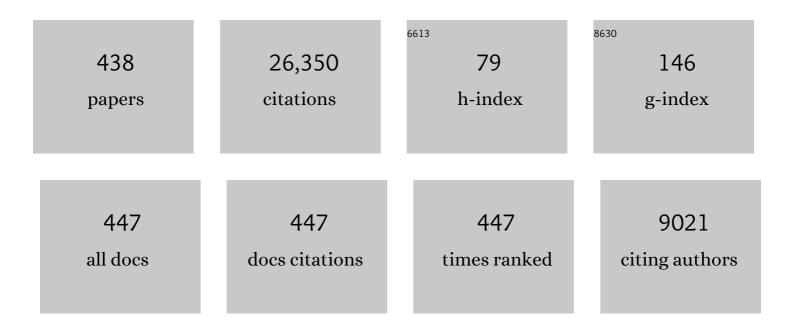
Hiroaki Okamoto

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Regulatory function of interferonâ€inducible 44â€like for hepatitis B virus covalently closed circular DNA in primary human hepatocytes. Hepatology Research, 2022, 52, 141-152.	3.4	4
2	Identification of hepatitis E virus in wild sika deer in Japan. Virus Research, 2022, 308, 198645.	2.2	10
3	Chronic Infection with Hepatitis C Virus Subtype 1 g in a Japanese Patient Successfully Treated with Glecaprevir/Pibrentasvir. Internal Medicine, 2022, , .	0.7	0
4	Development of Recombinant Infectious Hepatitis E Virus Harboring the nanoKAZ Gene and Its Application in Drug Screening. Journal of Virology, 2022, 96, jvi0190621.	3.4	9
5	Favipiravir Inhibits Hepatitis A Virus Infection in Human Hepatocytes. International Journal of Molecular Sciences, 2022, 23, 2631.	4.1	12
6	Genomic characterization and the prevalence of a novel copiparvovirus in wild sika deer (Cervus) Tj ETQq0 0 0 r	gBT_/Overl 2.2	ock 10 Tf 50 !
7	First detection and characterization of rat hepatitis E Virus (HEV-C1) in Japan. Virus Research, 2022, 314, 198766.	2.2	7
8	The Capsid (ORF2) Protein of Hepatitis E Virus in Feces Is C-Terminally Truncated. Pathogens, 2022, 11, 24.	2.8	8
9	Subclinical hepatitis E virus (HEV) infection detected by nucleic acid amplification test on blood donation: short-term positivity for immunoglobulin G class of antibody against HEV. Clinical Journal of Gastroenterology, 2022, 15, 750-754.	0.8	2
10	Hepatitis B, C, and D Virus Infections and AFP Tumor Marker Prevalence Among the Elderly Population in Mongolia: A Nationwide Survey. Journal of Preventive Medicine and Public Health, 2022, 55, 263-272.	1.9	3
11	Evaluation of Potential Anti-Hepatitis A Virus 3C Protease Inhibitors Using Molecular Docking. International Journal of Molecular Sciences, 2022, 23, 6044.	4.1	6
12	Persistent infection with a rabbit hepatitis E virus created by a reverse genetics system. Transboundary and Emerging Diseases, 2021, 68, 615-625.	3.0	10
13	Complete Genome Sequences of Two Rat Pegivirus Strains in Indonesia. Microbiology Resource Announcements, 2021, 10, .	0.6	0
14	Loss of antibodies to hepatitis E virus in organ transplant patients with hepatitis E. Hepatology Research, 2021, 51, 538-547.	3.4	2
15	Spontaneous reactivation of hepatitis B virus with a frameshift mutation in the precore region in an elderly hepatitis B virus carrier with lifestyle-related diseases. Clinical Journal of Gastroenterology, 2021, 14, 1202-1210.	0.8	1
16	Mechanism of Cross-Species Transmission, Adaptive Evolution and Pathogenesis of Hepatitis E Virus. Viruses, 2021, 13, 909.	3.3	23
17	Three cases of hepatitis E infected with subgenotype 3e hepatitis E virus indigenous to Mie Prefecture, which reemerged for the first time in six years. Acta Hepatologica Japonica, 2021, 62, 384-386.	0.1	1
18	An Autopsy Case of Primary Biliary Cholangitis with Histological Submassive Hepatic Necrosis Caused by Acute Hepatitis E Virus Infection. Internal Medicine, 2021, 60, 1863-1870.	0.7	1

#	Article	IF	CITATIONS
19	Knockdown of Mitogen-Activated Protein Kinase Kinase 3 Negatively Regulates Hepatitis A Virus Replication. International Journal of Molecular Sciences, 2021, 22, 7420.	4.1	8
20	Identification and a full genome analysis of novel camel hepatitis E virus strains obtained from Bactrian camels in Mongolia. Virus Research, 2021, 299, 198355.	2.2	14
21	Identification of a novel pegivirus in pet cats (Felis silvestris catus) in Japan. Virus Research, 2021, 301, 198452.	2.2	3
22	Taxonomic updates for the genus Gyrovirus (family Anelloviridae): recognition of several new members and establishment of species demarcation criteria. Archives of Virology, 2021, 166, 2937-2942.	2.1	18
23	Taxonomic update for mammalian anelloviruses (family Anelloviridae). Archives of Virology, 2021, 166, 2943-2953.	2.1	55
24	Production of capsid proteins of rat hepatitis E virus in Escherichia coli and characterization of self-assembled virus-like particles. Virus Research, 2021, 302, 198483.	2.2	4
25	Japanese Man with HCV Genotype 4 Infection and Cirrhosis Who was Successfully Treated by the Combination of Glecaprevir and Pibrentasvir. Internal Medicine, 2021, 60, 2061-2066.	0.7	1
26	Successful treatment of Japanese hemophilia patient co-infected with HIV and HCV genotype 4a by glecaprevir/pibrentasvir therapy. Clinical Journal of Gastroenterology, 2021, 14, 1725-1732.	0.8	2
27	The spontaneous clearance of hepatitis E virus (HEV) and emergence of HEV antibodies in a transfusion-transmitted chronic hepatitis E case after completion of chemotherapy for acute myeloid leukemia. Clinical Journal of Gastroenterology, 2020, 13, 252-259.	0.8	10
28	IFN-λ3 as a host immune response in acute hepatitis E virus infection. Cytokine, 2020, 125, 154816.	3.2	14
29	Molecular signature of hepatitisÂB virus regulation by interferonâ€Î³ in primary human hepatocytes. Hepatology Research, 2020, 50, 292-302.	3.4	3
30	A Nationwide Survey of Hepatitis E Virus Infection and Chronic Hepatitis in Heart and Kidney Transplant Recipients in Japan. Transplantation, 2020, 104, 437-444.	1.0	25
31	Hepatitis B virus reactivation after successful treatment of hepatitis C virus with sofosbuvir and ribavirin. Medicine (United States), 2020, 99, e22650.	1.0	4
32	Prevalence and genotype/subtype distribution of hepatitis E virus (HEV) among wild boars in Japan: Identification of a genotype 5 HEV strain. Virus Research, 2020, 287, 198106.	2.2	19
33	Additive Effects of Zinc Chloride on the Suppression of Hepatitis A Virus Replication by Interferon in Human Hepatoma Huh7 Cells. In Vivo, 2020, 34, 3301-3308.	1.3	12
34	Clinical and virologic features of hepatitis E virus infection at a university hospital in Japan between 2000 and 2019. Journal of Medical Virology, 2020, 92, 3572-3583.	5.0	3
35	Treatment Rationale and Design of a Phase III Study of Afatinib or Chemotherapy in Patients with Non–small-cell Lung Cancer Harboring Sensitizing Uncommon Epidermal Growth Factor Receptor Mutations (ACHILLES/TORG1834). Clinical Lung Cancer, 2020, 21, e592-e596.	2.6	5
36	Multivesicular body sorting and the exosomal pathway are required for the release of rat hepatitis E virus from infected cells. Virus Research, 2020, 278, 197868.	2.2	16

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37	Update: proposed reference sequences for subtypes of hepatitis E virus (species Orthohepevirus A). Journal of General Virology, 2020, 101, 692-698.	2.9	221
38	A case of autochthonous hepatitis E infected with subgenotype 4a hepatitis E virus endemic in China without overseas travel. Acta Hepatologica Japonica, 2020, 61, 270-272.	0.1	0
39	A fatal case of exacerbated liver cirrhosis caused by acute hepatitis E virus infection: atypical dynamics of anti-hepatitis E virus antibody titers. Acta Hepatologica Japonica, 2020, 61, 326-334.	0.1	1
40	Characteristics of cases of hepatitis E in 2019 in Gunma prefecture: a small epidemic caused by the same subgenotype 3a strain. Acta Hepatologica Japonica, 2020, 61, 478-481.	0.1	1
41	Successful prolonged treatment of glecaprevir/pibrentasvir for chronic hepatitis C patient with treatment failure after 8-week therapy: a case report. Clinical Journal of Gastroenterology, 2019, 12, 592-597.	0.8	3
42	Antiviral activity of zinc sulfate against hepatitis A virus replication. Future Virology, 2019, 14, 399-406.	1.8	12
43	Antiviral candidates against the hepatitis E virus (HEV) and their combinations inhibit HEV growth in in vitro. Antiviral Research, 2019, 170, 104570.	4.1	15
44	Acute Liver Failure Caused by the Transmission of Hepatitis B Virus from the Spouse after 38 Years of Marriage. Internal Medicine, 2019, 58, 2963-2968.	0.7	2
45	Superinfection of hepatitis A virus in hepatocytes infected with hepatitis B virus. International Journal of Medical Sciences, 2019, 16, 1366-1370.	2.5	9
46	Sofosbuvir/Ribavirin therapy for patients experiencing failure of ombitasvir/paritaprevir/ritonavir + ribavirin therapy: Two cases report and review of literature. World Journal of Clinical Cases, 2019, 7, 1043-1052.	0.8	1
47	Genetic Variability and Evolution of Hepatitis E Virus. Viruses, 2019, 11, 456.	3.3	71
48	Correlation between Helicobacter pylori Infection and Gastric Atrophy Examined in the Sera of Mongolian People. Gastrointestinal Disorders, 2019, 1, 241-252.	0.8	3
49	Screening of novel drugs for inhibiting hepatitis E virus replication. Journal of Virological Methods, 2019, 270, 1-11.	2.1	30
50	Changing clinical and molecular characteristics of hepatitis E virus infection in Mie Prefecture, Japan: Disappearance of indigenous subtype 3e strains. Hepatology Research, 2019, 49, 1003-1014.	3.4	4
51	Very low prevalence of anti-HAV in Japan: high potential for future outbreak. Scientific Reports, 2019, 9, 1493.	3.3	17
52	Dynamics of hepatitis B virus serum markers in an acute hepatitis B patient in the incubation phase. Clinical Journal of Gastroenterology, 2019, 12, 218-222.	0.8	0
53	Fullâ€genome characterization of the RIVMâ€HAV16â€090â€like hepatitis A virus strains recovered from Japanese men who have sex with men, with sporadic acute hepatitis A. Hepatology Research, 2019, 49, 521-530.	3.4	11
54	Prevalence and characteristics of hepatitis E virus infection in kidney transplant recipients: A singleâ€center experience in Japan. Transplant Infectious Disease, 2019, 21, e13033.	1.7	3

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55	An analysis of two open reading frames (ORF3 and ORF4) of rat hepatitis E virus genome using its infectious cDNA clones with mutations in ORF3 or ORF4. Virus Research, 2018, 249, 16-30.	2.2	16
56	The identification and characterization of novel rat hepatitis E virus strains in Bali and Sumbawa, Indonesia. Archives of Virology, 2018, 163, 1345-1349.	2.1	11
57	Clinical significance of changes in Torque teno virus DNA titer after chemotherapy in patients with primary lung cancer. Respiratory Investigation, 2018, 56, 173-178.	1.8	12
58	Hepatitis E virus subtype 3f strains isolated from Japanese hepatitis patients with no history of travel to endemic areas – The origin analyzed by molecular evolution. Virology, 2018, 513, 146-152.	2.4	12
59	A Case Report of Living Donor Liver Transplantation for Fulminant Hepatitis Related to Hepatitis E Virus Infection. Progress in Transplantation, 2018, 28, 91-92.	0.7	Ο
60	Distinct changing profiles of hepatitis A and E virus infection among patients with acute hepatitis in Mongolia: The first report of the full genome sequence of a novel genotype 1 hepatitis E virus strain. Journal of Medical Virology, 2018, 90, 84-92.	5.0	6
61	A 79-year-old Woman with Acute Hepatitis B Caused by the Infection of Subgenotype D1 Hepatitis B Virus in Japan: A Case Study. Internal Medicine, 2018, 57, 3099-3104.	0.7	Ο
62	A case of advanced gastric cancer with severe thrombocytopenia after administration of Nivolumab, following acute hepatitis E with decreased platelet count. Acta Hepatologica Japonica, 2018, 59, 497-500.	0.1	3
63	Identification and full-genome characterization of novel circoviruses in masked palm civets (Paguma) Tj ETQq1	1 0. <u>78</u> 4314	4 rgBT /Overlo
64	Identification and whole genome characterization of novel anelloviruses in masked palm civets (Paguma larvata): Segregation into four distinct clades. Virus Research, 2018, 256, 183-191.	2.2	8
65	Hepatitis E during Tocilizumab Therapy in a Patient with Rheumatoid Arthritis: Case Report and Literature Review. Case Reports in Rheumatology, 2018, 2018, 1-5.	0.6	8
66	Inhibitory effect of Japanese rice-koji miso extracts on hepatitis A virus replication in association with the elevation of glucose-regulated protein 78 expression. International Journal of Medical Sciences, 2018, 15, 1153-1159.	2.5	19
67	2. Hepatitis E: Frontier of Medical Practice. The Journal of the Japanese Society of Internal Medicine, 2018, 107, 1826-1832.	0.0	2
68	Three pairs of six sporadic acute hepatitis E cases infected with three different genotype 3b hepatitis E virus strains occurred within the same economic bloc in the central region of Japan. Acta Hepatologica Japonica, 2018, 59, 700-703.	0.1	0
69	Clinical and virological features of acute hepatitis E in Gunma prefecture, Japan between 2004 and 2015. Hepatology Research, 2017, 47, 435-445.	3.4	16
70	Unique clinical courses of transfusionâ€ŧransmitted hepatitis E in patients with immunosuppression. Transfusion, 2017, 57, 280-288.	1.6	62
71	Autochthonous sporadic acute hepatitis E caused by two distinct subgenotype 3b hepatitis E virus strains with only 90% nucleotide identity. Clinical Journal of Gastroenterology, 2017, 10, 168-173.	0.8	4
72	Glucose-regulated protein 78 is an antiviral against hepatitis A virus replication. Experimental and Therapeutic Medicine, 2017, 13, 3305-3308.	1.8	20

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73	Asymptomatic acute hepatitis E in a female patient with ulcerative colitis. Clinical Journal of Gastroenterology, 2017, 10, 255-260.	0.8	8
74	Fullâ€length genomic sequence analysis of new subtype 3k hepatitis E virus isolates with 99.97% nucleotide identity obtained from two consecutive acute hepatitis patients in a city in northeast Japan. Journal of Medical Virology, 2017, 89, 1116-1120.	5.0	20
75	Free fatty acids or high-concentration glucose enhances hepatitis A virus replication in association with a reduction in glucose-regulated protein 78 expression. Biochemical and Biophysical Research Communications, 2017, 483, 694-699.	2.1	12
76	Full-length genomic sequences of new subtype 1g hepatitis E virus strains obtained from four patients with imported or autochthonous acute hepatitis E in Japan. Infection, Genetics and Evolution, 2017, 55, 343-349.	2.3	7
77	Full-length genome of a novel genotype 3 hepatitis E virus strain obtained from domestic pigs in Japan. Virus Research, 2017, 240, 147-153.	2.2	3
78	Fatal acute hepatic failure in a family infected with the hepatitis A virus subgenotype IB. Medicine (United States), 2017, 96, e7847.	1.0	11
79	Characterization of the Quasi-Enveloped Hepatitis E Virus Particles Released by the Cellular Exosomal Pathway. Journal of Virology, 2017, 91, .	3.4	151
80	Occult hepatitis B virus infection in immunized children born to carrier mothers. Pediatrics International, 2017, 59, 1010-1016.	0.5	10
81	Hepatitis A virus genotype IA-infected patient with marked elevation of aspartate aminotransferase levels. Clinical Journal of Gastroenterology, 2017, 10, 52-56.	0.8	5
82	An autochthonous case of acute hepatitis E in Mie, Japan who was infected with a rare hepatitis E virus strain of subgenotype 3f. Acta Hepatologica Japonica, 2017, 58, 53-59.	0.1	2
83	Combination Therapy with Ombitasvir/Paritaprevir/Ritonavir for Dialysis Patients Infected with Hepatitis C Virus: A Prospective Multi-Institutional Study. Tohoku Journal of Experimental Medicine, 2017, 241, 45-53.	1.2	12
84	Ruptured dissecting aneurysm of the recurrent artery of Heubner: Consideration of pathological findings. Neurology India, 2017, 65, 623.	0.4	5
85	Molecular analysis of hepatitis A virus strains obtained from patients with acute hepatitis A in Mongolia, 2004–2013. Journal of Medical Virology, 2016, 88, 622-630.	5.0	4
86	Treatment with ribavirin for chronic hepatitis E following living donor liver transplantation: A case report. Hepatology Research, 2016, 46, 1058-1059.	3.4	10
87	Prevalence of hepatitis viruses in patients with acute hepatitis and characterization of the detected genotype 4 hepatitis E virus sequences in Mongolia. Journal of Medical Virology, 2016, 88, 282-291.	5.0	4
88	Chronic hepatitis E virus infection after living donor liver transplantation via blood transfusion: a case report. Surgical Case Reports, 2016, 2, 32.	0.6	21
89	Analysis of adaptive mutations selected during the consecutive passages of hepatitis E virus produced from an infectious cDNA clone. Virus Research, 2016, 223, 170-180.	2.2	9
90	Production of monoclonal antibodies against the ORF3 protein of rat hepatitis E virus (HEV) and demonstration of the incorporation of the ORF3 protein into enveloped rat HEV particles. Archives of Virology, 2016, 161, 3391-3404.	2.1	7

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91	Comparison of test results of serogrouping and core region PCR-based genotyping in patients with chronic hepatitis C virus infection: Analysis of inderminate or discrepant cases and identification of a 2b/1b recombinant HCV. Acta Hepatologica Japonica, 2016, 57, 447-456.	0.1	7
92	Two cases of sporadic acute hepatitis E in Mie, Japan who were infected with subgenotype 1a hepatitis E virus. Acta Hepatologica Japonica, 2016, 57, 81-88.	0.1	3
93	Fatal fulminant hepatitis caused by infection with subgenotype A1 hepatitis B virus with C1766T/T1768A core promoter mutations. Clinical Journal of Gastroenterology, 2016, 9, 160-167.	0.8	3
94	Possible involvement of PCSK9 overproduction in hyperlipoproteinemia associated with hepatocellular carcinoma: A case report. Journal of Clinical Lipidology, 2016, 10, 1045-1049.	1.5	14
95	Investigating the origin and global dispersal history of hepatitis E virus genotype 4 using phylogeographical analysis. Liver International, 2016, 36, 31-41.	3.9	18
96	Characterization and epitope mapping of monoclonal antibodies raised against rat hepatitis E virus capsid protein: An evaluation of their neutralizing activity in a cell culture system. Journal of Virological Methods, 2016, 233, 78-88.	2.1	17
97	Proposed reference sequences for hepatitis E virus subtypes. Journal of General Virology, 2016, 97, 537-542.	2.9	339
98	Enhanced pregenomic RNA levels and lowered precore mRNA transcription efficiency in a genotype A hepatitis B virus genome with C1766T and T1768A mutations obtained from a fulminant hepatitis patient. Journal of General Virology, 2016, 97, 2643-2656.	2.9	7
99	A case of acute hepatitis E who was suspected to have contracted hepatitis E virus infection via consumption of dried gall bladders from wild boars. Acta Hepatologica Japonica, 2016, 57, 606-613.	0.1	3
100	Clinical Features of Hepatitis E Virus Infection in Ibaraki, Japan: Autochthonous Hepatitis E and Acute-on-Chronic Liver Failure. Tohoku Journal of Experimental Medicine, 2015, 235, 275-282.	1.2	18
101	Serum Torque Teno Virus DNA Titer in Idiopathic Pulmonary Fibrosis Patients with Acute Respiratory Worsening. Internal Medicine, 2015, 54, 1015-1019.	0.7	11
102	Characteristics of 20 Patients with Autochthonous Acute Hepatitis E in Hokkaido, Japan: First Report of Bilateral Facial Palsy Following the Infection with Genotype 4 Hepatitis E Virus. Tohoku Journal of Experimental Medicine, 2015, 236, 263-271.	1.2	23
103	Recent Trend of Hepatitis E Virus Infection in Chiba Area, Japan: 3 of 5 Cases with Rheumatoid Arthritis. Case Reports in Gastroenterology, 2015, 9, 317-326.	0.6	8
104	Metachronous occurrence of two cases of acute hepatitis E after eating raw pig liver and heart at the same restaurant at Gifu city in Japan. Acta Hepatologica Japonica, 2015, 56, 617-620.	0.1	0
105	Full genome analysis of a European-type genotype 3 hepatitis E virus variant obtained from a Japanese patient with autochthonous acute hepatitis E. Journal of Medical Virology, 2015, 87, 1067-1071.	5.0	10
106	Horizontal transmission of de novo hepatitis <scp>B</scp> between spouses: A case report. Hepatology Research, 2015, 45, 933-938.	3.4	2
107	Spontaneous reactivation of hepatitis B virus (HBV) infection in patients with resolved or occult HBV infection. Journal of Medical Virology, 2015, 87, 589-600.	5.0	31
108	An expanded taxonomy of hepatitis C virus genotype 6: Characterization of 22 new full-length viral genomes. Virology, 2015, 476, 355-363.	2.4	27

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109	Acute infection by hepatitis E virus with a slight immunoglobulin M antibody response. Clinical Journal of Gastroenterology, 2015, 8, 247-252.	0.8	3
110	The JAK2 inhibitor AZD1480 inhibits hepatitis A virus replication inÂHuh7 cells. Biochemical and Biophysical Research Communications, 2015, 458, 908-912.	2.1	30
111	The sirtuin inhibitor sirtinol inhibits hepatitis A virus (HAV) replication by inhibiting HAV internal ribosomal entry site activity. Biochemical and Biophysical Research Communications, 2015, 466, 567-571.	2.1	25
112	A Nationwide Survey of Hepatitis E Virus Infection and Chronic Hepatitis E in Liver Transplant Recipients in Japan. EBioMedicine, 2015, 2, 1607-1612.	6.1	48
113	Clinical and molecular analyses of sporadic acute hepatitis <scp>A</scp> and <scp>E</scp> and the specific viral genotypes isolated in <scp>I</scp> wate and three neighboring prefectures in the northern part of <scp>H</scp> onshu, <scp>J</scp> apan, between 2004 and 2013. Hepatology Research, 2015. 45, 714-727.	3.4	5
114	Screening of hepatitis E virus infection should be included in the current diagnostic scale for drug-induced liver injury in Japan. Acta Hepatologica Japonica, 2014, 55, 325-334.	0.1	1
115	Changing trends of genotypic distribution of hepatitis B virus in Mito, Japan (2001-2013). Acta Hepatologica Japonica, 2014, 55, 626-629.	0.1	1
116	Full genome analysis of <scp>P</scp> hilippine indigenous subgenotype <scp>IA</scp> hepatitis <scp>A</scp> virus strains from <scp>J</scp> apanese patients with imported acute hepatitis <scp>A</scp> . Hepatology Research, 2014, 44, 270-279.	3.4	9
117	Consensus proposals for classification of the family Hepeviridae. Journal of General Virology, 2014, 95, 2223-2232.	2.9	570
118	High genomic similarity between <scp>E</scp> uropean type hepatitis <scp>E</scp> virus subgenotype 3e strains isolated from an acute hepatitis patient and a wild boar in <scp>M</scp> ie, <scp>J</scp> apan. Hepatology Research, 2014, 44, 694-699.	3.4	15
119	Characterization of sporadic acute hepatitis <scp>E</scp> and comparison of hepatitis <scp>E</scp> virus genomes in acute hepatitis patients and pig liver sold as food in <scp>M</scp> ie, <scp>J</scp> apan. Hepatology Research, 2014, 44, E63-E76.	3.4	30
120	The complete genomes of subgenotype IA hepatitis A virus strains from four different islands in Indonesia form a phylogenetic cluster. Archives of Virology, 2014, 159, 935-945.	2.1	10
121	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.	2.2	62
122	Origin of hepatitis C virus genotype 3 in Africa as estimated through an evolutionary analysis of the full-length genomes of nine subtypes, including the newly sequenced 3d and 3e. Journal of General Virology, 2014, 95, 1677-1688.	2.9	34
123	Marked genomic heterogeneity of rat hepatitis E virus strains in Indonesia demonstrated on a full-length genome analysis. Virus Research, 2014, 179, 102-112.	2.2	43
124	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.	2.1	69
125	Hepatitis E virus egress depends on the exosomal pathway, with secretory exosomes derived from multivesicular bodies. Journal of General Virology, 2014, 95, 2166-2175.	2.9	163
126	Molecular analysis of the interspousal transmission of hepatitis B virus in two Japanese patients who acquired fulminant hepatitis B after 50 and 49 years of marriage. Journal of Medical Virology, 2014, 86, 1851-1860.	5.0	5

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#	Article	IF	CITATIONS
127	Rat hepatitis E virus derived from wild rats (Rattus rattus) propagates efficiently in human hepatoma cell lines. Virus Research, 2014, 185, 92-102.	2.2	43
128	Features of hepatitis <scp>E</scp> virus infection in humans and animals in <scp>J</scp> apan. Hepatology Research, 2014, 44, 43-58.	3.4	91
129	Two Cases of Subfrontal Schwannoma, Including a Rare Case Located between the Endosteal and Meningeal Layers of the Dura. Neurologia Medico-Chirurgica, 2014, 54, 681-685.	2.2	5
130	Suppression of La Antigen Exerts Potential Antiviral Effects against Hepatitis A Virus. PLoS ONE, 2014, 9, e101993.	2.5	21
131	Occurrence of acute hepatitis E virus infection in the Hakodate district: A prospective study of four hospitals in Hakodate City. Acta Hepatologica Japonica, 2014, 55, 349-359.	0.1	1
132	A case of acute hepatitis E in Mie prefecture infected with genotype 4 hepatitis E virus strain endemic in Aichi and Shizuoka prefectures (Aichi/Shizuoka strain), without a history of eating wild animal meat. Acta Hepatologica Japonica, 2014, 55, 405-408.	0.1	5
133	European-type subgenotype 3e/3sp hepatitis E virus strain identified in a pig as a possible origin of hepatitis E cases in Mie prefecture during 2004-2012. Acta Hepatologica Japonica, 2014, 55, 553-555.	0.1	4
134	Frequent detection and characterization of hepatitis E virus variants in wild rats (Rattus rattus) in Indonesia. Archives of Virology, 2013, 158, 87-96.	2.1	43
135	Culture systems for hepatitis E virus. Journal of Gastroenterology, 2013, 48, 147-158.	5.1	83
136	Identification of European-type hepatitis E virus subtype 3e isolates in Japanese wild boars: Molecular tracing of HEV from swine to wild boars. Infection, Genetics and Evolution, 2013, 18, 287-298.	2.3	25
137	Full-length genome sequences of five hepatitis C virus isolates representing subtypes 3g, 3h, 3i and 3k, and a unique genotype 3 variant. Journal of General Virology, 2013, 94, 543-548.	2.9	23
138	Development of a Highly Sensitive Bioluminescent Enzyme Immunoassay for Hepatitis B Virus Surface Antigen Capable of Detecting Divergent Mutants. Vaccine Journal, 2013, 20, 1255-1265.	3.1	8
139	Transmission of genotype A (A2) hepatitis B virus from patients who developed acute hepatitis B to their sexual partners, via interspousal or homosexual infection. Acta Hepatologica Japonica, 2013, 54, 373-380.	0.1	1
140	A Marked Decrease in CD4-positive Lymphocytes at the Onset of Hepatitis in a Patient With Hepatitis-associated Aplastic Anemia. Journal of Pediatric Hematology/Oncology, 2012, 34, 375-377.	0.6	18
141	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.	2.2	43
142	Identification of four novel subgenotypes (C13–C16) and two inter-genotypic recombinants (C12/G and) Tj ET	Qq0,0 0 rį 2.2	gBT /Overlock
143	New findings regarding the epidemic history and population dynamics of Japanâ€indigenous genotype 3 hepatitis E virus inferred by molecular evolution. Liver International, 2012, 32, 675-688.	3.9	23

144Icteric acute hepatitis E with no response of immunoglobulin M class antiâ€hepatitis E virus antibody.3.46Hepatology Research, 2012, 42, 1146-1149.6

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#	Article	IF	CITATIONS
145	Molecular epidemiology and genetic history of European-type genotype 3 hepatitis E virus indigenized in the central region of Japan. Infection, Genetics and Evolution, 2012, 12, 1524-1534.	2.3	21
146	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.	2.1	57
147	Two cases with imported hepatitis A who were presumed to have contracted the disease while visiting Indonesia based on molecular analysis of hepatitis A virus strains. Acta Hepatologica Japonica, 2012, 53, 754-762.	0.1	1
148	Small epidemic of hepatitis E in the fall 2009 in Sapporo, Hokkaido. Acta Hepatologica Japonica, 2012, 53, 78-89.	0.1	6
149	Hepatitis E virus cell culture models. Virus Research, 2011, 161, 65-77.	2.2	80
150	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.	2.1	103
151	Identification and characterization of a natural inter-genotypic (2b/1b) recombinant hepatitis C virus in Japan. Archives of Virology, 2011, 156, 1591-1601.	2.1	23
152	Efficient cell culture systems for hepatitis E virus strains in feces and circulating blood. Reviews in Medical Virology, 2011, 21, 18-31.	8.3	42
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HLA DRB1 and DQB1 alleles and haplotypes influencing the progression of hepatitis C. , 1996, 49, 274-278. 360

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