Hiroaki Okamoto

List of Publications by Year in descending order

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		6592	8835
438	26,350	79	145
papers	citations	h-index	g-index

447	447	447	9021
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Consensus proposals for a unified system of nomenclature of hepatitis C virus genotypes. Hepatology, 2005, 42, 962-973.	3.6	1,303
2	A Novel DNA Virus (TTV) Associated with Elevated Transaminase Levels in Posttransfusion Hepatitis of Unknown Etiology. Biochemical and Biophysical Research Communications, 1997, 241, 92-97.	1.0	1,143
3	Transmission of Hepatitis C Virus from Mothers to Infants. New England Journal of Medicine, 1994, 330, 744-750.	13.9	679
4	Full-length sequence of a hepatitis C virus genome having poor homology to reported isolates: Comparative study of four distinct genotypes. Virology, 1992, 188, 331-341.	1.1	603
5	Consensus proposals for classification of the family Hepeviridae. Journal of General Virology, 2014, 95, 2223-2232.	1.3	570
6	Molecular cloning and characterization of a novel DNA virus (TTV) associated with posttransfusion hepatitis of unknown etiology. Hepatology Research, 1998, 10, 1-16.	1.8	541
7	Sporadic acute or fulminant hepatitis E in Hokkaido, Japan, may be food-borne, as suggested by the presence of hepatitis E virus in pig liver as food. Journal of General Virology, 2003, 84, 2351-2357.	1.3	531
8	Infection with Hepatitis GB Virus C in Patients on Maintenance Hemodialysis. New England Journal of Medicine, 1996, 334, 1485-1491.	13.9	461
9	Genetic drift of hepatitis C virus during an 8.2-year infection in a chimpanzee: Variability and stability. Virology, 1992, 190, 894-899.	1.1	409
10	Hepatitis C virus infection in medical personnel after needlestick accident. Hepatology, 1992, 16, 1109-1114.	3.6	402
11	Proposed reference sequences for hepatitis E virus subtypes. Journal of General Virology, 2016, 97, 537-542.	1.3	339
12	The degree of variability in the amino terminal region of the E2/NS1 protein of hepatitis C virus correlates with responsiveness to interferon therapy in viremic patients. Hepatology, 1992, 16, 619-624.	3.6	319
13	Fecal excretion of a nonenveloped DNA virus (TTV) associated with posttransfusion non-A-G hepatitis. , 1998, 56, 128-132.		300
14	Genetic variability and evolution of hepatitis E virus. Virus Research, 2007, 127, 216-228.	1.1	300
15	HCV genotypes in chronic hepatitis C and response to interferon. Lancet, The, 1992, 339, 1543.	6.3	298
16	Influence of the genotypes of hepatitis C virus on the severity of recurrent liver disease after liver transplantation. Gastroenterology, 1995, 108, 1088-1096.	0.6	296
17	Serological detection of hepatitis B virus genotypes by ELISA with monoclonal antibodies to type-specific epitopes in the preS2-region product. Journal of Virological Methods, 1999, 80, 97-112.	1.0	276
18	Fulminant hepatitis B: Induction by hepatitis B virus mutants defective in the precore region and incapable of encoding E antigen. Gastroenterology, 1991, 100, 1087-1094.	0.6	259

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19	Polyphyletic Strains of Hepatitis E Virus Are Responsible for Sporadic Cases of Acute Hepatitis in Japan. Journal of Clinical Microbiology, 2002, 40, 3209-3218.	1.8	253
20	Non-A, non-B hepatitis specific antibodies directed at host-derived epitope: implication for an autoimmune process. Lancet, The, 1990, 336, 1400-1403.	6.3	247
21	Genotypes and titers of hepatitis C virus for predicting response to interferon in patients with chronic hepatitis C. Journal of Medical Virology, 1994, 42, 299-305.	2.5	240
22	Marked Genomic Heterogeneity and Frequent Mixed Infection of TT Virus Demonstrated by PCR with Primers from Coding and Noncoding Regions. Virology, 1999, 259, 428-436.	1.1	237
23	Hepatitis E Virus (HEV) Strains in Serum Samples Can Replicate Efficiently in Cultured Cells Despite the Coexistence of HEV Antibodies: Characterization of HEV Virions in Blood Circulation. Journal of Clinical Microbiology, 2010, 48, 1112-1125.	1.8	229
24	Development and evaluation of an efficient cell-culture system for Hepatitis E virus. Journal of General Virology, 2007, 88, 903-911.	1.3	224
25	Update: proposed reference sequences for subtypes of hepatitis E virus (species Orthohepevirus A). Journal of General Virology, 2020, 101, 692-698.	1.3	221
26	Swine hepatitis E virus strains in Japan form four phylogenetic clusters comparable with those of Japanese isolates of human hepatitis E virus. Journal of General Virology, 2003, 84, 851-862.	1.3	220
27	Analysis of the full-length genome of a hepatitis E virus isolate obtained from a wild boar in Japan that is classifiable into a novel genotype. Journal of General Virology, 2011, 92, 902-908.	1.3	202
28	Hepatitis G Infection in Drug Abusers with Chronic Hepatitis C. New England Journal of Medicine, 1996, 334, 195-196.	13.9	198
29	Correlation between Antiâ€HBc Titers and HBV DNA in Blood Units without Detectable HBsAg. Vox Sanguinis, 1992, 63, 107-111.	0.7	196
30	ORF3 protein of hepatitis E virus is essential for virion release from infected cells. Journal of General Virology, 2009, 90, 1880-1891.	1.3	194
31	Possible risk factors for the transmission of hepatitis E virus and for the severe form of hepatitis E acquired locally in Hokkaido, Japan. Journal of Medical Virology, 2005, 76, 341-349.	2.5	192
32	A Structurally Flexible and Antigenically Variable N-Terminal Domain of the Hepatitis C Virus E2/NS1 Protein: Implication for an Escape from Antibody. Virology, 1993, 195, 297-301.	1.1	191
33	Analysis of the Complete Genome of Indigenous Swine Hepatitis E Virus Isolated in Japan. Biochemical and Biophysical Research Communications, 2001, 289, 929-936.	1.0	188
34	Hepatitis C virus variants from Vietnam are classifiable into the seventh, eighth, and ninth major genetic groups.. Proceedings of the National Academy of Sciences of the United States of America, 1994, 91, 11022-11026.	3.3	187
35	Mutations within the S Gene of Hepatitis B Virus Transmitted from Mothers to Babies Immunized with Hepatitis B Immune Globulin and Vaccine. Pediatric Research, 1992, 32, 264-268.	1.1	186
36	Simultaneous Detection of Immunoglobulin A (IgA) and IgM Antibodies against Hepatitis E Virus (HEV) Is Highly Specific for Diagnosis of Acute HEV Infection. Journal of Clinical Microbiology, 2005, 43, 49-56.	1.8	185

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37	Genomic characterization of TT viruses (TTVs) in pigs, cats and dogs and their relatedness with species-specific TTVs in primates and tupaias. Journal of General Virology, 2002, 83, 1291-1297.	1.3	185
38	Prevalence of Hepatitis E Virus (HEV) Infection in Wild Boars and Deer and Genetic Identification of a Genotype 3 HEV from a Boar in Japan. Journal of Clinical Microbiology, 2004, 42, 5371-5374.	1.8	166
39	Hepatitis E virus egress depends on the exosomal pathway, with secretory exosomes derived from multivesicular bodies. Journal of General Virology, 2014, 95, 2166-2175.	1.3	163
40	Reduced Precore Transcription and Enhanced Core-Pregenome Transcription of Hepatitis B Virus DNA after Replacement of the Precore-Core Promoter with Sequences Associated with e Antigen-Seronegative Persistent Infections. Virology, 1996, 226, 269-280.	1.1	160
41	Cytotoxic T lymphocyte response and viral load in hepatitis C virus infection. Hepatology, 1997, 25, 705-712.	3.6	156
42	Characterization of the Quasi-Enveloped Hepatitis E Virus Particles Released by the Cellular Exosomal Pathway. Journal of Virology, 2017, 91, .	1.5	151
43	History of Discoveries and Pathogenicity of TT Viruses. Current Topics in Microbiology and Immunology, 2009, 331, 1-20.	0.7	150
44	Chronic active hepatitis with hepatitis B virus DNA and antibody against e antigen in the serum. Gastroenterology, 1990, 99, 1113-1119.	0.6	149
45	Prevalence of hepatitis E virus infection among hemodialysis patients in Japan: Evidence for infection with a genotype 3 HEV by blood transfusion. Journal of Medical Virology, 2004, 74, 563-572.	2.5	149
46	Characterization of Japanese swine and human hepatitis E virus isolates of genotype IV with 99â€Š% identity over the entire genome. Journal of General Virology, 2003, 84, 1245-1251.	1.3	148
47	Analysis of the entire genomes of thirteen TT virus variants classifiable into the fourth and fifth genetic groups, isolated from viremic infants. Archives of Virology, 2002, 147, 21-41.	0.9	146
48	Monoclonal antibodies raised against the ORF3 protein of hepatitis E virus (HEV) can capture HEV particles in culture supernatant and serum but not those in feces. Archives of Virology, 2008, 153, 1703-1713.	0.9	140
49	Superinfection of chimpanzees carrying hepatitis C virus of genotype II/1b with that of genotype III/2a or I/1a. Hepatology, 1994, 20, 1131-1136.	3.6	139
50	A PSAP motif in the ORF3 protein of hepatitis E virus is necessary for virion release from infected cells. Journal of General Virology, 2011, 92, 269-278.	1.3	132
51	Antibodies against synthetic oligopeptides deduced from the putative core gene for the diagnosis of hepatitis C virus infection. Hepatology, 1992, 15, 180-186.	3.6	126
52	Infection with an unenveloped DNA Virus (TTV) associated with posttransfusion non-A to G hepatitis in hepatitis patients and healthy blood donors in Thailand. , 1998, 56, 234-238.		123
53	Development of PCR Assays with Nested Primers Specific for Differential Detection of Three Human Anelloviruses and Early Acquisition of Dual or Triple Infection during Infancy. Journal of Clinical Microbiology, 2008, 46, 507-514.	1.8	117
54	Hepatitis B Virus Mutants with Precore-Region Defects in Two Babies with Fulminant Hepatitis and Their Mothers Positive for Antibody to Hepatitis B e Antigen. Pediatric Research, 1991, 29, 5-9.	1.1	115

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55	The Entire Nucleotide Sequence of a TT Virus Isolate from the United States (TUS01): Comparison with Reported Isolates and Phylogenetic Analysis. Virology, 1999, 259, 437-448.	1.1	113
56	Features of Hepatitis E Virus Infection in Japan. Internal Medicine, 2003, 42, 1065-1071.	0.3	111
57	Identification of two distinct genotypes of hepatitis E virus in a Japanese patient with acute hepatitis who had not travelled abroad. Journal of General Virology, 2002, 83, 1931-1940.	1.3	110
58	Inverse relationship between the titre of TT virus DNA and the CD4 cell count in patients infected with HIV. Aids, 2001, 15, 563-570.	1.0	107
59	Excretion into Bile of a Novel Unenveloped DNA Virus (TT Virus) Associated with Acute and Chronic Nonâ€Aâ€“G Hepatitis. Journal of Infectious Diseases, 1999, 179, 1245-1248.	1.9	106
60	Heterogeneous Distribution of TT Virus of Distinct Genotypes in Multiple Tissues from Infected Humans. Virology, 2001, 288, 358-368.	1.1	103
61	A nationwide survey of hepatitis E virus (HEV) infection in wild boars in Japan: identification of boar HEV strains of genotypes 3 and 4 and unrecognized genotypes. Archives of Virology, 2011, 156, 1345-1358.	0.9	103
62	Two distinct subtypes of hepatitis C virus defined by antibodies directed to the putative core protein. Hepatology, 1992, 16, 886-891.	3.6	98
63	Genomic and evolutionary characterization of TT virus (TTV) in tupaias and comparison with species-specific TTVs in humans and non-human primates. Journal of General Virology, 2001, 82, 2041-2050.	1.3	97
64	Defects in the precore region of the HBV genome in patients with chronic hepatitis B after sustained seroconversion from HBeAg to anti-HBe induced spontaneously or with interferon therapy. Hepatology, 1990, 12, 1284-1289.	3.6	96
65	Tumour susceptibility gene 101 and the vacuolar protein sorting pathway are required for the release of hepatitis E virions. Journal of General Virology, 2011, 92, 2838-2848.	1.3	95
66	Prevalence, genotypes, and an isolate (HC-C2) of hepatitis C virus in Chinese patients with liver disease. Journal of Medical Virology, 1993, 40, 254-260.	2.5	94
67	Infection with GB virus C (GBV-C) in patients with chronic liver disease or on maintenance hemodialysis in Indonesia. , 1996, 49, 248-252.		93
68	Circular Double-Stranded Forms of TT Virus DNA in the Liver. Journal of Virology, 2000, 74, 5161-5167.	1.5	93
69	Development and validation of an improved RT-PCR assay with nested universal primers for detection of hepatitis E virus strains with significant sequence divergence. Journal of Virological Methods, 2006, 137, 325-333.	1.0	93
70	A second-generation method of genotyping hepatitis C virus by the polymerase chain reaction with sense and antisense primers deduced from the core gene. Journal of Virological Methods, 1996, 57, 31-45.	1.0	92
71	Features of hepatitis <scp>E</scp> virus infection in humans and animals in <scp>J</scp>apan. Hepatology Research, 2014, 44, 43-58.	1.8	91
72	Suppression of hepatitis C virus RNA by interferon-Î±. Lancet, The, 1990, 336, 245.	6.3	90

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73	Infection with an unenveloped DNA virus (TTV) in patients with acute or chronic liver disease of unknown etiology and in those positive for hepatitis C virus RNA. Journal of Hepatology, 1999, 30, 205-212.	1.8	90
74	Phase II Study of Area Under the Plasma-Concentration-Versus-Time Curveâ€"Based Carboplatin Plus Standard-Dose Intravenous Etoposide in Elderly Patients With Small-Cell Lung Cancer. Journal of Clinical Oncology, 1999, 17, 3540-3545.	0.8	89
75	Risk factors for the development of hepatocellular carcinoma among patients with chronic hepatitis C who achieved a sustained virological response to interferon therapy. Journal of Gastroenterology and Hepatology (Australia), 2005, 20, 752-758.	1.4	89
76	p26 protein and 33-nm particle associated with nucleocapsid of hepatitis C virus recovered from the circulation of infected hosts. Virology, 1992, 191, 431-434.	1.1	86
77	Fulminant Hepatitis E in Japan. New England Journal of Medicine, 2002, 347, 1456-1456.	13.9	86
78	Development and Characterization of a Genotype 4 Hepatitis E Virus Cell Culture System Using a HE-JF5/15F Strain Recovered from a Fulminant Hepatitis Patient. Journal of Clinical Microbiology, 2009, 47, 1906-1910.	1.8	86
79	GBV-C in the aetiology of fulminant hepatitis. Lancet, The, 1996, 347, 120-121.	6.3	83
80	Culture systems for hepatitis E virus. Journal of Gastroenterology, 2013, 48, 147-158.	2.3	83
81	Differentiation of hepatitis B virus genotypes D and E by ELISA using monoclonal antibodies to epitopes on the preS2-region product. Journal of Virological Methods, 2000, 87, 81-89.	1.0	82
82	Replicative Forms of TT Virus DNA in Bone Marrow Cells. Biochemical and Biophysical Research Communications, 2000, 270, 657-662.	1.0	82
83	Prolonged Fecal Shedding of Hepatitis E Virus (HEV) during Sporadic Acute Hepatitis E: Evaluation of Infectivity of HEV in Fecal Specimens in a Cell Culture System. Journal of Clinical Microbiology, 2007, 45, 3671-3679.	1.8	82
84	Mutation Rate of GB Virus C/Hepatitis G Virus over the Entire Genome and in Subgenomic Regions. Virology, 1997, 233, 43-50.	1.1	80
85	Quasispecies of TT Virus (TTV) with Sequence Divergence in Hypervariable Regions of the Capsid Protein in Chronic TTV Infection. Journal of Virology, 1999, 73, 9604-9608.	1.5	80
86	High Prevalence of Antibodies to Hepatitis A and E Viruses and Viremia of Hepatitis B, C, and D Viruses among Apparently Healthy Populations in Mongolia. Vaccine Journal, 2004, 11, 392-398.	2.6	80
87	Hepatitis E virus cell culture models. Virus Research, 2011, 161, 65-77.	1.1	80
88	Prevalence of antibodies to hepatitis E virus among Japanese blood donors: Identification of three blood donors infected with a genotype 3 hepatitis E virus. Journal of Medical Virology, 2004, 73, 554-561.	2.5	77
89	Reactivation of precore mutant hepatitis B virus leading to fulminant hepatic failure following cytotoxic treatment. Digestive Diseases and Sciences, 1992, 37, 1253-1259.	1.1	75
90	Influence of Antibodies to the Hypervariable Region of E2/NS1 Glycoprotein on the Selective Replication of Hepatitis C Virus in Chimpanzees. Virology, 1994, 204, 665-672.	1.1	75

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91	Transmission of HCV between spouses. Lancet, The, 1992, 339, 1059-1060.	6.3	74
92	Fulminant hepatitis related to transmission of hepatitis B variants with precore mutations between spouses. Hepatology, 1992, 16, 31-35.	3.6	74
93	Species-Specific TT Viruses in Humans and Nonhuman Primates and Their Phylogenetic Relatedness. Virology, 2000, 277, 368-378.	1.1	74
94	Contribution of hepatitis C virus to non-A, non-B fulminant hepatitis in Japan. Hepatology, 1994, 19, 829-835.	3.6	73
95	Identification and genomic characterization of a novel human torque teno virus of 3.2â€…kb. Journal of General Virology, 2007, 88, 1939-1944.	1.3	73
96	A nationwide molecular epidemiological study on hepatitis B virus in Indonesia: identification of two novel subgenotypes, B8 and C7. Archives of Virology, 2009, 154, 1047-1059.	0.9	72
97	TT Virus mRNAs Detected in the Bone Marrow Cells from an Infected Individual. Biochemical and Biophysical Research Communications, 2000, 279, 700-707.	1.0	71
98	Genetic Variability and Evolution of Hepatitis E Virus. Viruses, 2019, 11, 456.	1.5	71
99	Genetic Heterogeneity of Hepatitis C Virus. Intervirology, 1994, 37, 68-76.	1.2	70
100	Sporadic acute hepatitis E of a 47-year-old man whose pet cat was positive for antibody to hepatitis E virus. Hepatology Research, 2003, 26, 237-242.	1.8	70
101	Monitoring serum hepatitis C virus (HCV) RNA in patients with HCVâ€infected CD20â€positive Bâ€cell lymphoma undergoing rituximab combination chemotherapy. American Journal of Hematology, 2008, 83, 59-62.	2.0	70
102	Impaired induction of cytotoxic T lymphocytes by antagonism of a weak agonist borne by a variant hepatitis C virus epitope. European Journal of Immunology, 1997, 27, 1782-1787.	1.6	69
103	Correlation between positivity for immunoglobulin A antibodies and viraemia of swine hepatitis E virus observed among farm pigs in Japan. Journal of General Virology, 2005, 86, 1807-1813.	1.3	69
104	Ongoing subclinical infection of hepatitis E virus among blood donors with an elevated alanine aminotransferase level in Japan. Journal of Medical Virology, 2007, 79, 734-742.	2.5	69
105	The membrane on the surface of hepatitis E virus particles is derived from the intracellular membrane and contains trans-Golgi network protein 2. Archives of Virology, 2014, 159, 979-991.	0.9	69
106	Species-Specific TT Viruses and Cross-Species Infection in Nonhuman Primates. Journal of Virology, 2000, 74, 1132-1139.	1.5	68
107	Infection of a Japanese Patient by Genotype 4 Hepatitis E Virus While Traveling in Vietnam. Journal of Clinical Microbiology, 2004, 42, 3883-3885.	1.8	68
108	High prevalence of dual or triple infection of hepatitis B, C, and delta viruses among patients with chronic liver disease in Mongolia. Journal of Medical Virology, 2005, 77, 491-499.	2.5	68

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109	Distinct genotypes of a nonenveloped DNA virus associated with posttransfusion non-A to G hepatitis (TT virus) in plasma and peripheral blood mononuclear cells. , 1999, 57, 252-258.		67
110	Visualization of TT Virus Particles Recovered from the Sera and Feces of Infected Humans. Biochemical and Biophysical Research Communications, 2000, 279, 718-724.	1.0	63
111	Prevalence of antibodies to hepatitis E virus among apparently healthy humans and pigs in Bali, Indonesia: Identification of a pig infected with a genotype 4 hepatitis E virus. Journal of Medical Virology, 2004, 73, 38-44.	2.5	63
112	Construction of an infectious cDNA clone of hepatitis E virus strain JE03-1760F that can propagate efficiently in cultured cells. Journal of General Virology, 2009, 90, 457-462.	1.3	63
113	Molecular characterization of a novel hepatitis E virus (HEV) strain obtained from a wild boar in Japan that is highly divergent from the previously recognized HEV strains. Virus Research, 2014, 180, 59-69.	1.1	62
114	Unique clinical courses of transfusionâ€transmitted hepatitis E in patients with immunosuppression. Transfusion, 2017, 57, 280-288.	0.8	62
115	Production of monoclonal antibodies against hepatitis E virus capsid protein and evaluation of their neutralizing activity in a cell culture system. Archives of Virology, 2008, 153, 657-666.	0.9	61
116	Hepatitis C and B virus infections in populations at low or high risk in Ho Chi Minh and Hanoi, Vietnam. Journal of Gastroenterology and Hepatology (Australia), 1994, 9, 416-419.	1.4	60
117	Characterization of seven genotypes (A to E, G and H) of Hepatitis B virus recovered from Japanese patients infected with human immunodeficiency virus type 1. Journal of Medical Virology, 2005, 76, 24-32.	2.5	60
118	High TT virus load as an independent factor associated with the occurrence of hepatocellular carcinoma among patients with hepatitis C virus-related chronic liver disease. Journal of Medical Virology, 2002, 67, 501-509.	2.5	59
119	A nationwide survey of hepatitis E virus infection in the general population of Japan. Journal of Medical Virology, 2010, 82, 271-281.	2.5	59
120	Hepatitis C virus infection in patients with urticaria. Journal of the American Academy of Dermatology, 1996, 35, 195-198.	0.6	57
121	A549 and PLC/PRF/5 cells can support the efficient propagation of swine and wild boar hepatitis E virus (HEV) strains: demonstration of HEV infectivity of porcine liver sold as food. Archives of Virology, 2012, 157, 235-246.	0.9	57
122	Differences in the entire nucleotide sequence between hepatitis B virus genomes from carriers positive for antibody to hepatitis B e antigen with and without active disease. Journal of Medical Virology, 1994, 44, 96-103.	2.5	56
123	Hepatitis C virus infection in patients with chronic liver disease or chronic renal failure and blood donors in Thailand. Journal of Medical Virology, 1994, 44, 287-292.	2.5	56
124	Molecular investigation of hepatitis E virus infection in patients with acute hepatitis in Kathmandu, Nepal. Journal of Medical Virology, 2003, 69, 207-214.	2.5	56
125	Presence of Antibodies to Hepatitis E Virus in Japanese Pet Cats. Infection, 2004, 32, 57-58.	2.3	56
126	A solid-phase enzyme immunoassay for the common and subtypic determinants of hepatitis B surface antigen with monoclonal antibodies. Journal of Immunological Methods, 1986, 87, 203-210.	0.6	55

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127	A Novel Unenveloped DNA Virus (TT Virus) Associated with Acute and Chronic Non-A to G Hepatitis. Intervirology, 1999, 42, 196-204.	1.2	55
128	Analysis of the full-length genome of genotype 4 hepatitis E virus isolates from patients with fulminant or acute self-limited hepatitis E. Journal of Medical Virology, 2006, 78, 476-484.	2.5	55
129	Distinct changing profiles of hepatitis A and E virus infection among patients with acute hepatitis, patients on maintenance hemodialysis and healthy individuals in Japan. Journal of Medical Virology, 2006, 78, 1015-1024.	2.5	55
130	Taxonomic update for mammalian anelloviruses (family Anelloviridae). Archives of Virology, 2021, 166, 2943-2953.	0.9	55
131	Sequestration of TT Virus of Restricted Genotypes in Peripheral Blood Mononuclear Cells. Journal of Virology, 2000, 74, 10236-10239.	1.5	53
132	Molecular epidemiology of TT virus (TTV) and characterization of two novel TTV genotypes in Indonesia. Archives of Virology, 2001, 146, 1249-1266.	0.9	52
133	Infection of TT virus in patients with idiopathic pulmonary fibrosis. Respiratory Medicine, 2001, 95, 935-942.	1.3	51
134	Epidemiological and clinical study of sporadic acute hepatitis E caused by indigenous strains of hepatitis E virus in Japan compared with acute hepatitis A. Journal of Gastroenterology, 2004, 39, 640-648.	2.3	51
135	Analysis of the full-length genome of hepatitis E virus isolates obtained from wild boars in Japan. Journal of General Virology, 2005, 86, 3321-3326.	1.3	50
136	TT Viruses in Animals. Current Topics in Microbiology and Immunology, 2009, 331, 35-52.	0.7	50
137	Antibody to the receptor for polymerized human serum albumin in acute and persistent infection with hepatitis B virus. Hepatology, 1986, 6, 354-359.	3.6	49
138	HLA DRB1 and DQB1 alleles and haplotypes influencing the progression of hepatitis C. , 1996, 49, 274-278.		49
139	Determination of antibodies to TT virus (TTV) and application to blood donors and patients with post-transfusion non-A to G hepatitis in Japan. Journal of Virological Methods, 1999, 77, 199-206.	1.0	49
140	Transmission of Human TT Virus of Genotype 1a to Chimpanzees with Fecal Supernatant or Serum from Patients with Acute TTV Infection. Biochemical and Biophysical Research Communications, 2000, 278, 470-476.	1.0	49
141	Mutational events during the primary propagation and consecutive passages of hepatitis E virus strain JE03-1760F in cell culture. Virus Research, 2008, 137, 86-96.	1.1	49
142	Demonstration of Sugar Moiety on the Surface of Hepatitis C Virions Recovered from the Circulation of Infected Humans. Virology, 1993, 196, 354-357.	1.1	48
143	A common-source outbreak of fulminant hepatitis B in hemodialysis patients induced by precore mutant. Kidney International, 1995, 48, 1972-1978.	2.6	48
144	Infection with GB virus C (GBV-C) in patients with fulminant hepatitis. Journal of Hepatology, 1996, 25, 842-847.	1.8	48

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145	A Nationwide Survey of Hepatitis E Virus Infection and Chronic Hepatitis E in Liver Transplant Recipients in Japan. EBioMedicine, 2015, 2, 1607-1612.	2.7	48
146	Full-length nucleotide sequence of a Japanese hepatitis C virus isolate (HC-J1) with high homology to USA isolates. Nucleic Acids Research, 1992, 20, 6410-6410.	6.5	47
147	TT virus: virological and genomic characteristics and disease associations. Journal of Gastroenterology, 2001, 36, 519-529.	2.3	46
148	Identification of four novel subgenotypes (C13â€“C16) and two inter-genotypic recombinants (C12/G and) Tj ETQq0 0 0 rgBT /Overlock	1.1	46
149	High carrier rate after hepatitis B virus infection in the elderly. Hepatology, 1993, 18, 768-774.	3.6	45
150	Molecular and serological characterization of sporadic acute hepatitis E in a Japanese patient infected with a genotype III hepatitis E virus in 1993. Journal of General Virology, 2003, 84, 421-427.	1.3	45
151	Serological and molecular studies on subclinical hepatitis E virus infection using periodic serum samples obtained from healthy individuals. Journal of Medical Virology, 2005, 76, 526-533.	2.5	45
152	Virulent Strain of Hepatitis E Virus Genotype 3, Japan. Emerging Infectious Diseases, 2009, 15, 704-709.	2.0	45
153	High prevalence of hepatitis delta virus infection detectable by enzyme immunoassay among apparently healthy individuals in Mongolia. Journal of Medical Virology, 2005, 76, 333-340.	2.5	44
154	Infection with hepatitis A, B, C, and delta viruses among patients with acute hepatitis in Mongolia. Journal of Medical Virology, 2006, 78, 542-550.	2.5	43
155	Determination of the 5â€²-terminal sequence of subgenomic RNA of hepatitis E virus strains in cultured cells. Archives of Virology, 2009, 154, 1945-1951.	0.9	43
156	Molecular analysis of hepatitis E virus from farm rabbits in Inner Mongolia, China and its successful propagation in A549 and PLC/PRF/5 cells. Virus Research, 2012, 170, 126-137.	1.1	43
157	Frequent detection and characterization of hepatitis E virus variants in wild rats (Rattus rattus) in Indonesia. Archives of Virology, 2013, 158, 87-96.	0.9	43
158	Marked genomic heterogeneity of rat hepatitis E virus strains in Indonesia demonstrated on a full-length genome analysis. Virus Research, 2014, 179, 102-112.	1.1	43
159	Rat hepatitis E virus derived from wild rats (Rattus rattus) propagates efficiently in human hepatoma cell lines. Virus Research, 2014, 185, 92-102.	1.1	43
160	Efficient cell culture systems for hepatitis E virus strains in feces and circulating blood. Reviews in Medical Virology, 2011, 21, 18-31.	3.9	42
161	Molecular epidemiology of hepatitis B virus in Indonesia. Journal of Gastroenterology and Hepatology (Australia), 1991, 6, 491-498.	1.4	41
162	Influence of Primers on the Detection of TT Virus DNA by Polymerase Chain Reaction. Journal of Infectious Diseases, 1999, 180, 1750-1751.	1.9	41

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163	TT virus infection duringâ€ƒchildhood. Transfusion, 2002, 42, 892-898.	0.8	41
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165	Imported Hepatitis C Virus Genotypes in Japanese Hemophiliacs. Journal of Infectious Diseases, 1993, 168, 249-250.	1.9	38
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