Mitsuru Emi

List of Publications by Year in descending order

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161 papers	6,210 citations	41 h-index	95218 68 g-index
165	165	165	6484
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Risk prediction for metastasis of clear cell renal cell carcinoma using digital multiplex ligationâ€dependent probe amplification. Cancer Science, 2022, 113, 297-307.	1.7	4
2	Medical and Surgical Care of Patients With Mesothelioma and Their Relatives Carrying Germline BAP1 Mutations. Journal of Thoracic Oncology, 2022, 17, 873-889.	0.5	44
3	Recent Advances in the Genomic and Proteomic Researches on Mesothelioma: What Are Novel Insights into Mesothelioma Biology?. Respiratory Disease Series, 2021, , 137-149.	0.1	O
4	Mesothelioma developing in carriers of inherited genetic mutations. Translational Lung Cancer Research, 2020, 9, S67-S76.	1.3	19
5	Heterozygous germline <i>BLM</i> mutations increase susceptibility to asbestos and mesothelioma. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 33466-33473.	3.3	30
6	A Subset of Mesotheliomas With Improved Survival Occurring in Carriers of <i>BAP1</i> and Other Germline Mutations. Journal of Clinical Oncology, 2018, 36, 3485-3494.	0.8	104
7	Whole-exome sequencing and digital PCR identified a novel compound heterozygous mutation in the NPHP1 gene in a case of Joubert syndrome and related disorders. BMC Medical Genetics, 2017, 18, 37.	2.1	7
8	Uterine Leiomyoma, Clinical Oncology. , 2017, , 4756-4759.		0
9	High-density array-CGH with targeted NGS unmask multiple noncontiguous minute deletions on chromosome 3p21 in mesothelioma. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 13432-13437.	3.3	130
10	Somatic alteration and depleted nuclear expression of BAP 1 in human esophageal squamous cell carcinoma. Cancer Science, 2015, 106, 1118-1129.	1.7	20
11	Combined Genetic and Genealogic Studies Uncover a Large BAP1 Cancer Syndrome Kindred Tracing Back Nine Generations to a Common Ancestor from the 1700s. PLoS Genetics, 2015, 11, e1005633.	1.5	76
12	Frequent genomic rearrangements of BRCA1 associated protein-1 (BAP1) gene in Japanese malignant mesotheliomaâ€"characterization of deletions at exon level. Journal of Human Genetics, 2015, 60, 647-649.	1.1	9
13	High Incidence of Somatic BAP1 Alterations in Sporadic Malignant Mesothelioma. Journal of Thoracic Oncology, 2015, 10, 565-576.	0.5	282
14	Uterine Leiomyoma, Clinical Oncology. , 2014, , 1-5.		0
15	Segmental copy number loss in the region of <i>Semaphorin 4D</i> gene in patients with acetabular dysplasia. Journal of Orthopaedic Research, 2013, 31, 957-961.	1.2	11
16	Genome-wide copy number analysis in primary breast cancer. Expert Opinion on Therapeutic Targets, 2012, 16, S31-S35.	1.5	22
17	Frequent inactivation of the <scp><i>BAP1</i></scp> gene in epithelioidâ€type malignant mesothelioma. Cancer Science, 2012, 103, 868-874.	1.7	159
18	Segmental Copy Number Loss of SFMBT1 Gene in Elderly Individuals with Ventriculomegaly: A Community-Based Study. Internal Medicine, 2011, 50, 297-303.	0.3	33

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19	Copy number loss of (src homology 2 domain containing)-transforming protein 2 (SHC2) gene: discordant loss in monozygotic twins and frequent loss in patients with multiple system atrophy. Molecular Brain, 2011, 4, 24.	1.3	37
20	Frequent Loss of Genome Gap Region in 4p16.3 Subtelomere in Early-Onset Type 2 Diabetes Mellitus. Experimental Diabetes Research, 2011, 2011, 1-10.	3.8	9
21	Uterine Leiomyoma, Clinical Oncology. , 2011, , 3862-3865.		0
22	HNF1B alterations associated with congenital anomalies of the kidney and urinary tract. Pediatric Nephrology, 2010, 25, 1073-1079.	0.9	76
23	Various types of <i>LRP5</i> mutations in four patients with osteoporosisâ€pseudoglioma syndrome: Identification of a 7.2â€kb microdeletion using oligonucleotide tiling microarray. American Journal of Medical Genetics, Part A, 2010, 152A, 133-140.	0.7	31
24	The novel and independent association between single-point SNP of NPHP4 gene and renal function in non-diabetic Japanese population: the Takahata study. Journal of Human Genetics, 2010, 55, 791-795.	1.1	7
25	Segmental copy-number gain within the region of isopentenyl diphosphate isomerase genes in sporadic amyotrophic lateral sclerosis. Biochemical and Biophysical Research Communications, 2010, 402, 438-442.	1.0	17
26	Genetic polymorphisms of paraoxonase-1 are associated with chronic kidney disease in Japanese women. Kidney International, 2009, 76, 183-189.	2.6	12
27	Nucleotide variations in genes encoding carbonic anhydrase 8 and 10 associated with femoral bone mineral density in Japanese female with osteoporosis. Journal of Bone and Mineral Metabolism, 2009, 27, 213-216.	1.3	14
28	Association of CC chemokine ligand 5 genotype with urinary albumin excretion in the non-diabetic Japanese general population: the Takahata study. Journal of Human Genetics, 2008, 53, 267-274.	1.1	10
29	Association of the PIK3C2G gene polymorphisms with type 2 DM in a Japanese population. Biochemical and Biophysical Research Communications, 2008, 365, 466-471.	1.0	52
30	Salt consumption-dependent association of the GNB3 gene polymorphism with type 2 DM. Biochemical and Biophysical Research Communications, 2008, 374, 576-580.	1.0	22
31	Association of a single-nucleotide variation (A1330V) in the low-density lipoprotein receptor-related protein 5 gene (LRP5) with bone mineral density in adult Japanese women. Bone, 2007, 40, 997-1005.	1.4	49
32	Genetic Association of Low-density Lipoprotein Receptor-related Protein 2 (LRP2) with Plasma Lipid Levels. Journal of Atherosclerosis and Thrombosis, 2007, 14, 310-316.	0.9	16
33	Overexpressed in anaplastic thyroid carcinoma-1 (OEATC-1) as a novel gene responsible for anaplastic thyroid carcinoma. Cancer, 2005, 103, 1785-1790.	2.0	45
34	Association of nucleotide variations in the apolipoprotein B48 receptor gene (APOB48R) with hypercholesterolemia. Journal of Human Genetics, 2005, 50, 203-209.	1.1	8
35	Association of single nucleotide polymorphisms in the promoter region of the pro-opiomelanocortin gene (POMC) with low bone mineral density in adult women. Journal of Human Genetics, 2005, 50, 235-240.	1.1	7
36	Nucleotide variations in genes encoding plasminogen activator inhibitor-2 and serine proteinase inhibitor B10 associated with prostate cancer. Journal of Human Genetics, 2005, 50, 507-515.	1.1	11

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37	Upregulation and Overexpression of DVL1, the Human Counterpart of the Drosophila Dishevelled Gene, in Prostate Cancer. Tumori, 2005, 91, 546-551.	0.6	42
38	G-substrate gene promoter SNP (-1323T>C) modifies plasma total cholesterol and triglyceride phenotype in familial hypercholesterolemia: Intra-familial association study in an eight-generation hyperlipidemic kindred. Geriatrics and Gerontology International, 2004, 4, 71-76.	0.7	0
39	Association of a single-nucleotide polymorphism in the promoter region of leukemia inhibitory factor receptor gene with low bone mineral density in adult women. Geriatrics and Gerontology International, 2004, 4, 245-249.	0.7	1
40	Down-regulation of members of glycolipid-enriched membrane raft gene family, MAL and BENE, in cervical squamous cell cancers. Journal of Obstetrics and Gynaecology Research, 2004, 30, 53-58.	0.6	32
41	Expression profiling to predict postoperative prognosis for estrogen receptor-negative breast cancers by analysis of 25,344 genes on a cDNA microarray. Cancer Science, 2004, 95, 218-225.	1.7	190
42	Suppressor of cytokine signalling-1 gene silencing in acute myeloid leukaemia and human haematopoietic cell lines. British Journal of Haematology, 2004, 126, 726-735.	1.2	68
43	Association of Multiple Nucleotide Variations in the Pituitary Glutaminyl Cyclase Gene (QPCT) With Low Radial BMD in Adult Women. Journal of Bone and Mineral Research, 2004, 19, 1296-1301.	3.1	39
44	Hypercholesterolemia associated with splice-junction variation of inter-α-trypsin inhibitor heavy chain 4 (ITIH4) gene. Journal of Human Genetics, 2004, 49, 24-28.	1.1	25
45	Soluble epoxide hydrolase variant (Glu287Arg) modifies plasma total cholesterol and triglyceride phenotype in familial hypercholesterolemia: intrafamilial association study in an eight-generation hyperlipidemic kindred. Journal of Human Genetics, 2004, 49, 29-34.	1.1	78
46	Up-regulation of transcriptional factor E2F1 in papillary and anaplastic thyroid cancers. Journal of Human Genetics, 2004, 49, 312-318.	1.1	41
47	Amplification, up-regulation and over-expression of C3G (CRK SH3 domain-binding guanine) Tj ETQq1 1 0.78431 290-295.	4 rgBT /O [.] 1.1	
48	Functional impairment of two novel mutations detected in lipoprotein-associated phospholipase A2 (Lp-PLA2) deficiency patients. Journal of Human Genetics, 2004, 49, 302-307.	1.1	38
49	The important role for βVLDLs binding at the fourth cysteine of first ligand-binding domain in the low-density lipoprotein receptor. Journal of Human Genetics, 2004, 49, 622-628.	1.1	4
50	Association of a single-nucleotide polymorphism in low-density lipoprotein receptor-related protein 5 gene with bone mineral density. Journal of Bone and Mineral Metabolism, 2004, 22, 341-5.	1.3	77
51	Natural Selection and Population History in the Human Angiotensinogen Gene (AGT): 736 Complete AGT Sequences in Chromosomes from Around the World. American Journal of Human Genetics, 2004, 74, 898-916.	2.6	122
52	Association of a Haplotype (196Phe/532Ser) in the Interleukin-1-Receptor-Associated Kinase (IRAK1) Gene With Low Radial Bone Mineral Density in Two Independent Populations. Journal of Bone and Mineral Research, 2003, 18, 419-423.	3.1	26
53	Association of Molecular Variants, Haplotypes, and Linkage Disequilibrium Within the Human Vitamin D-Binding Protein (DBP) Gene With Postmenopausal Bone Mineral Density. Journal of Bone and Mineral Research, 2003, 18, 1642-1649.	3.1	43
54	Association of natural tooth loss with genetic variation at the human matrix Gla protein locus in elderly women. Journal of Human Genetics, 2003, 48, 288-292.	1.1	8

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55	Hypertriglyceridemia associated with amino acid variation Asn985Tyr of the RP1 gene. Journal of Human Genetics, 2003, 48, 305-308.	1.1	10
56	Association of genetic variation of the RIL gene, encoding a PDZ-LIM domain protein and localized in 5q31.1, with low bone mineral density in adult Japanese women. Journal of Human Genetics, 2003, 48, 342-345.	1.1	24
57	A promoter SNP (–1323T>C) in G-substrate gene (GSBS) correlates with hypercholesterolemia. Journal of Human Genetics, 2003, 48, 447-450.	1.1	16
58	Hypermethylation associated with inactivation of the SOCS-1 gene, a JAK/STAT inhibitor, in human hepatoblastomas. Journal of Human Genetics, 2003, 48, 0065-0069.	1.1	70
59	Growth hormone receptor variant (L526I) modifies plasma HDL cholesterol phenotype in familial hypercholesterolemia: Intra-familial association study in an eight-generation hyperlipidemic kindred. American Journal of Medical Genetics Part A, 2003, 121A, 136-140.	2.4	16
60	DNA alterations during multi-step development of human hepatocellular carcinomas revealed by laser capture microdissection. Hepatology Research, 2003, 26, 199-208.	1.8	3
61	Association of natural tooth loss with genetic variation at the SRC locus among elderly women in Japan. Geriatrics and Gerontology International, 2003, 3, 159-163.	0.7	1
62	Amplification, up-regulation and over-expression of DVL-1, the human counterpart of the Drosophila disheveled gene, in primary breast cancers. Cancer Science, 2003, 94, 515-518.	1.7	116
63	Association of a Trp16Ser variation in the gonadotropin releasing hormone signal peptide with bone mineral density, revealed by SNP-dependent PCR typing. Bone, 2003, 32, 185-190.	1.4	22
64	Association of a promoter haplotype (\hat{a} °1542G/ \hat{a} °525C) in the tumor necrosis factor receptor associated factor-interacting protein gene with low bone mineral density in Japanese women. Bone, 2003, 33, 237-241.	1.4	8
65	Up-regulation and overproduction of DVL-1, the human counterpart of the Drosophila dishevelled gene, in cervical squamous cell carcinoma. Oncology Reports, 2003, 10, 1219-23.	1.2	66
66	Accuracy of Genotyping for Single Nucleotide Polymorphisms by a Microarray-Based Single Nucleotide Polymorphism Typing Method Involving Hybridization of Short Allele-Specific Oligonucleotides. DNA Research, 2002, 9, 59-62.	1.5	15
67	Nucleotide Diversity and Haplotype Structure of the Human Angiotensinogen Gene in Two Populations. American Journal of Human Genetics, 2002, 70, 108-123.	2.6	118
68	Down-regulation of a novel gene, DRLM, in human liver malignancy from 4q22 that encodes a NAP-like protein. Gene, 2002, 296, 171-177.	1.0	20
69	Association of allelic loss at 8p22 with poor prognosis among breast cancer cases treated with high-dose adjuvant chemotherapy. Cancer Letters, 2002, 180, 75-82.	3.2	116
70	Combined hypermethylation and chromosome loss associated with inactivation of SSI-1/SOCS-1/JAB gene in human hepatocellular carcinomas. Cancer Letters, 2002, 186, 59-65.	3.2	57
71	Differential expression of multiple isoforms of the RNAs involved in a papillary thyroid carcinoma. Genes Chromosomes and Cancer, 2002, 35, 30-37.	1.5	25
72	Correlation of allelic losses and clinicopathological factors in 504 primary breast cancers. Breast Cancer, 2002, 9, 208-215.	1.3	35

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73	Eight novel mutations and functional impairments of the LDL receptor in familial hypercholesterolemia in the north of Japan. Journal of Human Genetics, 2002, 47, 80-87.	1.1	25
74	Clinical variant of Tangier disease in Japan: mutation of the ABCA1 gene in hypoalphalipoproteinemia with corneal lipidosis. Journal of Human Genetics, 2002, 47, 366-369.	1.1	25
75	Head-to-head juxtaposition of Fas-associated phosphatase-1 (FAP-1) and c-Jun NH2-terminal kinase 3 (JNK3) genes: genomic structure and seven polymorphisms of the FAP-1 gene. Journal of Human Genetics, 2002, 47, 0614-0619.	1.1	21
76	Interaction between the LDL-receptor gene bearing a novel mutation and a variant in the apolipoprotein A-II promoter: molecular study in a 1135-member familial hypercholesterolemia kindred. Journal of Human Genetics, 2002, 47, 0656-0664.	1.1	37
77	Ethnic difference in contribution of alleles of the interleukin-1 receptor antagonist gene to predisposition to osteoporosis. Geriatrics and Gerontology International, 2002, 2, 87-90.	0.7	1
78	Association of amino acid variation (Trp64Arg) in the beta3-adrenergic receptor gene with bone mineral density. Geriatrics and Gerontology International, 2002, 2, 138-142.	0.7	2
79	Differentially Regulated Genes as Putative Targets of Amplifications at 20q in Ovarian Cancers. Japanese Journal of Cancer Research, 2002, 93, 1114-1122.	1.7	54
80	TFDP1, CUL4A, and CDC16 identified as targets for amplification at 13q34 in hepatocellular carcinomas. Hepatology, 2002, 35, 1476-1484.	3.6	148
81	Inactivation of SSI-1, a JAK/STAT inhibitor, in human hepatocellular carcinomas, as revealed by two-dimensional electrophoresis. Journal of Hepatology, 2001, 34, 416-421.	1.8	28
82	Identification, tissue expression, and chromosomal position of a novel gene encoding human ubiquitin-conjugating enzyme E2-230k. Gene, 2001, 267, 95-100.	1.0	19
83	An 8-cM interstitial deletion on 4q21-q22 in DNA from an infant with hepatoblastoma overlaps with a commonly deleted region in adult liver cancers. American Journal of Medical Genetics Part A, 2001, 103, 176-180.	2.4	15
84	Three aberrant splicing variants of the HMGIC gene transcribed in uterine leiomyomas. Genes Chromosomes and Cancer, 2001, 30, 212-217.	1.5	11
85	Fusion of a Sequence fromHEI10(14q11) to theHMGICGene at 12ql5 in a Uterine Leiomyoma. Japanese Journal of Cancer Research, 2001, 92, 135-139.	1.7	36
86	Down-regulation in Multiple Human Cancers of a Novel Gene, DMHC, from 17q25.1 That Encodes an Integral Membrane Protein. Japanese Journal of Cancer Research, 2001, 92, 417-422.	1.7	1
87	Association of Allelic Losses at 3p25.1, 13q12, or 17p13.3 with Poor Prognosis in Breast Cancers with Lymph Node Metastasis. Japanese Journal of Cancer Research, 2001, 92, 1199-1206.	1.7	5
88	Allelotyping of Follicular Thyroid Carcinoma: Frequent Allelic Losses in Chromosome Arms 7q, 11p, and 22q. Journal of Clinical Endocrinology and Metabolism, 2001, 86, 4268-4272.	1.8	38
89	Allelotyping of Follicular Thyroid Carcinoma: Frequent Allelic Losses in Chromosome Arms 7q, 11p, and 22q. Journal of Clinical Endocrinology and Metabolism, 2001, 86, 4268-4272.	1.8	9
90	骨粗鬆症äº^é~²ã®ãŸã,ã®éºä¼åãfžãf¼ã,«ãf¼ã®é–‹ç™º. Japanese Journal of Geriatrics, 2001, 38, 498-500.	0.0	2

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91	Allelic loss at the 8p22 region as a prognostic factor in large and estrogen receptor negative breast carcinomas., 2000, 88, 1410-1416.		20
92	Allelotyping of anaplastic thyroid carcinoma: Frequent allelic losses on $1q$, $9p$, 11 , 17 , $19p$, and $22q$., 2000 , 27 , 244 - 251 .		68
93	Novel gene fusion of COX6C at 8q22-23 to HMGIC at 12q15 in a uterine leiomyoma., 2000, 27, 303-307.		40
94	Identification of a 1-Mb common region at 16q24.1-24.2 deleted in hepatocellular carcinoma. , 2000, 28, 38-44.		21
95	Mapping of Target Regions of Allelic Loss in Primary Breast Cancers to 1-cM Intervals on Genomic Contigs at 6q21 and 6q25.3. Japanese Journal of Cancer Research, 2000, 91, 293-300.	1.7	26
96	Association of bone mineral density with polymorphism of the human matrix Gla protein locus in elderly women. Journal of Bone and Mineral Metabolism, 2000, 18, 27-30.	1.3	26
97	Association of Estrogen Receptor \hat{l}^2 Gene Polymorphism with Bone Mineral Density. Biochemical and Biophysical Research Communications, 2000, 269, 537-541.	1.0	130
98	Loss of heterozygosity at 3p24–p25 as a prognostic factor in breast cancer. Cancer Letters, 2000, 152, 63-69.	3.2	15
99	Fine Localization of a Major Disease-Susceptibility Locus for Diffuse Panbronchiolitis. American Journal of Human Genetics, 2000, 66, 501-507.	2.6	72
100	Allelotyping of anaplastic thyroid carcinoma: Frequent allelic losses on 1q, 9p, 11, 17, 19p, and 22q. Genes Chromosomes and Cancer, 2000, 27, 244.	1.5	3
101	Familial Hypercholesterolemia in Utah Kindred with Novel R103W Mutations in Exon 4 of the LDL Receptor Gene International Heart Journal, 1999, 40, 443-449.	0.6	2
102	Familial Hypercholesterolemia in Utah Kindred with Novel 2412-6 Ins G Mutations in Exon 17 of the LDL Receptor Gene. International Heart Journal, 1999, 40, 435-441.	0.6	4
103	Frameshift Mutations and a Length Polymorphism in thehMSH3Gene and the Spectrum of Microsatellite Instability in Sporadic Colon Cancer. Japanese Journal of Cancer Research, 1999, 90, 1310-1315.	1.7	7
104	PTEN/MMAC1Mutations in Hepatocellular Carcinomas: Somatic Inactivation of Both Alleles in Tumors. Japanese Journal of Cancer Research, 1999, 90, 413-418.	1.7	47
105	Localization of a Target Region of Allelic Loss to a 1-cM Interval on Chromosome 16p.13.13 in Hepatocellular Carcinoma. Japanese Journal of Cancer Research, 1999, 90, 951-956.	1.7	20
106	Two Target Regions of Allelic Loss on Chromosome 9 in Urinary-bladder Cancer. Japanese Journal of Cancer Research, 1999, 90, 957-964.	1.7	14
107	Correlation of allelic loss with poor postoperative survival in breast cancer. Breast Cancer, 1999, 6, 351-356.	1.3	6
108	A common lle796Val polymorphism of the human SREBP cleavage-activating protein (SCAP) gene. Journal of Human Genetics, 1999, 44, 421-422.	1.1	22

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109	Identification of a 1-cM region of common deletion on 13q14 associated with human prostate cancer. Genes Chromosomes and Cancer, 1999, 24, 183-190.	1.5	32
110	Frequent allelic loss at the TOC locus on 17q25.1 in primary breast cancers., 1999, 24, 345-350.		24
111	Localization of tumor suppressor gene associated with distant metastasis of urinary bladder cancer to a 1-Mb interval on 8p22. Genes Chromosomes and Cancer, 1999, 25, 1-5.	1.5	38
112	Fusion of a novel gene, ELKS, to RET due to translocation $t(10;12)(q11;p13)$ in a papillary thyroid carcinoma., 1999, 25, 97-103.		119
113	Identification of a 1-cM region of common deletion on 4q35 associated with progression of hepatocellular carcinoma., 1999, 25, 284-289.		39
114	Allelic loss at 1p34, 13q12, 17p13.3, and 17q21.1 correlates with poor postoperative prognosis in breast cancer., 1999, 26, 134-141.		26
115	Localization of a tumor suppressor gene associated with the progression of human breast carcinoma within a 1-cm interval of 8p22-p23.1., 1999, 85, 447-452.		60
116	Clinicopathologic analysis of BRCA1- or BRCA2-associated hereditary breast carcinoma in Japanese women., 1999, 85, 2200-2205.		77
117	Variant Manifestation of Cowden Disease in Japan: Hamartomatous Polyposis of the Digestive Tract with Mutation of the PTEN Gene. American Journal of Human Genetics, 1999, 64, 308-310.	2.6	32
118	Inactivation of both alleles of the DPC4/SMAD4 gene in advanced colorectal cancers: identification of seven novel somatic mutations in tumors from Japanese patients. Mutation Research - Mutation Research Genomics, 1999, 406, 71-77.	1.2	38
119	Frequent multiplication of chromosomal region 8q24.1 associated with aggressive histologic types of breast cancers. Cancer Letters, 1999, 139, 7-13.	3.2	20
120	Frequent multiplication of chromosome 1q in non-invasive and papillotubular carcinoma of the breast. Cancer Letters, 1999, 141, 21-28.	3.2	4
121	Somatic mutation of the PTEN/MMAC1 gene in breast cancers with microsatellite instability. Cancer Letters, 1999, 144, 9-16.	3.2	12
122	Analysis of the MEN1 gene in sporadic pituitary adenomas from Japanese patients. Cancer Letters, 1999, 144, 85-92.	3.2	17
123	Localization of a tumor suppressor gene associated with the progression of human breast carcinoma within a 1â€cm interval of 8p22–p23.1. Cancer, 1999, 85, 447-452.	2.0	1
124	Clinicopathologic analysis of BRCA1―or BRCA2â€associated hereditary breast carcinoma in Japanese women. Cancer, 1999, 85, 2200-2205.	2.0	5
125	Identification of a 1 M region of common deletion on 4q35 associated with progression of hepatocellular carcinoma. Genes Chromosomes and Cancer, 1999, 25, 284-289.	1.5	5
126	Multiplex Mutation Screening of the BRCAl Gene in 1000 Japanese Breast Cancers. Japanese Journal of Cancer Research, 1998, 89, 12-16.	1.7	22

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127	Somatic Mutations of the PTEN/MMAC1Gene in Fifteen Japanese Endometrial Cancers: Evidence for Inactivation of Both Alleles. Japanese Journal of Cancer Research, 1998, 89, 842-848.	1.7	33
128	Allelic Loss on Chromosome 9q Is Associated with Lymph Node Metastasis of Primary Breast Cancer. Japanese Journal of Cancer Research, 1998, 89, 916-922.	1.7	21
129	Frequent Allelic Loss at 7p14-15 Associated with Aggressive Histologic Types of Breast Cancer. Japanese Journal of Cancer Research, 1998, 89, 533-538.	1.7	9
130	Allelic loss on chromosome 1p is Associated with progression and lymph node metastasis of primary breast carcinoma., 1998, 82, 317-322.		54
131	Mapping of a new target region of allelic loss to a 2-cM interval at 22q13.1 in primary breast cancer. Genes Chromosomes and Cancer, 1998, 21, 108-112.	1.5	78
132	Mapping of a new target region of allelic loss to a 6-cM interval at 21q21 in primary breast cancers. Genes Chromosomes and Cancer, 1998, 23, 244-247.	1.5	28
133	Frequent Allelic Loss at 6q26-27 in Breast Carcinomas of the Solid-tubular Histologic Type. Breast Cancer, 1998, 5, 127-130.	1.3	2
134	A novel missense mutation and frameshift mutations in the type II receptor of transforming growth factor- \hat{l}^2 gene in sporadic colon cancer with microsatellite instability. Mutation Research - Mutation Research Genomics, 1998, 382, 115-120.	1.2	4
135	Contribution of HLA Genes to Genetic Predisposition in Diffuse Panbronchiolitis. American Journal of Respiratory and Critical Care Medicine, 1998, 158, 846-850.	2.5	60
136	Molecular Genetic Diagnosis of a Family with Hypercholesterolemia by a Mismatched PCR-RFLP Method for Genotyping Single Base Substitution of the LDL Receptor Gene International Heart Journal, 1998, 39, 681-686.	0.6	5
137	Familial Hypercholesterolemia Kindred in Utah with Novel C54S Mutations of the LDL Receptor Gene International Heart Journal, 1998, 39, 785-789.	0.6	2
138	Mapping of a Breast Cancer Tumor Suppressor Gene Locus to a 4-cM Interval on Chromosome 18q21. Japanese Journal of Cancer Research, 1997, 88, 959-964.	1.7	32
139	PRLTS Gene Alterations in Human Prostate Cancer. Japanese Journal of Cancer Research, 1997, 88, 389-393.	1.7	17
140	Correlation of allelic losses and clinicopathological factors in primary breast cancers. Breast Cancer, 1997, 4, 243-246.	1.3	14
141	Allelic losses on $18q21$ are associated with progression and metastasis in human prostate cancer. , $1997, 20, 140-147.$		62
142	Detailed deletion mapping of chromosome arm 3p in breast cancers: A 2-cM region on 3p14.3-21.1 and a 5-cM region on 3p24.3-25.1 commonly deleted in tumors. Genes Chromosomes and Cancer, 1997, 20, 268-274.	1.5	52
143	Two distinct commonly deleted regions on chromosome 13q suggest involvement of BRCA2 and retinoblastoma genes in sporadic breast carcinomas., 1996, 78, 1929-1934.		26
144	Mutations in the BRCA1 gene in Japanese breast cancer patients. Human Mutation, 1996, 7, 334-339.	1.1	47

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145	Three distinct commonly deleted regions of chromosome arm 16q in human primary and metastatic prostate cancers. Genes Chromosomes and Cancer, 1996, 17, 225-233.	1.5	118
146	Allelic losses at loci on chromosome 10 are associated with metastasis and progression of human prostate cancer., 1996, 17, 245-253.		63
147	Chromosome mapping of human (ZNF147) and mouse genes for estrogen-responsive finger protein (efp), a member of the RING finger family. Genomics, 1995, 25, 581-583.	1.3	17
148	A human gene that restores the DNA-repair defect in SCID mice is located on 8p11.1?q11.1. Human Genetics, 1994, 93, 21-6.	1.8	34
149	Genetic studies of 457 breast cancers. Clinicopathologic parameters compared with genetic alterations. Cancer, 1994, 74, 2281-2286.	2.0	61
150	A 3-Mb physical map of the chromosome region 8p21.3-p22, including a 600-kb region commonly deleted in human hepatocellular carcinoma, colorectal cancer, and non-small cell lung cancer. Genes Chromosomes and Cancer, 1994, 10, 7-14.	1.5	76
151	Deletion mapping of the short arm of chromosome 8 in non-small cell lung carcinoma. Genes Chromosomes and Cancer, 1993, 7, 85-88.	1.5	67
152	Allelic loss at chromosome band 8p21.3-p22 is associated with progression of hepatocellular carcinoma. Genes Chromosomes and Cancer, 1993, 7, 152-157.	1.5	95
153	A novel metalloprotease/disintegrin–like gene at 17q21.3 is somatically rearranged in two primary breast cancers. Nature Genetics, 1993, 5, 151-157.	9.4	94
154	A Primary Genetic Linkage Map of 14 Polymorphic Loci for the Short Arm of Human Chromosome 8. Genomics, 1993, 15, 530-534.	1.3	10
155	Isolation and mapping of 88 new RFLP markers on human chromosome 8. Genomics, 1992, 13, 1261-1266.	1.3	19
156	Cloning and characterization of a third type of human α-amylase gene, AMY2B. Gene, 1990, 90, 281-286.	1.0	22
157	Lipoprotein(a) and Plasminogen: Linkage Analysis. , 1990, , 129-139.		3
158	Genetic linkage between lipoprotein(a) phenotype and a DNA polymorphism in the plasminogen gene. Genomics, 1988, 3, 230-236.	1.3	98
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