

Yoshinori Murakami

List of Publications by Year in descending order

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Version: 2024-02-01

129
papers

8,676
citations

41323

49
h-index

60583

81
g-index

141
all docs

141
docs citations

141
times ranked

10581
citing authors

#	ARTICLE	IF	CITATIONS
1	A cross-population atlas of genetic associations for 220 human phenotypes. <i>Nature Genetics</i> , 2021, 53, 1415-1424.	9.4	560
2	Overview of the BioBank Japan Project: Study design and profile. <i>Journal of Epidemiology</i> , 2017, 27, S2-S8.	1.1	451
3	TSLC1 is a tumor-suppressor gene in human non-small-cell lung cancer. <i>Nature Genetics</i> , 2001, 27, 427-430.	9.4	402
4	The Polygenic and Monogenic Basis of Blood Traits and Diseases. <i>Cell</i> , 2020, 182, 1214-1231.e11.	13.5	388
5	Trans-ethnic and Ancestry-Specific Blood-Cell Genetics in 746,667 Individuals from 5 Global Populations. <i>Cell</i> , 2020, 182, 1198-1213.e14.	13.5	353
6	The power of genetic diversity in genome-wide association studies of lipids. <i>Nature</i> , 2021, 600, 675-679.	13.7	353
7	Large-scale genome-wide association study in a Japanese population identifies novel susceptibility loci across different diseases. <i>Nature Genetics</i> , 2020, 52, 669-679.	9.4	304
8	Population-specific and trans-ancestry genome-wide analyses identify distinct and shared genetic risk loci for coronary artery disease. <i>Nature Genetics</i> , 2020, 52, 1169-1177.	9.4	206
9	Genetic predisposition to mosaic Y chromosome loss in blood. <i>Nature</i> , 2019, 575, 652-657.	13.7	198
10	Overexpression of a cell adhesion molecule, TSLC1, as a possible molecular marker for acute-type adult T-cell leukemia. <i>Blood</i> , 2004, 105, 1204-1213.	0.6	164
11	Identification of 28 new susceptibility loci for type 2 diabetes in the Japanese population. <i>Nature Genetics</i> , 2019, 51, 379-386.	9.4	164
12	CRTAM determines the CD4+ cytotoxic T lymphocyte lineage. <i>Journal of Experimental Medicine</i> , 2016, 213, 123-138.	4.2	155
13	Involvement of a cell adhesion molecule, TSLC1/IGSF4, in human oncogenesis. <i>Cancer Science</i> , 2005, 96, 543-552.	1.7	152
14	The Tumor Suppressor Protein TSLC1 Is Involved in Cell-Cell Adhesion. <i>Journal of Biological Chemistry</i> , 2002, 277, 31014-31019.	1.6	148
15	Direct association of TSLC1 and DAL-1, two distinct tumor suppressor proteins in lung cancer. <i>Cancer Research</i> , 2002, 62, 5129-33.	0.4	148
16	Cross-sectional analysis of BioBank Japan clinical data: A large cohort of 200,000 patients with 47 common diseases. <i>Journal of Epidemiology</i> , 2017, 27, S9-S21.	1.1	133
17	Improving the trans-ancestry portability of polygenic risk scores by prioritizing variants in predicted cell-type-specific regulatory elements. <i>Nature Genetics</i> , 2020, 52, 1346-1354.	9.4	126
18	Characterizing rare and low-frequency height-associated variants in the Japanese population. <i>Nature Communications</i> , 2019, 10, 4393.	5.8	123

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19	Hematopoietic mosaic chromosomal alterations increase the risk for diverse types of infection. <i>Nature Medicine</i> , 2021, 27, 1012-1024.	15.2	109
20	Leveraging fine-mapping and multipopulation training data to improve cross-population polygenic risk scores. <i>Nature Genetics</i> , 2022, 54, 450-458.	9.4	109
21	A 2-Mb Sequence-Ready Contig Map and a Novel Immunoglobulin Superfamily Gene IGSF4 in the LOH Region of Chromosome 11q23.2. <i>Genomics</i> , 1999, 62, 139-146.	1.3	108
22	Promoter methylation of the TSLC1 gene in advanced lung tumors and various cancer cell lines. <i>International Journal of Cancer</i> , 2003, 107, 53-59.	2.3	105
23	Lung cancer with loss of BRG1/BRM, shows epithelial mesenchymal transition phenotype and distinct histologic and genetic features. <i>Cancer Science</i> , 2013, 104, 266-273.	1.7	103
24	Chromosomal alterations among age-related haematopoietic clones in Japan. <i>Nature</i> , 2020, 584, 130-135.	13.7	102
25	Promoter Methylation of TSLC1 and Tumor Suppression by Its Gene Product in Human Prostate Cancer. <i>Japanese Journal of Cancer Research</i> , 2002, 93, 605-609.	1.7	94
26	Disruption of Spermatogenic Cell Adhesion and Male Infertility in Mice Lacking TSLC1/IGSF4, an Immunoglobulin Superfamily Cell Adhesion Molecule. <i>Molecular and Cellular Biology</i> , 2006, 26, 3610-3624.	1.1	91
27	Identification of CCDC6-RET Fusion in the Human Lung Adenocarcinoma Cell Line, LC-2/ad. <i>Journal of Thoracic Oncology</i> , 2012, 7, 1872-1876.	0.5	90
28	Involvement of TSLC1 in progression of esophageal squamous cell carcinoma. <i>Cancer Research</i> , 2003, 63, 6320-6.	0.4	85
29	Associations of autozygosity with a broad range of human phenotypes. <i>Nature Communications</i> , 2019, 10, 4957.	5.8	84
30	Loss of YAP1 defines neuroendocrine differentiation of lung tumors. <i>Cancer Science</i> , 2016, 107, 1527-1538.	1.7	82
31	GWAS of 165,084 Japanese individuals identified nine loci associated with dietary habits. <i>Nature Human Behaviour</i> , 2020, 4, 308-316.	6.2	80
32	Combined landscape of single-nucleotide variants and copy number alterations in clonal hematopoiesis. <i>Nature Medicine</i> , 2021, 27, 1239-1249.	15.2	78
33	Promoter Methylation of DAL-1/4.1B Predicts Poor Prognosis in Non-Small Cell Lung Cancer. <i>Clinical Cancer Research</i> , 2005, 11, 2954-2961.	3.2	76
34	Hypermethylation of the TSLC1/IGSF4 promoter is associated with tobacco smoking and a poor prognosis in primary nonsmall cell lung carcinoma. <i>Cancer</i> , 2006, 106, 1751-1758.	2.0	75
35	Trans-biobank analysis with 676,000 individuals elucidates the association of polygenic risk scores of complex traits with human lifespan. <i>Nature Medicine</i> , 2020, 26, 542-548.	15.2	74
36	Association of a lung tumor suppressor TSLC1 with MPP3, a human homologue of Drosophila tumor suppressor Dlg. <i>Oncogene</i> , 2003, 22, 6160-6165.	2.6	72

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37	Promoter hypermethylation of the potential tumor suppressor DAL-1/4.1B gene in renal clear cell carcinoma. <i>International Journal of Cancer</i> , 2006, 118, 916-923.	2.3	71
38	Expansion of Cancer Risk Profile for <i>BRCA1</i> and <i>BRCA2</i> Pathogenic Variants. <i>JAMA Oncology</i> , 2022, 8, 871.	3.4	70
39	Isolation of the <i>TSL1</i> and <i>TSL2</i> genes, members of the tumor suppressor <i>TSLC1</i> gene family encoding transmembrane proteins. <i>Oncogene</i> , 2001, 20, 5401-5407.	2.6	66
40	Aberrant expression of tumor suppressors <i>CADM1</i> and <i>4.1B</i> in invasive lesions of primary breast cancer. <i>Breast Cancer</i> , 2012, 19, 242-252.	1.3	66
41	Expression of <i>PRMT5</i> in lung adenocarcinoma and its significance in epithelial-mesenchymal transition. <i>Human Pathology</i> , 2014, 45, 1397-1405.	1.1	66
42	Genome-wide meta-analysis identifies multiple novel loci associated with serum uric acid levels in Japanese individuals. <i>Communications Biology</i> , 2019, 2, 115.	2.0	66
43	Hypermethylation of the <i>TSLC1</i> Gene Promoter in Primary Gastric Cancers and Gastric Cancer Cell Lines. <i>Japanese Journal of Cancer Research</i> , 2002, 93, 857-860.	1.7	65
44	Loss of Tumor Suppressor in Lung Cancer-1 (<i>TSLC1</i>) Expression in Meningioma Correlates with Increased Malignancy Grade and Reduced Patient Survival. <i>Journal of Neuropathology and Experimental Neurology</i> , 2004, 63, 1015-1027.	0.9	61
45	<i>CADM1</i> Interacts with <i>Tiam1</i> and Promotes Invasive Phenotype of Human T-cell Leukemia Virus Type I-transformed Cells and Adult T-cell Leukemia Cells. <i>Journal of Biological Chemistry</i> , 2010, 285, 15511-15522.	1.6	61
46	Dimensionality reduction reveals fine-scale structure in the Japanese population with consequences for polygenic risk prediction. <i>Nature Communications</i> , 2020, 11, 1569.	5.8	58
47	Tumor suppressor <i>CADM1</i> is involved in epithelial cell structure. <i>Biochemical and Biophysical Research Communications</i> , 2009, 390, 977-982.	1.0	56
48	Functional cloning of a tumor suppressor gene, <i>TSLC1</i> , in human non-small cell lung cancer. <i>Oncogene</i> , 2002, 21, 6936-6948.	2.6	54
49	Aberrations of a cell adhesion molecule <i>CADM4</i> in renal clear cell carcinoma. <i>International Journal of Cancer</i> , 2012, 130, 1329-1337.	2.3	54
50	Development of a Highly Sensitive Device for Counting the Number of Disease-Specific Exosomes in Human Sera. <i>Clinical Chemistry</i> , 2018, 64, 1463-1473.	1.5	53
51	GWAS identifies two novel colorectal cancer loci at 16q24.1 and 20q13.12. <i>Carcinogenesis</i> , 2018, 39, 652-660.	1.3	52
52	GWAS of mosaic loss of chromosome Y highlights genetic effects on blood cell differentiation. <i>Nature Communications</i> , 2019, 10, 4719.	5.8	50
53	Expression of <i>TSLC1</i> , a candidate tumor suppressor gene mapped to chromosome 11q23, is downregulated in unfavorable neuroblastoma without promoter hypermethylation. <i>International Journal of Cancer</i> , 2008, 123, 2087-2094.	2.3	48
54	Overview of BioBank Japan follow-up data in 32 diseases. <i>Journal of Epidemiology</i> , 2017, 27, S22-S28.	1.1	47

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55	Fine mapping of the 11q22-q23 tumor suppressive region and involvement of TSLC1 in nasopharyngeal carcinoma. <i>International Journal of Cancer</i> , 2004, 112, 628-635.	2.3	46
56	Transethnic Meta-Analysis of Genome-Wide Association Studies Identifies Three New Loci and Characterizes Population-Specific Differences for Coronary Artery Disease. <i>Circulation Genomic and Precision Medicine</i> , 2020, 13, e002670.	1.6	44
57	The cytoplasmic domain is critical to the tumor suppressor activity of TSLC1 in non-small cell lung cancer. <i>Cancer Research</i> , 2003, 63, 7979-85.	0.4	43
58	Loss of TSLC1 expression in lung adenocarcinoma: Relationships with histological subtypes, sex and prognostic significance. <i>Cancer Science</i> , 2005, 96, 480-486.	1.7	42
59	The ERK signaling target RNF126 regulates anoikis resistance in cancer cells by changing the mitochondrial metabolic flux. <i>Cell Discovery</i> , 2016, 2, 16019.	3.1	40
60	Characterization of KIF11 as a novel prognostic biomarker and therapeutic target for oral cancer. <i>International Journal of Oncology</i> , 2018, 52, 155-165.	1.4	39
61	Genome-wide association study identifies gastric cancer susceptibility loci at 12q24.11 and 20q11.21. <i>Cancer Science</i> , 2018, 109, 4015-4024.	1.7	39
62	Genetic characterization of pancreatic cancer patients and prediction of carrier status of germline pathogenic variants in cancer-predisposing genes. <i>EBioMedicine</i> , 2020, 60, 103033.	2.7	39
63	Increased ectodomain shedding of lung epithelial cell adhesion molecule 1 as a cause of increased alveolar cell apoptosis in emphysema. <i>Thorax</i> , 2014, 69, 223-231.	2.7	37
64	Tumor suppressor cell adhesion molecule 1 (CADM1) is cleaved by a disintegrin and metalloprotease 10 (ADAM10) and subsequently cleaved by β -secretase complex. <i>Biochemical and Biophysical Research Communications</i> , 2012, 417, 462-467.	1.0	36
65	Tumor Suppressor in Lung Cancer (TSLC)1 Suppresses Epithelial Cell Scattering and Tubulogenesis. <i>Journal of Biological Chemistry</i> , 2005, 280, 42164-42171.	1.6	34
66	Genome-wide association meta-analysis identifies GP2 gene risk variants for pancreatic cancer. <i>Nature Communications</i> , 2020, 11, 3175.	5.8	34
67	Trans-Homophilic Interaction of CADM1 Activates PI3K by Forming a Complex with MAGuK-Family Proteins MPP3 and Dlg. <i>PLoS ONE</i> , 2014, 9, e82894.	1.1	34
68	Genetic analysis of right heart structure and function in 40,000 people. <i>Nature Genetics</i> , 2022, 54, 792-803.	9.4	34
69	Expression of a Soluble Isoform of Cell Adhesion Molecule 1 in the Brain and Its Involvement in Directional Neurite Outgrowth. <i>American Journal of Pathology</i> , 2009, 174, 2278-2289.	1.9	33
70	Measles virus selectively blind to signaling lymphocyte activity molecule has oncolytic efficacy against nectin-4-expressing pancreatic cancer cells. <i>Cancer Science</i> , 2016, 107, 1647-1652.	1.7	32
71	Xanthohumol inhibits STAT3 activation pathway leading to growth suppression and apoptosis induction in human cholangiocarcinoma cells. <i>Oncology Reports</i> , 2016, 35, 2065-2072.	1.2	32
72	GWAS of five gynecologic diseases and cross-trait analysis in Japanese. <i>European Journal of Human Genetics</i> , 2020, 28, 95-107.	1.4	32

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73	Genetic and phenotypic landscape of the mitochondrial genome in the Japanese population. <i>Communications Biology</i> , 2020, 3, 104.	2.0	32
74	NECAB3 Promotes Activation of Hypoxia-inducible factor-1 during Normoxia and Enhances Tumourigenicity of Cancer Cells. <i>Scientific Reports</i> , 2016, 6, 22784.	1.6	30
75	Cell adhesion molecule 1 is a new osteoblastic cell adhesion molecule and a diagnostic marker for osteosarcoma. <i>Life Sciences</i> , 2013, 92, 91-99.	2.0	29
76	Random Segregation of DNA Strands in Epidermal Basal Cells. <i>Japanese Journal of Cancer Research</i> , 1989, 80, 637-642.	1.7	27
77	A One-Pot Three-Component Double-Click Method for Synthesis of [67Cu]-Labeled Biomolecular Radiotherapeutics. <i>Scientific Reports</i> , 2017, 7, 1912.	1.6	25
78	A measles virus selectively blind to signaling lymphocytic activation molecule shows anti-tumor activity against lung cancer cells. <i>Oncotarget</i> , 2015, 6, 24895-24903.	0.8	25
79	Adhesion molecule CADM1 contributes to gap junctional communication among pancreatic islet β -cells and prevents their excessive secretion of glucagon. <i>Islets</i> , 2012, 4, 49-55.	0.9	24
80	Control of metastatic niche formation by targeting APBA3/Mint3 in inflammatory monocytes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E4416-E4424.	3.3	24
81	Genomic and transcriptional alterations of cholangiocarcinoma. <i>Journal of Hepato-Biliary-Pancreatic Sciences</i> , 2014, 21, 380-387.	1.4	23
82	Accumulation of genetic alterations and their significance in each primary human cancer and cell line. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 1998, 400, 421-437.	0.4	22
83	Expression of a splicing variant of the <i>CADM1</i> specific to small cell lung cancer. <i>Cancer Science</i> , 2012, 103, 1051-1057.	1.7	22
84	Endogenization and excision of human herpesvirus 6 in human genomes. <i>PLoS Genetics</i> , 2020, 16, e1008915.	1.5	22
85	A Mendelian randomization study identified obesity as a causal risk factor of uterine endometrial cancer in Japanese. <i>Cancer Science</i> , 2020, 111, 4646-4651.	1.7	22
86	Genome-Wide Natural Selection Signatures Are Linked to Genetic Risk of Modern Phenotypes in the Japanese Population. <i>Molecular Biology and Evolution</i> , 2020, 37, 1306-1316.	3.5	22
87	Cell division cycle associated 1 as a novel prognostic biomarker and therapeutic target for oral cancer. <i>International Journal of Oncology</i> , 2016, 49, 1385-1393.	1.4	20
88	Population-based Screening for Hereditary Colorectal Cancer Variants in Japan. <i>Clinical Gastroenterology and Hepatology</i> , 2022, 20, 2132-2141.e9.	2.4	20
89	Dynamic Regulation of a Cell Adhesion Protein Complex Including CADM1 by Combinatorial Analysis of FRAP with Exponential Curve-Fitting. <i>PLoS ONE</i> , 2015, 10, e0116637.	1.1	20
90	Mechanistic insights into ectodomain shedding: susceptibility of CADM1 adhesion molecule is determined by alternative splicing and O-glycosylation. <i>Scientific Reports</i> , 2017, 7, 46174.	1.6	19

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91	Decreased expression of CADM1 and CADM4 are associated with advanced stage breast cancer. <i>Oncology Letters</i> , 2017, 15, 2401-2406.	0.8	19
92	Comparison of effects of UGT1A1*6 and UGT1A1*28 on irinotecan-induced adverse reactions in the Japanese population: analysis of the Biobank Japan Project. <i>Journal of Human Genetics</i> , 2019, 64, 1195-1202.	1.1	19
93	EXOSC9 depletion attenuates P-body formation, stress resistance, and tumorigenicity of cancer cells. <i>Scientific Reports</i> , 2020, 10, 9275.	1.6	18
94	Establishment of highly metastatic KRAS mutant lung cancer cell sublines in long-term three-dimensional low attachment cultures. <i>PLoS ONE</i> , 2017, 12, e0181342.	1.1	17
95	Mint3 depletion restricts tumor malignancy of pancreatic cancer cells by decreasing SKP2 expression via HIF-1. <i>Oncogene</i> , 2020, 39, 6218-6230.	2.6	16
96	Mint3/Apba3 depletion ameliorates severe murine influenza pneumonia and macrophage cytokine production in response to the influenza virus. <i>Scientific Reports</i> , 2016, 6, 37815.	1.6	15
97	Involvement of the SgIGSF/Necl-2 adhesion molecule in degranulation of mesenteric mast cells. <i>Journal of Neuroimmunology</i> , 2007, 184, 209-213.	1.1	14
98	CADM 1 associates with Hippo pathway core kinases; membranous co-expression of CADM 1 and LATS 2 in lung tumors predicts good prognosis. <i>Cancer Science</i> , 2019, 110, 2284-2295.	1.7	14
99	Functional variants in ADH1B and ALDH2 are non-additively associated with all-cause mortality in Japanese population. <i>European Journal of Human Genetics</i> , 2020, 28, 378-382.	1.4	14
100	<i>HLA-B*51:01</i> and <i>CYP2C9*3</i> Are Risk Factors for Phenytoin-Induced Eruption in the Japanese Population: Analysis of Data From the Biobank Japan Project. <i>Clinical Pharmacology and Therapeutics</i> , 2020, 107, 1170-1178.	2.3	13
101	Usefulness of circulating tumor DNA by targeting human papilloma virus-derived sequences as a biomarker in p16-positive oropharyngeal cancer. <i>Scientific Reports</i> , 2022, 12, 572.	1.6	13
102	Transcriptional regulation of the CADM1 gene by retinoic acid during the neural differentiation of murine embryonal carcinoma P19 cells. <i>Genes To Cells</i> , 2011, 16, 791-802.	0.5	12
103	Mathematical analysis of gefitinib resistance of lung adenocarcinoma caused by MET amplification. <i>Biochemical and Biophysical Research Communications</i> , 2019, 511, 544-550.	1.0	11
104	Mint3 in bone marrow-derived cells promotes lung metastasis in breast cancer model mice. <i>Biochemical and Biophysical Research Communications</i> , 2017, 490, 688-692.	1.0	10
105	Reciprocal expression of trefoil factor-1 and thyroid transcription factor-1 in lung adenocarcinomas. <i>Cancer Science</i> , 2020, 111, 2183-2195.	1.7	10
106	Identification of two novel breast cancer loci through large-scale genome-wide association study in the Japanese population. <i>Scientific Reports</i> , 2019, 9, 17332.	1.6	9
107	Genome-wide association study (GWAS) of ovarian cancer in Japanese predicted regulatory variants in 22q13.1. <i>PLoS ONE</i> , 2018, 13, e0209096.	1.1	8
108	Quantitative Analysis of Interaction Between CADM1 and Its Binding Cell-Surface Proteins Using Surface Plasmon Resonance Imaging. <i>Frontiers in Cell and Developmental Biology</i> , 2018, 6, 86.	1.8	8

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109	Susceptibility loci and polygenic architecture highlight population specific and common genetic features in inguinal hernias. <i>EBioMedicine</i> , 2021, 70, 103532.	2.7	8
110	Progression of Pulmonary Emphysema and Continued Increase in Ectodomain Shedding of Cell Adhesion Molecule 1 After Cessation of Cigarette Smoke Exposure in Mice. <i>Frontiers in Cell and Developmental Biology</i> , 2018, 6, 52.	1.8	7
111	Fine Mapping of the Major Histocompatibility Complex Region and Association of the HLA-B*52:01 Allele With Cervical Cancer in Japanese Women. <i>JAMA Network Open</i> , 2020, 3, e2023248.	2.8	7
112	Detection of Lung Tumors in Mice Using a 1-Tesla Compact Magnetic Resonance Imaging System. <i>PLoS ONE</i> , 2014, 9, e94945.	1.1	7
113	CADM1 promotes malignant features of small-cell lung cancer by recruiting 4.1R to the plasma membrane. <i>Biochemical and Biophysical Research Communications</i> , 2021, 534, 172-178.	1.0	6
114	Short somatic alterations at the site of copy number variation in breast cancer. <i>Cancer Science</i> , 2021, 112, 444-453.	1.7	6
115	Circulating Tumor DNA Harboring the <i>BRAF</i> ^{V600E} Mutation May Predict Poor Outcomes of Primary Papillary Thyroid Cancer Patients. <i>Thyroid</i> , 2021, 31, 1822-1828.	2.4	6
116	Genetic alterations in human pancreatic cancer. <i>Journal of Hepato-Biliary-Pancreatic Surgery</i> , 1997, 4, 283-290.	2.0	5
117	CADM1 suppresses c-Src activation by binding with Cbp on membrane lipid rafts and intervenes colon carcinogenesis. <i>Biochemical and Biophysical Research Communications</i> , 2020, 529, 854-860.	1.0	5
118	Genome-wide association study of colorectal polyps identified highly overlapping polygenic architecture with colorectal cancer. <i>Journal of Human Genetics</i> , 2022, 67, 149-156.	1.1	5
119	Expression profile of CADM1 and CADM4 in triple negative breast cancer with primary systemic therapy. <i>Oncology Letters</i> , 2018, 17, 921-926.	0.8	4
120	Family trees representing the finitely proliferative nature of cultured rat liver cells.. <i>Cell Structure and Function</i> , 1983, 8, 293-301.	0.5	4
121	Pharmacological inhibition of Mint3 attenuates tumour growth, metastasis, and endotoxic shock. <i>Communications Biology</i> , 2021, 4, 1165.	2.0	4
122	<i>Trans</i> homophilic interaction of CADM1 promotes organ infiltration of T cell lymphoma by adhesion to vascular endothelium. <i>Cancer Science</i> , 2022, , .	1.7	4
123	Mathematical modeling and analysis of ErbB3 and EGFR dimerization process for the gefitinib resistance. <i>JSIAM Letters</i> , 2018, 10, 33-36.	0.3	3
124	Highlights of topic "Etiology and epidemiology of cholangiocarcinoma". <i>Journal of Hepato-Biliary-Pancreatic Sciences</i> , 2014, 21, 299-300.	1.4	2
125	Mint3 is dispensable for pancreatic and kidney functions in mice. <i>Biochemistry and Biophysics Reports</i> , 2020, 24, 100872.	0.7	2
126	A case of an elderly patient with high-grade colorectal cancer in poor general condition who showed near complete response to chemotherapy and achieved long-term survival. <i>International Journal of Surgery Case Reports</i> , 2019, 58, 186-189.	0.2	1

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127	Preface to topic "International collaboration to control cholangiocarcinoma", Journal of Hepato-Biliary-Pancreatic Sciences, 2014, 21, 297-298.	1.4	0
128	Highlights of topic "Biochemical and molecular pathological aspects of cholangiocarcinoma", Journal of Hepato-Biliary-Pancreatic Sciences, 2014, 21, 359-361.	1.4	0
129	Mathematical Modeling of the Dimerization of EGFR and ErbB3 in Lung Adenocarcinoma. Springer Proceedings in Mathematics and Statistics, 2021, , 195-202.	0.1	0