

Hiromu Suzuki

List of Publications by Year in descending order

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Version: 2024-02-01

103
papers

7,736
citations

94269

37
h-index

51492

86
g-index

105
all docs

105
docs citations

105
times ranked

10113
citing authors

#	ARTICLE	IF	CITATIONS
1	Characterization of <i>RNF43</i> frameshift mutations that drive Wnt ligand and R-spondin-dependent colon cancer. <i>Journal of Pathology</i> , 2022, 257, 39-52.	2.1	17
2	Genome-wide analysis of mRNA and microRNA expression in colorectal cancer and adjacent normal mucosa. <i>Journal of Pathology: Clinical Research</i> , 2022, , .	1.3	0
3	Cribriform type adenocarcinoma of the colorectum: comprehensive molecular analyses of a distinctive histologic subtype of colorectal cancer. <i>Carcinogenesis</i> , 2022, , .	1.3	2
4	A genome-wide study of the relationship between chromosomal abnormalities and gene expression in colorectal tumors. <i>Genes Chromosomes and Cancer</i> , 2021, 60, 250-262.	1.5	13
5	An Integrated Epigenome and Transcriptome Analysis to Clarify the Effect of Epigenetic Inhibitors on GIST. <i>Anticancer Research</i> , 2021, 41, 2817-2828.	0.5	1
6	A genome-wide analysis of the molecular alterations occurring in the adenomatous and carcinomatous components of the same tumor based on the adenoma-carcinoma sequence. <i>Pathology International</i> , 2021, 71, 582-593.	0.6	5
7	Activated macrophages promote invasion by early colorectal cancer via an interleukin 1 β -serum amyloid A1 axis. <i>Cancer Science</i> , 2021, 112, 4151-4165.	1.7	9
8	Dual EZH2 and G9a inhibition suppresses multiple myeloma cell proliferation by regulating the interferon signal and IRF4-MYC axis. <i>Cell Death Discovery</i> , 2021, 7, 7.	2.0	37
9	Immunohistochemical Examination is Highly Sensitive and Specific for Detection of the V600E BRAF Mutation in Colorectal Serrated Lesions. <i>Applied Immunohistochemistry and Molecular Morphology</i> , 2021, 29, 446-453.	0.6	1
10	DLEU1 promotes oral squamous cell carcinoma progression by activating interferon-stimulated genes. <i>Scientific Reports</i> , 2021, 11, 20438.	1.6	10
11	Comprehensive analyses of microRNA and mRNA expression in colorectal serrated lesions and colorectal cancer with a MSI phenotype. <i>Genes Chromosomes and Cancer</i> , 2021, , .	1.5	3
12	Sessile serrated adenoma/polyp showed rapid malignant transformation in the final 13 months. <i>Digestive Endoscopy</i> , 2020, 32, 979-983.	1.3	10
13	Integrated genetic and epigenetic analysis of cancer-related genes in non-ampullary duodenal adenomas and intramucosal adenocarcinomas. <i>Journal of Pathology</i> , 2020, 252, 330-342.	2.1	16
14	Aggressive variant of splenic marginal zone lymphoma characterized using a cancer panel test and treated with rituximab-containing chemotherapy. <i>Medicine (United States)</i> , 2020, 99, e21938.	0.4	0
15	Traditional serrated adenoma has two distinct genetic pathways for molecular tumorigenesis with potential neoplastic progression. <i>Journal of Gastroenterology</i> , 2020, 55, 846-857.	2.3	12
16	The clinicopathological and molecular features of sporadic gastric foveolar type neoplasia. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2020, 477, 835-844.	1.4	8
17	Dysregulation of microRNA expression during the progression of colorectal tumors. <i>Pathology International</i> , 2020, 70, 633-643.	0.6	4
18	Upregulation of adipocyte enhancer-binding protein 1 in endothelial cells promotes tumor angiogenesis in colorectal cancer. <i>Cancer Science</i> , 2020, 111, 1631-1644.	1.7	26

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19	UHRF1 depletion and HDAC inhibition reactivate epigenetically silenced genes in colorectal cancer cells. <i>Clinical Epigenetics</i> , 2019, 11, 70.	1.8	27
20	DOT1L inhibition blocks multiple myeloma cell proliferation by suppressing IRF4-MYC signaling. <i>Haematologica</i> , 2019, 104, 155-165.	1.7	26
21	Integrated Analysis of the Endoscopic, Pathological and Molecular Characteristics of Colorectal Tumorigenesis. <i>Digestion</i> , 2019, 99, 33-38.	1.2	6
22	Stromal fibroblasts induce metastatic tumor cell clusters via epithelialâ€mesenchymal plasticity. <i>Life Science Alliance</i> , 2019, 2, e201900425.	1.3	48
23	Molecular profiling and comprehensive genome-wide analysis of somatic copy number alterations in gastric intramucosal neoplasias based on microsatellite status. <i>Gastric Cancer</i> , 2018, 21, 765-775.	2.7	12
24	Subtypes of the Type II Pit Pattern Reflect Distinct Molecular Subclasses in the Serrated Neoplastic Pathway. <i>Digestive Diseases and Sciences</i> , 2018, 63, 1920-1928.	1.1	9
25	Evaluation of Urinary DNA Methylation as a Marker for Recurrent Bladder Cancer: A 2-Center Prospective Study. <i>Urology</i> , 2018, 113, 71-78.	0.5	12
26	Molecular profiling and genome-wide analysis based on somatic copy number alterations in advanced colorectal cancers. <i>Molecular Carcinogenesis</i> , 2018, 57, 451-461.	1.3	13
27	Epigenetic silencing of miR-200b is associated with cisplatin resistance in bladder cancer. <i>Oncotarget</i> , 2018, 9, 24457-24469.	0.8	31
28	Frequent downregulation of LRRC26 by epigenetic alterations is involved in the malignant progression of triple-negative breast cancer. <i>International Journal of Oncology</i> , 2018, 52, 1539-1558.	1.4	8
29	Molecular Profiling Based on KRAS/BRAF Mutation, Methylation, and Microsatellite Statuses in Serrated Lesions. <i>Digestive Diseases and Sciences</i> , 2018, 63, 2626-2638.	1.1	14
30	Surface microstructures are associated with mutational intratumoral heterogeneity in colorectal tumors. <i>Journal of Gastroenterology</i> , 2018, 53, 1241-1252.	2.3	5
31	Screening for long noncoding RNAs associated with oral squamous cell carcinoma reveals the potentially oncogenic actions of DLEU1. <i>Cell Death and Disease</i> , 2018, 9, 826.	2.7	46
32	Dysregulation of miRNA in chronic hepatitis B is associated with hepatocellular carcinoma risk after nucleos(t)ide analogue treatment. <i>Cancer Letters</i> , 2018, 434, 91-100.	3.2	18
33	Analysis of the expression of cancer-associated fibroblast- and EMT-related proteins in submucosal invasive colorectal cancer. <i>Journal of Cancer</i> , 2018, 9, 2702-2712.	1.2	18
34	Epigenetic silencing of <i>SMOC1</i> in traditional serrated adenoma and colorectal cancer. <i>Oncotarget</i> , 2018, 9, 4707-4721.	0.8	21
35	Comprehensive molecular analysis based on somatic copy number alterations in intramucosal colorectal neoplasias and early invasive colorectal cancers. <i>Oncotarget</i> , 2018, 9, 22895-22906.	0.8	6
36	Genome-wide analysis of DNA copy number alterations in early and advanced gastric cancers. <i>Molecular Carcinogenesis</i> , 2017, 56, 527-537.	1.3	23

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37	Epigenetic silencing of diacylglycerol kinase gamma in colorectal cancer. <i>Molecular Carcinogenesis</i> , 2017, 56, 1743-1752.	1.3	27
38	Molecular alterations in colorectal adenomas and intramucosal adenocarcinomas defined by high-density single-nucleotide polymorphism arrays. <i>Journal of Gastroenterology</i> , 2017, 52, 1158-1168.	2.3	14
39	Endoscopic and molecular characterization of colorectal sessile serrated adenoma/polyps with cytologic dysplasia. <i>Gastrointestinal Endoscopy</i> , 2017, 86, 1131-1138.e4.	0.5	26
40	Molecular subtypes of colorectal cancers determined by PCR-based analysis. <i>Cancer Science</i> , 2017, 108, 427-434.	1.7	13
41	DNA and Histone Methylation in Colon Cancer. <i>Cancer Drug Discovery and Development</i> , 2017, , 461-487.	0.2	0
42	Identification and characterization of a metastatic suppressor <i>BRMS1L</i> as a target gene of p53. <i>Cancer Science</i> , 2017, 108, 2413-2421.	1.7	17
43	Analysis of the DNA methylation level of cancer-related genes in colorectal cancer and the surrounding normal mucosa. <i>Clinical Epigenetics</i> , 2017, 9, 55.	1.8	35
44	Analysis of molecular alterations in laterally spreading tumors of the colorectum. <i>Journal of Gastroenterology</i> , 2017, 52, 715-723.	2.3	9
45	TET1 Depletion Induces Aberrant CpG Methylation in Colorectal Cancer Cells. <i>PLoS ONE</i> , 2016, 11, e0168281.	1.1	13
46	Relationship Between Noncoding RNA Dysregulation and Epigenetic Mechanisms in Cancer. <i>Advances in Experimental Medicine and Biology</i> , 2016, 927, 109-135.	0.8	18
47	Molecular differences in the microsatellite stable phenotype between left-sided and right-sided colorectal cancer. <i>International Journal of Cancer</i> , 2016, 139, 2493-2501.	2.3	31
48	Clinical prognostic value of DNA methylation in hepatoblastoma: Four novel tumor suppressor candidates. <i>Cancer Science</i> , 2016, 107, 812-819.	1.7	44
49	A genomic screen for long noncoding RNA genes epigenetically silenced by aberrant DNA methylation in colorectal cancer. <i>Scientific Reports</i> , 2016, 6, 26699.	1.6	34
50	Clinicopathological and molecular alterations in early gastric cancers with the microsatellite instability-high phenotype. <i>International Journal of Cancer</i> , 2016, 138, 1689-1697.	2.3	34
51	Epigenetic activation of LY6K predicts the presence of metastasis and poor prognosis in breast carcinoma. <i>Oncotarget</i> , 2016, 7, 55677-55689.	0.8	11
52	Low-Frequency IL23R Coding Variant Associated with Crohn's Disease Susceptibility in Japanese Subjects Identified by Personal Genomics Analysis. <i>PLoS ONE</i> , 2015, 10, e0137801.	1.1	8
53	Comparative analysis of methods to determine DNA methylation levels of a tumor-related microRNA gene. <i>Analytical Biochemistry</i> , 2015, 484, 66-71.	1.1	4
54	Aberrant TET1 Methylation Closely Associated with CpG Island Methylator Phenotype in Colorectal Cancer. <i>Cancer Prevention Research</i> , 2015, 8, 702-711.	0.7	47

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55	Molecular analysis of isolated tumor glands from endometrial endometrioid adenocarcinomas. <i>Pathology International</i> , 2015, 65, 240-249.	0.6	5
56	A Screen for Epigenetically Silenced microRNA Genes in Gastrointestinal Stromal Tumors. <i>PLoS ONE</i> , 2015, 10, e0133754.	1.1	14
57	Epigenetic silencing of <i>NTSR1</i> is associated with lateral and noninvasive growth of colorectal tumors. <i>Oncotarget</i> , 2015, 6, 29975-29990.	0.8	32
58	Epigenetic Regulation of microRNA Genes in Colorectal Cancer. , 2014, , 199-211.		1
59	Colorectal Carcinomas With CpG Island Methylator Phenotype 1 Frequently Contain Mutations in Chromatin Regulators. <i>Gastroenterology</i> , 2014, 146, 530-538.e5.	0.6	76
60	Aberrant methylation of microRNA-34b/c is a predictive marker of metachronous gastric cancer risk. <i>Journal of Gastroenterology</i> , 2014, 49, 1135-1144.	2.3	45
61	Targeting of miR34a "NOTCH1 Axis Reduced Breast Cancer Stemness and Chemoresistance. <i>Cancer Research</i> , 2014, 74, 7573-7582.	0.4	179
62	Analysis of DNA Methylation in Bowel Lavage Fluid for Detection of Colorectal Cancer. <i>Cancer Prevention Research</i> , 2014, 7, 1002-1010.	0.7	38
63	Biological significance of the CpG island methylator phenotype. <i>Biochemical and Biophysical Research Communications</i> , 2014, 455, 35-42.	1.0	86
64	An updated review of gastric cancer in the next-generation sequencing era: Insights from bench to bedside and vice versa. <i>World Journal of Gastroenterology</i> , 2014, 20, 3927.	1.4	72
65	Gastric cancers with microsatellite instability sharing clinical features, chemoresistance and germline MSH6 variants. <i>Clinical Journal of Gastroenterology</i> , 2013, 6, 122-126.	0.4	2
66	Association between genomic alterations and metastatic behavior of colorectal cancer identified by array-based comparative genomic hybridization. <i>Genes Chromosomes and Cancer</i> , 2013, 52, 140-149.	1.5	20
67	Methylation of a Panel of MicroRNA Genes Is a Novel Biomarker for Detection of Bladder Cancer. <i>European Urology</i> , 2013, 63, 1091-1100.	0.9	83
68	Epigenetic alteration and microRNA dysregulation in cancer. <i>Frontiers in Genetics</i> , 2013, 4, 258.	1.1	144
69	Upregulation of miR-196a and <i>HOTAIR</i> Drive Malignant Character in Gastrointestinal Stromal Tumors. <i>Cancer Research</i> , 2012, 72, 1126-1136.	0.4	336
70	A Novel Pit Pattern Identifies the Precursor of Colorectal Cancer Derived From Sessile Serrated Adenoma. <i>American Journal of Gastroenterology</i> , 2012, 107, 460-469.	0.2	164
71	Aberrant Methylation of <i>RASGRF1</i> Is Associated with an Epigenetic Field Defect and Increased Risk of Gastric Cancer. <i>Cancer Prevention Research</i> , 2012, 5, 1203-1212.	0.7	32
72	DNA methylation and microRNA dysregulation in cancer. <i>Molecular Oncology</i> , 2012, 6, 567-578.	2.1	228

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73	Molecular Dissection of Premalignant Colorectal Lesions Reveals Early Onset of the CpG Island Methylator Phenotype. <i>American Journal of Pathology</i> , 2012, 181, 1847-1861.	1.9	74
74	Long noncoding RNA involvement in cancer. <i>BMB Reports</i> , 2012, 45, 604-611.	1.1	178
75	Hypermethylation of Sox17 gene is useful as a molecular diagnostic application in early gastric cancer. <i>Tumor Biology</i> , 2012, 33, 383-393.	0.8	45
76	Emerging links between epigenetic alterations and dysregulation of noncoding RNAs in cancer. <i>Tumor Biology</i> , 2012, 33, 277-285.	0.8	33
77	Role of DNA Methylation in the Development of Diffuse-Type Gastric Cancer. <i>Digestion</i> , 2011, 83, 241-249.	1.2	60
78	Genome-wide Profiling of Chromatin Signatures Reveals Epigenetic Regulation of MicroRNA Genes in Colorectal Cancer. <i>Cancer Research</i> , 2011, 71, 5646-5658.	0.4	156
79	Epigenetic Alteration of DNA in Mucosal Wash Fluid Predicts Invasiveness of Colorectal Tumors. <i>Cancer Prevention Research</i> , 2011, 4, 674-683.	0.7	35
80	Methylation-associated silencing of microRNA-34b/c in gastric cancer and its involvement in an epigenetic field defect. <i>Carcinogenesis</i> , 2010, 31, 2066-2073.	1.3	188
81	IGFBP7 is a p53-responsive gene specifically silenced in colorectal cancer with CpG island methylator phenotype. <i>Carcinogenesis</i> , 2010, 31, 342-349.	1.3	90
82	A Novel Correlation between <i>LINE-1</i> Hypomethylation and the Malignancy of Gastrointestinal Stromal Tumors. <i>Clinical Cancer Research</i> , 2010, 16, 5114-5123.	3.2	83
83	Genomic Screening for Genes Silenced by DNA Methylation Revealed an Association between RASD1 Inactivation and Dexamethasone Resistance in Multiple Myeloma. <i>Clinical Cancer Research</i> , 2009, 15, 4356-4364.	3.2	69
84	Inflammation-Related Aberrant Patterns of DNA Methylation: Detection and Role in Epigenetic Dereglulation of Cancer Cell Transcriptome. <i>Methods in Molecular Biology</i> , 2009, 512, 55-69.	0.4	26
85	Frequent epigenetic inactivation of SFRP genes in hepatocellular carcinoma. <i>Journal of Gastroenterology</i> , 2008, 43, 378-389.	2.3	109
86	Epigenetic Silencing of <i>MicroRNA-34b/c</i> and <i>B-Cell Translocation Gene 4</i> Is Associated with CpG Island Methylation in Colorectal Cancer. <i>Cancer Research</i> , 2008, 68, 4123-4132.	0.4	593
87	A Requirement for DICER to Maintain Full Promoter CpG Island Hypermethylation in Human Cancer Cells. <i>Cancer Research</i> , 2008, 68, 2570-2575.	0.4	46
88	LINE-1 Hypomethylation Is Associated with Increased CpG Island Methylation in <i>Helicobacter pylori</i> -Related Enlarged-Fold Gastritis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 2555-2564.	1.1	62
89	DNA methylation and cancer pathways in gastrointestinal tumors. <i>Pharmacogenomics</i> , 2008, 9, 1917-1928.	0.6	43
90	Frequent epigenetic inactivation of DICKKOPF family genes in human gastrointestinal tumors. <i>Carcinogenesis</i> , 2007, 28, 2459-2466.	1.3	191

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91	Comparing the DNA Hypermethylome with Gene Mutations in Human Colorectal Cancer. PLoS Genetics, 2007, 3, e157.	1.5	307
92	Genetic, epigenetic, and clinicopathologic features of gastric carcinomas with the CpG island methylator phenotype and an association with Epstein-Barr virus. Cancer, 2006, 106, 1467-1479.	2.0	178
93	Roles and causes of abnormal DNA methylation in gastrointestinal cancers. Asian Pacific Journal of Cancer Prevention, 2006, 7, 177-85.	0.5	13
94	The Ras Effector RASSF2 Is a Novel Tumor-Suppressor Gene in Human Colorectal Cancer. Gastroenterology, 2005, 129, 156-169.	0.6	132
95	Epigenetic inactivation of SFRP genes allows constitutive WNT signaling in colorectal cancer. Nature Genetics, 2004, 36, 417-422.	9.4	976
96	GATA-4 and GATA-5 Transcription Factor Genes and Potential Downstream Antitumor Target Genes Are Epigenetically Silenced in Colorectal and Gastric Cancer. Molecular and Cellular Biology, 2003, 23, 8429-8439.	1.1	234
97	Increased expression of T-fimbrin gene after DNA damage in CHO cells and inactivation of T-fimbrin by CpG methylation in human colorectal cancer cells. International Journal of Cancer, 2002, 97, 211-216.	2.3	13
98	Aberrant methylation and histone deacetylation of cyclooxygenase 2 in gastric cancer. International Journal of Cancer, 2002, 97, 272-277.	2.3	75
99	A genomic screen for genes upregulated by demethylation and histone deacetylase inhibition in human colorectal cancer. Nature Genetics, 2002, 31, 141-149.	9.4	820
100	Quantitative DNA methylation analysis by fluorescent polymerase chain reaction single-strand conformation polymorphism using an automated DNA sequencer. Electrophoresis, 2000, 21, 904-908.	1.3	23
101	Frequent hypermethylation of CpG islands and loss of expression of the 14-3-3 σ gene in human hepatocellular carcinoma. Oncogene, 2000, 19, 5298-5302.	2.6	227
102	Distinct methylation pattern and microsatellite instability in sporadic gastric cancer. , 1999, 83, 309-313.		149
103	Epigenetic inactivation of SFRP genes in oral squamous cell carcinoma. International Journal of Oncology, 1992, 32, 1253-1261.	1.4	42