. Clinical and Experimental Medicine, 2011, 11, 211-217.	1.9	21

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73	Effect of polymorphisms of ILâ€1β and TNFâ€Î± genes on CpG island hyper methylation (CIHM) in the nonneoplastic gastric mucosa. Molecular Carcinogenesis, 2011, 50, 835-845.	1.3	3
74	Effect of genetic polymorphisms related to DNA repair and the xenobiotic pathway on the prognosis and survival of gastric cancer patients. Anticancer Research, 2011, 31, 705-10.	0.5	12
75	Host genetic factors, related to inflammatory response, influence the CpG island methylation status in colonic mucosa in ulcerative colitis. Anticancer Research, 2011, 31, 933-8.	0.5	9
76	Association between genetic polymorphisms related to DNA repair or xenobiotic pathways and gastric premalignant conditions. Anticancer Research, 2011, 31, 1459-65.	0.5	4
77	Association between common genetic variants in pre-microRNAs and the clinicopathological characteristics and survival of gastric cancer patients. Experimental and Therapeutic Medicine, 2010, 1, 1035-1040.	0.8	10
78	Functional promoter polymorphisms of macrophage migration inhibitory factor in peptic ulcer diseases. International Journal of Molecular Medicine, 2010, 26, 707-11.	1.8	12
79	Association between IL-17A, -17F and MIF polymorphisms predispose to CpG island hyper-methylation in gastric cancer. International Journal of Molecular Medicine, 2010, 25, 471-7.	1.8	23
80	Homozygous TRPV1 315C Influences the Susceptibility to Functional Dyspepsia. Journal of Clinical Gastroenterology, 2010, 44, e1-e7.	1.1	55
81	Chronic Aspirin Use Suppresses CDH1 Methylation in Human Gastric Mucosa. Digestive Diseases and Sciences, 2010, 55, 54-59.	1.1	29
82	Association Between Cyclin D1 Polymorphism with CpG Island Promoter Methylation Status of Tumor Suppressor Genes in Gastric Cancer. Digestive Diseases and Sciences, 2010, 55, 3449-3457.	1.1	5
83	Genetic Polymorphisms of Molecules Associated with Innate Immune Responses, TRL2 and MBL2 Genes in Japanese Subjects with Functional Dyspepsia. Journal of Clinical Biochemistry and Nutrition, 2010, 47, 217-223.	0.6	18
84	Increased Number of CpG Island Hypermethylation in Tumor Suppressor Genes of Non-Neoplastic Gastric Mucosa Correlates with Higher Risk of Gastric Cancer. Digestion, 2010, 82, 27-36.	1.2	22
85	The mucosal pattern in the non-neoplastic gastric mucosa by using magnifying narrow-band imaging endoscopy significantly correlates with gastric cancer risk. Gastrointestinal Endoscopy, 2010, 71, 429-430.	0.5	9
86	Association between functional promoter polymorphisms of macrophage migration inhibitory factor (MIF) gene and ulcerative colitis in Japan. Cytokine, 2010, 51, 173-177.	1.4	31
87	Promoter Methylation Status of Multidrug Resistance 1 (MDR1) Gene in Noncancerous Gastric Mucosa Correlates With <i>Helicobacter Pylori</i> Infection and Gastric Cancer Occurrence. Cancer Investigation, 2010, 28, 711-716.	0.6	7
88	CpG island promoter methylation (CIHM) status of tumor suppressor genes correlates with morphological appearances of gastric cancer. Anticancer Research, 2010, 30, 239-44.	0.5	6
89	Promoter methylation of protease-activated receptor (PAR2) is associated with severe clinical phenotypes of ulcerative colitis (UC). Clinical and Experimental Medicine, 2009, 9, 125-130.	1.9	42
90	Effect of RANTES Promoter Genotype on the Severity of Intestinal Metaplasia in Helicobacter pylori-Infected Japanese Subjects. Digestive Diseases and Sciences, 2009, 54, 1247-1252.	1.1	7

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91	MTHFR 677T Carrier Influences the Methylation Status of H. Pylori-Infected Gastric Mucosa in Older Subjects. Digestive Diseases and Sciences, 2009, 54, 2391-2398.	1.1	11
92	Chronic nonsteroidal antiâ€inflammatory drug (NSAID) use suppresses multiple CpG islands hyper methylation (CIHM) of tumor suppressor genes in the human gastric mucosa. Cancer Science, 2009, 100, 1192-1197.	1.7	20
93	Genetic polymorphism of interleukin-17A and -17F genes in gastric carcinogenesis. Human Immunology, 2009, 70, 547-551.	1.2	120
94	Castric mucosal pattern by using magnifying narrow-band imaging endoscopy clearly distinguishes histological and serological severity of chronic gastritis. Gastrointestinal Endoscopy, 2009, 70, 246-253.	0.5	132
95	Association of polymorphism of thep22PHOXcomponent of NADPH oxidase in gastroduodenal diseases in Japan. Scandinavian Journal of Gastroenterology, 2009, 44, 296-300.	0.6	7
96	Mannan-binding lectin B allele is associated with a risk of developing more severe gastric mucosal atrophy in Helicobacter pylori-infected Japanese patients. European Journal of Gastroenterology and Hepatology, 2009, 21, 781-786.	0.8	6
97	A genetic variant of the p22PHOX component of NADPH oxidase C242T is associated with reduced risk of functional dyspepsia in Helicobacter pylori-infected Japanese individuals. European Journal of Gastroenterology and Hepatology, 2009, 21, 1363-1368.	0.8	13
98	Increased number of methylated CpG islands correlates with Helicobacter pylori infection, histological and serological severity of chronic gastritis. European Journal of Gastroenterology and Hepatology, 2009, 21, 613-619.	0.8	31
99	Effect of MDR1 gene promoter methylation in patients with ulcerative colitis. International Journal of Molecular Medicine, 2009, 23, 521-7.	1.8	56
100	Effect of polymorphisms of IL-17A, -17F and MIF genes on CpG island hyper-methylation (CIHM) in the human gastric mucosa. International Journal of Molecular Medicine, 2009, 24, 563-9.	1.8	14
101	Effect of promoter methylation of multidrug resistance 1 (MDR1) gene in gastric carcinogenesis. Anticancer Research, 2009, 29, 337-41.	0.5	16
102	Effect of polymorphisms in the 3'-untranslated region (3'-UTR) of VEGF gene on gastric pre-malignant condition. Anticancer Research, 2009, 29, 485-9.	0.5	11
103	COMT gene Val158Met polymorphism influences the severity of intestinal metaplasia in H. pylori infected older subjects. Hepato-Gastroenterology, 2009, 56, 411-5.	0.5	1
104	Impact of catechol-O-methyltransferase (COMT) gene polymorphism on promoter methylation status in gastric mucosa. Anticancer Research, 2009, 29, 2857-61.	0.5	7
105	The BB genotype of heat-shock protein (HSP) 70-2 gene is associated with gastric pre-malignant condition in H. pylori-infected older patients. Anticancer Research, 2009, 29, 3453-8.	0.5	9
106	Association between cyclin D1 (CCND1) polymorphism and gastric cancer risk in Japanese population. Hepato-Gastroenterology, 2009, 56, 1232-5.	0.5	2
107	779 TC of CCK-1 intron 1 is associated with postprandial syndrome (PDS) in Japanese male subjects. Hepato-Gastroenterology, 2009, 56, 1245-8.	0.5	9
108	RANTES promoter genotype and gastric cancer risk in a Japanese population. Anticancer Research, 2009, 29, 4265-9.	0.5	6

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109	The Influence of Polymorphisms of Interleukin-17A and Interleukin-17F Genes on the Susceptibility to Ulcerative Colitis. Journal of Clinical Immunology, 2008, 28, 44-49.	2.0	179
110	Homozygous 825T Allele of the GNB3 Protein Influences the Susceptibility of Japanese to Dyspepsia. Digestive Diseases and Sciences, 2008, 53, 642-646.	1.1	47
111	Influence of Peroxisome Proliferator-activated Receptor (PPAR)Î <sup>3</sup> Plo12Ala Polymorphism as a Shared Risk Marker for Both Gastric Cancer and Impaired Fasting Glucose (IFG) in Japanese. Digestive Diseases and Sciences, 2008, 53, 614-621.	1.1	23
112	Toll-like Receptor 2 (TLR) â^196 to 174del Polymorphism in Gastro-duodenal Diseases in Japanese Population. Digestive Diseases and Sciences, 2008, 53, 919-924.	1.1	30
113	Mannan-binding Lectin (MBL) Polymorphism and Gastric Cancer Risk in Japanese Population. Digestive Diseases and Sciences, 2008, 53, 2904-2908.	1.1	15
114	Genetic Polymorphisms of Cyclooxygenase-1 (COX-1) Are Associated with Functional Dyspepsia in Japanese Women. Journal of Women's Health, 2008, 17, 1039-1043.	1.5	27
115	Effect of cyclin D1 (CCND1) polymorphism on gastric premalignant condition. Clinical Chemistry and Laboratory Medicine, 2008, 46, 1696-701.	1.4	2
116	The influence of promoter polymorphism of nuclear factor-erythroid 2-related factor 2 gene on the aberrant DNA methylation in gastric epithelium. Oncology Reports, 2008, , .	1.2	6
117	The Role of Mannan-Binding Lectin (MBL) Gene Polymorphism in Ulcerative Colitis. Journal of Clinical Biochemistry and Nutrition, 2008, 42, 54-58.	0.6	9
118	Aberrant DNA methylation in ulcerative colitis without neoplasia. Hepato-Gastroenterology, 2008, 55, 62-5.	0.5	16
119	COMT gene val158met polymorphism in patients with dyspeptic symptoms. Hepato-Gastroenterology, 2008, 55, 979-82.	0.5	11
120	No association between a genetic variant of the p22PHOX component of NADPH oxidase C242T and ulcerative colitis. Hepato-Gastroenterology, 2008, 55, 1573-7.	0.5	0
121	Serotonin-2A receptor gene T102C polymorphism in patients with dyspeptic symptoms. Hepato-Gastroenterology, 2008, 55, 1921-4.	0.5	4
122	Genetic variant of the p22PHOX component of NADPH oxidase C242T and the incidence of gastric cancer in Japan. Hepato-Gastroenterology, 2008, 55, 2273-6.	0.5	1
123	Risk Prediction of Gastric Cancer by Analysis of Aberrant DNA Methylation in Non-Neoplastic Gastric Epithelium. Digestion, 2007, 75, 54-61.	1.2	38
124	Promoter hypomethylation of protease-activated receptor 2 associated with carcinogenesis in the stomach. Journal of Gastroenterology and Hepatology (Australia), 2007, 22, 943-948.	1.4	11
125	Multi-drug resistance 1 polymorphism is associated with reduced risk of gastric cancer in the Japanese population. Journal of Gastroenterology and Hepatology (Australia), 2007, 22, 1678-1682.	1.4	22
126	Tollâ€like receptor 2 –196 to 174del polymorphism influences the susceptibility of Japanese people to gastric cancer. Cancer Science, 2007, 98, 1790-1794.	1.7	124

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127	Association between genetic polymorphisms in the cyclooxygenase-1 gene promoter and peptic ulcers in Japan. International Journal of Molecular Medicine, 2007, 20, 373-8.	1.8	24
128	Mutation spectrum of TP53 gene predicts clinicopathological features and survival of gastric cancer. Oncotarget, 0, 7, 42252-42260.	0.8	41