## Jan Vinje

## List of Publications by Year in descending order

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16791 14012 19,419 211 66 133 citations h-index g-index papers 219 219 219 9290 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Secretor Status Strongly Influences the Incidence of Symptomatic Norovirus Infection in a Genotype-Dependent Manner in a Nicaraguan Birth Cohort. Journal of Infectious Diseases, 2022, 225, 105-115.	1.9	18
2	Norovirus Outbreaks in Long-term Care Facilities in the United States, 2009–2018: A Decade of Surveillance. Clinical Infectious Diseases, 2022, 74, 113-119.	2.9	17
3	Development and evaluation of a ligation-free sequence-independent, single-primer amplification (LF-SISPA) assay for whole genome characterization of viruses. Journal of Virological Methods, 2022, 299, 114346.	1.0	1
4	Descriptive evaluation of antibody responses to severe acute respiratory coronavirus virus 2 (SARS-CoV-2) infection in plasma and gingival crevicular fluid in a nursing home cohort—Arkansas, June–August 2020. Infection Control and Hospital Epidemiology, 2022, 43, 1610-1617.	1.0	3
5	Advances in understanding of the innate immune response to human norovirus infection using organoid models. Journal of General Virology, 2022, 103, .	1.3	14
6	Development and Validation of an Enzyme Immunoassay for Detection and Quantification of SARS-CoV-2 Salivary IgA and IgG. Journal of Immunology, 2022, 208, 1500-1508.	0.4	19
7	Hygienic monitoring in long-term care facilities using ATP, crAssphage, and human noroviruses to direct environmental surface cleaning. American Journal of Infection Control, 2022, 50, 289-294.	1.1	4
8	Efficacy of EPA-registered disinfectants against two human norovirus surrogates and Clostridioides difficile endospores. Journal of Applied Microbiology, 2022, 132, 4289-4299.	1.4	2
9	Gut Microbiome Changes Occurring with Norovirus Infection and Recovery in Infants Enrolled in a Longitudinal Birth Cohort in Leon, Nicaragua. Viruses, 2022, 14, 1395.	1.5	3
10	Characteristics of GII.4 Norovirus Versus Other Genotypes in Sporadic Pediatric Infections in Davidson County, Tennessee, USA. Clinical Infectious Diseases, 2021, 73, e1525-e1531.	2.9	24
11	Single-step RT-PCR assay for dual genotyping of GI and GII norovirus strains. Journal of Clinical Virology, 2021, 134, 104689.	1.6	34
12	Human Calicivirus Typing tool: A web-based tool for genotyping human norovirus and sapovirus sequences. Journal of Clinical Virology, 2021, 134, 104718.	1.6	29
13	Detection of Norovirus Variant GII.4 Hong Kong in Asia and Europe, 2017â^2019. Emerging Infectious Diseases, 2021, 27, 289-293.	2.0	21
14	Homotypic and Heterotypic Protection and Risk of Reinfection Following Natural Norovirus Infection in a Highly Endemic Setting. Clinical Infectious Diseases, 2021, 72, 222-229.	2.9	25
15	Risk Factors and Clinical Profile of Sapovirus-associated Acute Gastroenteritis in Early Childhood. Pediatric Infectious Disease Journal, 2021, 40, 220-226.	1.1	18
16	Pediatric Respiratory and Enteric Virus Acquisition and Immunogenesis in US Mothers and Children Aged 0-2: PREVAIL Cohort Study. JMIR Research Protocols, 2021, 10, e22222.	0.5	11
17	Rare Norovirus GIV Foodborne Outbreak, Wisconsin, USA. Emerging Infectious Diseases, 2021, 27, 1151-1154.	2.0	8
18	Global Trends in Norovirus Genotype Distribution among Children with Acute Gastroenteritis. Emerging Infectious Diseases, 2021, 27, 1438-1445.	2.0	85

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19	Detection of SARS-CoV-2 on Surfaces in Households of Persons with COVID-19. International Journal of Environmental Research and Public Health, 2021, 18, 8184.	1.2	37
20	Global distribution of sporadic sapovirus infections: A systematic review and meta-analysis. PLoS ONE, 2021, 16, e0255436.	1.1	10
21	Viral etiology of acute gastroenteritis among Forcibly Displaced Myanmar Nationals and adjacent host population in Bangladesh. Journal of Infectious Diseases, 2021, , .	1.9	2
22	Rotaviruses, astroviruses, and sapoviruses as foodborne infections., 2021,, 327-344.		2
23	The Changing Landscape of Pediatric Viral Enteropathogens in the Post–Rotavirus Vaccine Era. Clinical Infectious Diseases, 2021, 72, 576-585.	2.9	26
24	Hospital-based Surveillance for Pediatric Norovirus Gastroenteritis in Bangladesh, 2012–2016. Pediatric Infectious Disease Journal, 2021, 40, 215-219.	1.1	4
25	Prevalence and genetic characterization of noroviruses in children with acute gastroenteritis in Senegal, 2007â€2010. Journal of Medical Virology, 2021, , .	2.5	0
26	Gastrointestinal Tract Infections: Viruses. , 2021, , .		0
27	Trends in Incidence of Norovirus-associated Acute Gastroenteritis in 4 Veterans Affairs Medical Center Populations in the United States, 2011–2015. Clinical Infectious Diseases, 2020, 70, 40-48.	2.9	11
28	Infectious Causes of Acute Gastroenteritis in US Children Undergoing Allogeneic Hematopoietic Cell Transplant: A Longitudinal, Multicenter Study. Journal of the Pediatric Infectious Diseases Society, 2020, 9, 421-427.	0.6	5
29	Temporal and Genotypic Associations of Sporadic Norovirus Gastroenteritis and Reported Norovirus Outbreaks in Middle Tennessee, 2012–2016. Clinical Infectious Diseases, 2020, 71, 2398-2404.	2.9	8
30	Preadaptation of pandemic GII.4Ânoroviruses in unsampled virus reservoirs years before emergence. Virus Evolution, 2020, 6, veaa067.	2.2	22
31	Virus–Host Interactions Between Nonsecretors and Human Norovirus. Cellular and Molecular Gastroenterology and Hepatology, 2020, 10, 245-267.	2.3	24
32	Human Intestinal Enteroids to Evaluate Human Norovirus GII.4 Inactivation by Aged-Green Tea. Frontiers in Microbiology, 2020, 11, 1917.	1.5	29
33	CrAssphage as a Novel Tool to Detect Human Fecal Contamination on Environmental Surfaces and Hands. Emerging Infectious Diseases, 2020, 26, 1731-1739.	2.0	34
34	Recent advances in human norovirus research and implications for candidate vaccines. Expert Review of Vaccines, 2020, 19, 539-548.	2.0	46
35	Incidence, etiology, and severity of acute gastroenteritis among prospectively enrolled patients in 4 Veterans Affairs hospitals and outpatient centers, 2016–18. Clinical Infectious Diseases, 2020, 73, e2729-e2738.	2.9	16
36	Norovirus Seroprevalence among Adults in the United States: Analysis of NHANES Serum Specimens from 1999–2000 and 2003–2004. Viruses, 2020, 12, 179.	1.5	5

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37	Norovirus Outbreak Surveillance, China, 2016–2018. Emerging Infectious Diseases, 2020, 26, 437-445.	2.0	53
38	Molecular epidemiology of norovirus outbreaks in Argentina, 2013â€2018. Journal of Medical Virology, 2020, 92, 1330-1333.	2.5	14
39	Humoral and Mucosal Immune Responses to Human Norovirus in the Elderly. Journal of Infectious Diseases, 2020, 221, 1864-1874.	1.9	14
40	Comparison of Illumina MiSeq and the Ion Torrent PGM and S5 platforms for whole-genome sequencing of picornaviruses and caliciviruses. Journal of Virological Methods, 2020, 280, 113865.	1.0	20
41	Evaluation of viral co-infections among patients with community-associated Clostridioides difficile infection. PLoS ONE, 2020, 15, e0240549.	1.1	2
42	<i>Notes from the Field:</i> Multiple Cruise Ship Outbreaks of Norovirus Associated with Frozen Fruits and Berries — United States, 2019. Morbidity and Mortality Weekly Report, 2020, 69, 501-502.	9.0	14
43	Incidence and etiology of infectious diarrhea from a facility-based surveillance system in Guatemala, 2008–2012. BMC Public Health, 2019, 19, 1340.	1,2	8
44	A new solid matrix for preservation of viral nucleic acid from clinical specimens at ambient temperature. Journal of Virological Methods, 2019, 274, 113732.	1.0	6
45	Sera Antibody Repertoire Analyses Reveal Mechanisms of Broad and Pandemic Strain Neutralizing Responses after Human Norovirus Vaccination. Immunity, 2019, 50, 1530-1541.e8.	6.6	71
46	Emerging Novel GII.P16 Noroviruses Associated with Multiple Capsid Genotypes. Viruses, 2019, 11, 535.	1.5	53
47	Prevalence and genetic diversity of viral gastroenteritis viruses in children younger than 5 years of age in Guatemala, 2014–2015. Journal of Clinical Virology, 2019, 114, 6-11.	1.6	16
48	Complete Genome Sequences of Human Astrovirus Prototype Strains (Types $1\ \rm to\ 8$ ). Microbiology Resource Announcements, 2019, $8, .$	0.3	3
49	Near-Complete Human Sapovirus Genome Sequences from Kenya. Microbiology Resource Announcements, 2019, 8, .	0.3	16
50	Active Surveillance for Norovirus in a US Veterans Affairs Patient Population, Houston, Texas, 2015–2016. Open Forum Infectious Diseases, 2019, 6, ofz115.	0.4	6
51	Impact of long-term storage of clinical samples collected from 1996 to 2017 on RT-PCR detection of norovirus. Journal of Virological Methods, 2019, 267, 35-41.	1.0	10
52	Birth Cohort Studies Assessing Norovirus Infection and Immunity in Young Children: A Review. Clinical Infectious Diseases, 2019, 69, 357-365.	2.9	43
53	Sapovirus: an important cause of acute gastroenteritis in children. The Lancet Child and Adolescent Health, 2019, 3, 758-759.	2.7	27
54	Viral Etiology of Acute Gastroenteritis in <2-Year-Old US Children in the Post–Rotavirus Vaccine Era. Journal of the Pediatric Infectious Diseases Society, 2019, 8, 414-421.	0.6	53

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55	Molecular epidemiology of noroviruses in children under 5 years of age with acute gastroenteritis in Yaoundé, Cameroon. Journal of Medical Virology, 2019, 91, 738-743.	2.5	8
56	The Norovirus Epidemiologic Triad: Predictors of Severe Outcomes in US Norovirus Outbreaks, 2009–2016. Journal of Infectious Diseases, 2019, 219, 1364-1372.	1.9	52
57	Epidemiologic and Genotypic Distribution of Noroviruses Among Children With Acute Diarrhea and Healthy Controls in a Low-income Rural Setting. Clinical Infectious Diseases, 2019, 69, 505-513.	2.9	17
58	Minimally Invasive Saliva Testing to Monitor Norovirus Infection in Community Settings. Journal of Infectious Diseases, 2019, 219, 1234-1242.	1.9	22
59	Updated classification of norovirus genogroups and genotypes. Journal of General Virology, 2019, 100, 1393-1406.	1.3	535
60	ICTV Virus Taxonomy Profile: Caliciviridae. Journal of General Virology, 2019, 100, 1469-1470.	1.3	117
61	Enteropathogen antibody dynamics and force of infection among children in low-resource settings. ELife, 2019, 8, .	2.8	26
62	High Hand Contamination Rates During Norovirus Outbreaks in Long-Term Care Facilities. Infection Control and Hospital Epidemiology, 2018, 39, 219-221.	1.0	6
63	Antigenic Characterization of a Novel Recombinant GII.P16-GII.4 Sydney Norovirus Strain With Minor Sequence Variation Leading to Antibody Escape. Journal of Infectious Diseases, 2018, 217, 1145-1152.	1.9	30
64	1648. Incidence of Norovirus and Rotavirus From Multisite Active Surveillance in Veteran's Affairs Hospitals, December 2016–February 2018: Results From the SUPERNOVA Network. Open Forum Infectious Diseases, 2018, 5, S49-S49.	0.4	3
65	Water quality, availability, and acute gastroenteritis on the Navajo Nation – a pilot case-control study. Journal of Water and Health, 2018, 16, 1018-1028.	1.1	4
66	Near-Complete Genome Sequences of Several New Norovirus Genogroup II Genotypes. Genome Announcements, 2018, 6, .	0.8	19
67	Prevalence of Human Noroviruses in Commercial Food Establishment Bathrooms. Journal of Food Protection, 2018, 81, 719-728.	0.8	12
68	Genetic Diversity of Noroviruses Circulating in a Pediatric Cohort in Bangladesh. Journal of Infectious Diseases, 2018, 218, 1937-1942.	1.9	13
69	Epidemiology of Foodborne Norovirus Outbreaks – United States, 2009–2015. Food Safety (Tokyo,) Tj ETQq1	1.0.7843	14 rgBT /
70	Human Norovirus Replication in Human Intestinal Enteroids as Model to Evaluate Virus Inactivation. Emerging Infectious Diseases, 2018, 24, 1453-1464.	2.0	179
71	Genetic diversity of human sapovirus across the Americas. Journal of Clinical Virology, 2018, 104, 65-72.	1.6	45
72	Synthesis and Evaluation of Biotinylated Bivalent HistoBlood Group Antigens for Capturing Human Noroviruses. FASEB Journal, 2018, 32, 544.22.	0.2	0

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73	Complete Genome Sequence of Human Norovirus GII.Pe-GII.4 Sydney from the United States. Genome Announcements, $2017, 5, \ldots$	0.8	2
74	Genetic and Epidemiologic Trends of Norovirus Outbreaks in the United States from 2013 to 2016 Demonstrated Emergence of Novel GII.4 Recombinant Viruses. Journal of Clinical Microbiology, 2017, 55, 2208-2221.	1.8	222
75	Pathogen-Specific Burden of Outpatient Diarrhea in Infants in Nepal: A Multisite Prospective Case-Control Study. Journal of the Pediatric Infectious Diseases Society, 2017, 6, e75-e85.	0.6	10
76	Genomic Characterization of Three Melon Necrotic Spot Viruses Detected in Human Stool Specimens. Genome Announcements, $2017, 5, .$	0.8	3
77	Swab Sampling Method for the Detection of Human Norovirus on Surfaces. Journal of Visualized Experiments, 2017, , .	0.2	27
78	Pediatric norovirus GII.4 infections in Nicaragua, 1999–2015. Infection, Genetics and Evolution, 2017, 55, 305-312.	1.0	26
79	Comparison of three multiplex gastrointestinal platforms for the detection of gastroenteritis viruses. Journal of Clinical Virology, 2017, 95, 66-71.	1.6	41
80	Can Use of Viral Load Improve Norovirus Clinical Diagnosis and Disease Attribution?. Open Forum Infectious Diseases, 2017, 4, ofx131.	0.4	21
81	Association of GII.P16-GII.2 Recombinant Norovirus Strain with Increased Norovirus Outbreaks, Guangdong, China, 2016. Emerging Infectious Diseases, 2017, 23, 1188-1190.	2.0	50
82	Global Spread of Norovirus GII.17 Kawasaki 308, 2014–2016. Emerging Infectious Diseases, 2017, 23, 1359-1354.	2.0	71
83	Near Real-Time Surveillance of U.S. Norovirus Outbreaks by the Norovirus Sentinel Testing and Tracking Network — United States, August 2009–July 2015. Morbidity and Mortality Weekly Report, 2017, 66, 185-189.	9.0	26
84	Critical role of RIG-I and MDA5 in early and late stages of Tulane virus infection. Journal of General Virology, 2017, 98, 1016-1026.	1.3	11
85	Norovirus and Sapovirus Epidemiology and Strain Characteristics among Navajo and Apache Infants. PLoS ONE, 2017, 12, e0169491.	1.1	13
86	Detection of human norovirus in intestinal biopsies from immunocompromised transplant patients. Journal of General Virology, 2016, 97, 2291-2300.	1.3	85
87	Strain-Specific Virolysis Patterns of Human Noroviruses in Response to Alcohols. PLoS ONE, 2016, 11, e0157787.	1.1	14
88	Synthesis and Evaluation of Biotinylated Bivalent HistoBlood Group Antigens for Capturing Human Noroviruses. Bioconjugate Chemistry, 2016, 27, 1822-1829.	1.8	4
89	The effect of diarrheal disease on bivalent oral polio vaccine (bOPV) immune response in infants in Nepal. Vaccine, 2016, 34, 2519-2526.	1.7	11
90	Molecular Detection Methods of Foodborne Viruses. , 2016, , 303-333.		5

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91	Complete Genome Sequence of Human Norovirus Strain GII.P7-GII.6 Detected in a Patient in the United States in 2014. Genome Announcements, 2016, 4, .	0.8	12
92	Genetic characterization of norovirus strains in hospitalized children from Pakistan. Journal of Medical Virology, 2016, 88, 216-223.	2.5	20
93	Using Multiplex Molecular Testing to Determine the Etiology of AcuteÂGastroenteritis in Children. Journal of Pediatrics, 2016, 176, 50-56.e2.	0.9	52
94	Norovirus in a United States virgin islands resort: outbreak investigation, response, and costs. Journal of Travel Medicine, 2016, 23, taw040.	1.4	7
95	Characterization of a Salivirus ( $\mbox{Picornaviridae}) from a Diarrheal Child in Guatemala. Genome Announcements, 2016, 4, .$	0.8	4
96	Multicenter Evaluation of the Xpert Norovirus Assay for Detection of Norovirus Genogroups I and II in Fecal Specimens. Journal of Clinical Microbiology, 2016, 54, 142-147.	1.8	32
97	Epidemiologic, Virologic, and Host Genetic Factors of Norovirus Outbreaks in Long-term Care Facilities. Clinical Infectious Diseases, 2016, 62, 1-10.	2.9	196
98	Comparison of norovirus genogroup I, II and IV seroprevalence among children in the Netherlands, 1963, 1983 and 2006. Journal of General Virology, 2016, 97, 2255-2264.	1.3	26
99	Population-Based Incidence Rates of Diarrheal Disease Associated with Norovirus, Sapovirus, and Astrovirus in Kenya. PLoS ONE, 2016, 11, e0145943.	1.1	37
100	Transmission of Norovirus Within Households in Quininde, Ecuador. Pediatric Infectious Disease Journal, 2015, 34, 1031-1033.	1.1	16
101	Epidemiology and molecular characteristics of norovirus GII.4 Sydney outbreaks in Taiwan, January 2012-December 2013. Journal of Medical Virology, 2015, 87, 1462-1470.	2.5	17
102	Differences in Norovirus-Associated Hospital Visits Between Jewish and Bedouin Children in Southern Israel. Pediatric Infectious Disease Journal, 2015, 34, 1036-1038.	1.1	14
103	Development of a Nucleic Acid Extraction Procedure for Simultaneous Recovery of DNA and RNA from Diverse Microbes in Water. Pathogens, 2015, 4, 335-354.	1.2	36
104	A diverse group of small circular ssDNA viral genomes in human and non-human primate stools. Virus Evolution, 2015, 1, vev017.	2.2	49
105	Norovirus Vaccine Against Experimental Human GII.4 Virus Illness: A Challenge Study in Healthy Adults. Journal of Infectious Diseases, 2015, 211, 870-878.	1.9	223
106	Evaluation of a New Environmental Sampling Protocol for Detection of Human Norovirus on Inanimate Surfaces. Applied and Environmental Microbiology, 2015, 81, 5987-5992.	1.4	44
107	Serological Correlates of Protection against a GII.4 Norovirus. Vaccine Journal, 2015, 22, 923-929.	3.2	109
108	Noroviruses: epidemiology, immunity and prospects for prevention. Future Microbiology, 2015, 10, 53-67.	1.0	78

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109	Innate Susceptibility to Norovirus Infections Influenced by FUT2 Genotype in a United States Pediatric Population. Clinical Infectious Diseases, 2015, 60, 1631-1638.	2.9	98
110	Human norovirus culture in B cells. Nature Protocols, 2015, 10, 1939-1947.	5 <b>.</b> 5	202
111	Norovirus Genotype Profiles Associated with Foodborne Transmission, 1999–2012. Emerging Infectious Diseases, 2015, 21, 592-599.	2.0	136
112	Antimicrobial activity of bismuth subsalicylate on <i>Clostridium difficile</i> , <i>Escherichia coli</i> O157:H7, norovirus, and other common enteric pathogens. Gut Microbes, 2015, 6, 93-100.	4.3	41
113	Norovirus Infection and Disease in an Ecuadorian Birth Cohort: Association of Certain Norovirus Genotypes With Host FUT2 Secretor Status. Journal of Infectious Diseases, 2015, 211, 1813-1821.	1.9	106
114	Advances in Laboratory Methods for Detection and Typing of Norovirus. Journal of Clinical Microbiology, 2015, 53, 373-381.	1.8	639
115	Incidence of Medically-Attended Norovirus-Associated Acute Gastroenteritis in Four Veteran's Affairs Medical Center Populations in the United States, 2011-2012. PLoS ONE, 2015, 10, e0126733.	1.1	13
116	Genotypic and Epidemiologic Trends of Norovirus Outbreaks in the United States, 2009 to 2013. Journal of Clinical Microbiology, 2014, 52, 147-155.	1.8	265
117	Seroprevalence of Canine Norovirus in 14 European Countries. Vaccine Journal, 2014, 21, 898-900.	3.2	14
118	RNA Populations in Immunocompromised Patients as Reservoirs for Novel Norovirus Variants. Journal of Virology, 2014, 88, 14184-14196.	1.5	44
119	Comprehensive Comparison of Cultivable Norovirus Surrogates in Response to Different Inactivation and Disinfection Treatments. Applied and Environmental Microbiology, 2014, 80, 5743-5751.	1.4	164
120	Viral gastroenteritis in rotavirus negative hospitalized children <5 years of age from the independent states of the former Soviet Union. Infection, Genetics and Evolution, 2014, 28, 283-288.	1.0	21
121	Etiology of Childhood Diarrhea After Rotavirus Vaccine Introduction. Pediatric Infectious Disease Journal, 2014, 33, 1156-1163.	1.1	98
122	Infection control for norovirus. Clinical Microbiology and Infection, 2014, 20, 731-740.	2.8	132
123	Epidemiologic Implications of Asymptomatic Reinfection: A Mathematical Modeling Study of Norovirus. American Journal of Epidemiology, 2014, 179, 507-512.	1.6	70
124	Enteric bacteria promote human and mouse norovirus infection of B cells. Science, 2014, 346, 755-759.	6.0	689
125	Fluorinated TiO2 as an ambient light-activated virucidal surface coating material for the control of human norovirus. Journal of Photochemistry and Photobiology B: Biology, 2014, 140, 315-320.	1.7	59
126	Feline fecal virome reveals novel and prevalent enteric viruses. Veterinary Microbiology, 2014, 171, 102-111.	0.8	83

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127	Divergent Picobirnaviruses in Human Feces. Genome Announcements, 2014, 2, .	0.8	15
128	Noroviruses, Sapoviruses, and Astroviruses. , 2014, , 479-499.		2
129	Presence of Antibodies against Genogroup VI Norovirus in Humans. Virology Journal, 2013, 10, 176.	1.4	43
130	Emergence of New Pandemic GII.4 Sydney Norovirus Strain Correlates With Escape From Herd Immunity. Journal of Infectious Diseases, 2013, 208, 1877-1887.	1.9	151
131	Diagnostic performance of rectal swab versus bulk stool specimens for the detection of rotavirus and norovirus: Implications for outbreak investigations. Journal of Clinical Virology, 2013, 58, 678-682.	1.6	22
132	Norovirus and Medically Attended Gastroenteritis in U.S. Children. New England Journal of Medicine, 2013, 368, 1121-1130.	13.9	518
133	Proposal for a unified norovirus nomenclature and genotyping. Archives of Virology, 2013, 158, 2059-2068.	0.9	488
134	Human Norovirus Detection and Production, Quantification, and Storage of Virusâ€Like Particles. Current Protocols in Microbiology, 2013, 31, 15K.1.1-15K.1.45.	6.5	27
135	Genotype Gl.6 Norovirus, United States, 2010–2012. Emerging Infectious Diseases, 2013, 19, 1317-1320.	2.0	26
136	Effects and Clinical Significance of GII.4 Sydney Norovirus, United States, 2012–2013. Emerging Infectious Diseases, 2013, 19, 1231-1238.	2.0	67
137	Clinical Profile of Children with Norovirus Disease in Rotavirus Vaccine Era. Emerging Infectious Diseases, 2013, 19, 1691-1693.	2.0	33
138	Etiology of Viral Gastroenteritis in Children <5 Years of Age in the United States, 2008–2009. Journal of Infectious Diseases, 2013, 208, 790-800.	1.9	184
139	Emergence of a Norovirus GII.4 Strain Correlates with Changes in Evolving Blockade Epitopes. Journal of Virology, 2013, 87, 2803-2813.	1.5	140
140	Prevalence and genetic diversity of norovirus among patients with acute diarrhea in Guatemala. Journal of Medical Virology, 2013, 85, 1293-1298.	2.5	21
141	Sapovirus Gastroenteritis in Preschool Center, Puerto Rico, 2011. Emerging Infectious Diseases, 2013, 19, 174-175.	2.0	12
142	Norovirus Disease in the United States. Emerging Infectious Diseases, 2013, 19, 1198-1205.	2.0	478
143	Challenges of Culturing Human Norovirus in Three-Dimensional Organoid Intestinal Cell Culture Models. PLoS ONE, 2013, 8, e63485.	1.1	102
144	Sapovirus. , 2013, , 313-319.		2

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145	Risk Factors for Death among Children Less than 5 Years Old Hospitalized with Diarrhea in Rural Western Kenya, 2005–2007: A Cohort Study. PLoS Medicine, 2012, 9, e1001256.	3.9	79
146	The Etiology of Severe Acute Gastroenteritis Among Adults Visiting Emergency Departments in the United States. Journal of Infectious Diseases, 2012, 205, 1374-1381.	1.9	155
147	Antiviral Activity of Nucleoside Analogues against Norovirus. Antiviral Therapy, 2012, 17, 981-991.	0.6	63
148	Environmental transmission of norovirus gastroenteritis. Current Opinion in Virology, 2012, 2, 96-102.	2.6	244
149	Norovirus outbreak of probable waterborne transmission with high attack rate in a Guatemalan resort. Journal of Clinical Virology, 2012, 55, 8-11.	1.6	31
150	Experimental Inoculation of Juvenile Rhesus Macaques with Primate Enteric Caliciviruses. PLoS ONE, 2012, 7, e37973.	1.1	40
151	Sapovirus Outbreaks in Long-Term Care Facilities, Oregon and Minnesota, USA, 2002–2009. Emerging Infectious Diseases, 2012, 18, 873-876.	2.0	70
152	Monoclonal Antibody-Based Antigenic Mapping of Norovirus GII.4-2002. Journal of Virology, 2012, 86, 873-883.	1.5	113
153	Epidemiologic and Clinical Features of Other Enteric Viruses Associated with Acute Gastroenteritis in American Indian Infants. Journal of Pediatrics, 2012, 161, 110-115.e1.	0.9	33
154	Detection and molecular characterization of noroviruses and sapoviruses in children admitted to hospital with acute gastroenteritis in Vietnam. Journal of Medical Virology, 2012, 84, 290-297.	2.5	29
155	Development and evaluation of novel one-step TaqMan realtime RT-PCR assays for the detection and direct genotyping of genogroup I and II norovirusesâ <sup>†</sup> t. Journal of Clinical Virology, 2011, 50, 230-234.	1.6	22
156	An automated genotyping tool for enteroviruses and noroviruses. Journal of Clinical Virology, 2011, 51, 121-125.	1.6	673
157	Epidemiological and genetic characteristics of norovirus outbreaks in long-term care facilities, 2003–2006. Epidemiology and Infection, 2011, 139, 286-294.	1.0	33
158	Incidence of Acute Gastroenteritis and Role of Norovirus, Georgia, USA, 2004-2005. Emerging Infectious Diseases, 2011, 17, 1381-8.	2.0	124
159	Novel Surveillance Network for Norovirus Gastroenteritis Outbreaks, United States. Emerging Infectious Diseases, 2011, 17, 1389-95.	2.0	198
160	Novel GII.12 Norovirus Strain, United States, 2009-2010. Emerging Infectious Diseases, 2011, 17, 1516-8.	2.0	57
161	Impact of an Emergent Norovirus Variant in 2009 on Norovirus Outbreak Activity in the United States. Clinical Infectious Diseases, 2011, 53, 568-571.	2.9	105
162	Detection of GI and GII Noroviruses in Ground Water Using Ultrafiltration and TaqMan Real-time RT-PCR. Food and Environmental Virology, 2010, 2, 218-224.	1.5	40

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163	Comparative Efficacy of Seven Hand Sanitizers against Murine Norovirus, Feline Calicivirus, and Gll.4 Norovirus. Journal of Food Protection, 2010, 73, 2232-2238.	0.8	131
164	Redefining Outcome of First Seizures by Acute Illness. Pediatrics, 2010, 126, e1477-e1484.	1.0	16
165	A Norovirus Vaccine on the Horizon?. Journal of Infectious Diseases, 2010, 202, 1623-1625.	1.9	26
166	Molecular Epidemiology of Genogroup II-Genotype 4 Noroviruses in the United States between 1994 and 2006. Journal of Clinical Microbiology, 2010, 48, 168-177.	1.8	165
167	Diagnostic Accuracy and Analytical Sensitivity of IDEIA Norovirus Assay for Routine Screening of Human Norovirus. Journal of Clinical Microbiology, 2010, 48, 2770-2778.	1.8	62
168	Novel Norovirus in Dogs with Diarrhea. Emerging Infectious Diseases, 2010, 16, 980-982.	2.0	125
169	Norovirus Illness Is a Global Problem: Emergence and Spread of Norovirus GII.4 Variants, 2001–2007. Journal of Infectious Diseases, 2009, 200, 802-812.	1.9	596
170	Herd Immunity to GII.4 Noroviruses Is Supported by Outbreak Patient Sera. Journal of Virology, 2009, 83, 5363-5374.	1.5	92
171	Identification of a Novel Astrovirus (Astrovirus VA1) Associated with an Outbreak of Acute Gastroenteritis. Journal of Virology, 2009, 83, 10836-10839.	1.5	190
172	Gene Mapping and Phylogenetic Analysis of the Complete Genome from 30 Single-Stranded RNA Male-Specific Coliphages (Family <i>Leviviridae</i> ). Journal of Virology, 2009, 83, 11233-11243.	1.5	32
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