

Helene Norder

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

110
papers

6,034
citations

40
h-index

77
g-index

113
ext. papers

6,677
ext. citations

8.7
avg, IF

5.38
L-index

#	Paper	IF	Citations
110	Thalassemia Patients from Baluchistan in Pakistan Are Infected with Multiple Hepatitis B or C Virus Strains. <i>American Journal of Tropical Medicine and Hygiene</i> , 2021 ,	3.2	1
109	Four novel picornaviruses detected in Magellanic Penguins (<i>Spheniscus magellanicus</i>) in Chile. <i>Virology</i> , 2021 , 560, 116-123	3.6	2
108	Analysis of Multiple Liver Explant Pieces Reveals that Levels of Hepatitis Delta Virus RNA Are Independent of Hepatitis B Virus Expression. <i>Hepatology Communications</i> , 2021 , 5, 1964-1966	6	
107	A new assay for quantitative detection of hepatitis A virus. <i>Journal of Virological Methods</i> , 2021 , 288, 114010	2.6	4
106	Surveillance of wastewater revealed peaks of SARS-CoV-2 preceding those of hospitalized patients with COVID-19. <i>Water Research</i> , 2021 , 189, 116620	12.5	53
105	An Evaluation of Hepatitis E Virus Molecular Typing Methods.. <i>Clinical Chemistry</i> , 2021 , 68, 181-191	5.5	2
104	Quantification of viral RNA in multiple pieces of explant liver tissue shows distinct focal differences of hepatitis B infection. <i>Journal of Infectious Diseases</i> , 2021 ,	7	1
103	Deep sequencing of liver explant transcriptomes reveals extensive expression from integrated hepatitis B virus DNA. <i>Journal of Viral Hepatitis</i> , 2020 , 27, 1162-1170	3.4	7
102	Hepatitis E virus is an infrequent but potentially serious infection in allogeneic hematopoietic stem cell transplant recipients. <i>Bone Marrow Transplantation</i> , 2020 , 55, 1255-1263	4.4	1
101	Update: proposed reference sequences for subtypes of hepatitis E virus (species). <i>Journal of General Virology</i> , 2020 , 101, 692-698	4.9	92
100	Serum-IgG responses to SARS-CoV-2 after mild and severe COVID-19 infection and analysis of IgG non-responders. <i>PLoS ONE</i> , 2020 , 15, e0241104	3.7	101
99	Variations among Viruses in Influent Water and Effluent Water at a Wastewater Plant over One Year as Assessed by Quantitative PCR and Metagenomics. <i>Applied and Environmental Microbiology</i> , 2020 , 86,	4.8	9
98	Molecular evolution of coxsackievirus A24v in Cuba over 23-years, 1986-2009. <i>Scientific Reports</i> , 2020 , 10, 13761	4.9	1
97	Hepatitis E virus genotype 3 strains and a plethora of other viruses detected in raw and still in tap water. <i>Water Research</i> , 2020 , 168, 115141	12.5	15
96	Serum-IgG responses to SARS-CoV-2 after mild and severe COVID-19 infection and analysis of IgG non-responders 2020 , 15, e0241104		
95	Serum-IgG responses to SARS-CoV-2 after mild and severe COVID-19 infection and analysis of IgG non-responders 2020 , 15, e0241104		
94	Serum-IgG responses to SARS-CoV-2 after mild and severe COVID-19 infection and analysis of IgG non-responders 2020 , 15, e0241104		

93	Serum-IgG responses to SARS-CoV-2 after mild and severe COVID-19 infection and analysis of IgG non-responders 2020 , 15, e0241104		
92	Serum-IgG responses to SARS-CoV-2 after mild and severe COVID-19 infection and analysis of IgG non-responders 2020 , 15, e0241104		
91	Serum-IgG responses to SARS-CoV-2 after mild and severe COVID-19 infection and analysis of IgG non-responders 2020 , 15, e0241104		
90	Hepatitis E virus genotype 3 is associated with gallstone-related disease. <i>Scandinavian Journal of Gastroenterology</i> , 2019 , 54, 1269-1273	2.4	1
89	High Frequency of Either Altered Pre-Core StartCodon or Weakened Kozak Sequence in the CorePromoter Region in Hepatitis B Virus A1 Strainsfrom Rwanda. <i>Genes</i> , 2019 , 10,	4.2	2
88	Hepatitis A and E virus infections have different epidemiological patterns in Rwanda. <i>International Journal of Infectious Diseases</i> , 2019 , 86, 12-14	10.5	3
87	Hepatitis E virus strains infecting Swedish domestic pigs are unique for each pig farm and remain in the farm for at least 2years. <i>Transboundary and Emerging Diseases</i> , 2019 , 66, 1314-1323	4.2	9
86	Genetically similar hepatitis E virus strains infect both humans and wild boars in the Barcelona area, Spain, and Sweden. <i>Transboundary and Emerging Diseases</i> , 2019 , 66, 978-985	4.2	19
85	Differential removal of human pathogenic viruses from sewage by conventional and ozone treatments. <i>International Journal of Hygiene and Environmental Health</i> , 2018 , 221, 479-488	6.9	45
84	Hepatitis E Virus Genotype 3 Genomes from RNA-Positive but Serologically Negative Plasma Donors Have CUG as the Start Codon for ORF3. <i>Intervirology</i> , 2018 , 61, 96-103	2.5	12
83	Inosine Triphosphate Pyrophosphatase Dephosphorylates Ribavirin Triphosphate and Reduced Enzymatic Activity Potentiates Mutagenesis in Hepatitis C Virus. <i>Journal of Virology</i> , 2018 , 92,	6.6	14
82	Hepatitis B virus strains from Rwandan blood donors are genetically similar and form one clade within subgenotype A1. <i>BMC Infectious Diseases</i> , 2017 , 17, 32	4	8
81	High seroprevalence against hepatitis E virus in patients with chronic hepatitis C virus infection. <i>Journal of Clinical Virology</i> , 2017 , 88, 39-45	14.5	14
80	Low prevalence of hepatitis C virus RNA in blood donors with anti-hepatitis C virus reactivity in Rwanda. <i>Transfusion</i> , 2017 , 57, 2420-2432	2.9	7
79	Diagnostic Performance of Five Assays for Anti-Hepatitis E Virus IgG and IgM in a Large Cohort Study. <i>Journal of Clinical Microbiology</i> , 2016 , 54, 549-55	9.7	70
78	Markers for Ongoing or Previous Hepatitis E Virus Infection Are as Common in Wild Ungulates as in Humans in Sweden. <i>Viruses</i> , 2016 , 8,	6.2	27
77	A Single Lineage of Hepatitis E Virus Causes Both Outbreaks and Sporadic Hepatitis in Sudan. <i>Viruses</i> , 2016 , 8,	6.2	11
76	Molecular characterization of HCV in a Swedish county over 8 years (2002-2009) reveals distinct transmission patterns. <i>Infection Ecology and Epidemiology</i> , 2016 , 6, 30670	4.3	2

75	Can sequence data predict enterovirus D68 infection outcome?. <i>Lancet Infectious Diseases, The</i> , 2015 , 15, 620-1	25.5	5
74	Chronic hepatitis E infection with an emerging virus strain in a heart transplant recipient successfully treated with ribavirin: a case report. <i>Journal of Medical Case Reports</i> , 2015 , 9, 180	1.2	6
73	High incidence of the hepatitis C virus recombinant 2k/1b in Georgia: Recommendations for testing and treatment. <i>Hepatology Research</i> , 2015 , 45, 1292-8	5.1	23
72	Complete coding regions of the prototypes enterovirus B93 and C95: phylogenetic analyses of the P1 and P3 regions of EV-B and EV-C strains. <i>Journal of Medical Virology</i> , 2015 , 87, 485-97	19.7	7
71	High prevalence of hepatitis e virus in Swedish moose—a phylogenetic characterization and comparison of the virus from different regions. <i>PLoS ONE</i> , 2015 , 10, e0122102	3.7	22
70	Molecular analysis of enterovirus in Cameroon by partial 5'UTR-VP4 gene sequencing reveals a high genetic diversity and frequency of infections. <i>Journal of Medical Virology</i> , 2014 , 86, 2092-101	19.7	7
69	Novel hepatitis E like virus found in Swedish moose. <i>Journal of General Virology</i> , 2014 , 95, 557-570	4.9	50
68	Should we use oral polio vaccine in Europe?. <i>Lancet, The</i> , 2014 , 383, 1037-8	4.0	1
67	Reduced prevalence of HBsAg variants following a successful immunization program in China. <i>Journal of Virology</i> , 2014 , 88, 4605-6	6.6	6
66	Detection of pathogenic viruses in sewage provided early warnings of hepatitis A virus and norovirus outbreaks. <i>Applied and Environmental Microbiology</i> , 2014 , 80, 6771-81	4.8	224
65	Shift of Enterovirus species among children in Cameroon—identification of a new enterovirus, EV-A119. <i>Journal of Clinical Virology</i> , 2013 , 58, 227-32	14.5	22
64	Patient-to-Patient Transmission of Hepatitis C at Iranian Thalassemia Centers Shown by Genetic Characterization of Viral Strains. <i>Hepatitis Monthly</i> , 2013 , 13, e7699	1.8	11
63	Neonatal transfusion-transmitted hepatitis C virus infection following a pre-seroconversion window-phase donation in Sweden. <i>Scandinavian Journal of Infectious Diseases</i> , 2013 , 45, 796-9		7
62	3C protease of enterovirus 68: structure-based design of Michael acceptor inhibitors and their broad-spectrum antiviral effects against picornaviruses. <i>Journal of Virology</i> , 2013 , 87, 4339-51	6.6	65
61	Seasonal variation in TP53 R249S-mutated serum DNA with aflatoxin exposure and hepatitis B virus infection. <i>Environmental Health Perspectives</i> , 2011 , 119, 1635-40	8.4	27
60	Hepatitis B virus impairs TLR9 expression and function in plasmacytoid dendritic cells. <i>PLoS ONE</i> , 2011 , 6, e26315	3.7	109
59	Picornavirus non-structural proteins as targets for new anti-virals with broad activity. <i>Antiviral Research</i> , 2011 , 89, 204-18	10.8	59
58	Structural basis for antiviral inhibition of the main protease, 3C, from human enterovirus 93. <i>Journal of Virology</i> , 2011 , 85, 10764-73	6.6	16

57	Mutations in TP53 and CTNNB1 in Relation to Hepatitis B and C Infections in Hepatocellular Carcinomas from Thailand. <i>Hepatitis Research and Treatment</i> , 2011 , 2011, 697162		10
56	Protein Tpr is required for establishing nuclear pore-associated zones of heterochromatin exclusion. <i>EMBO Journal</i> , 2010 , 29, 1659-73	13	126
55	High prevalence of hepatitis B virus genotype E in Northern Madagascar indicates a West-African lineage. <i>Journal of Medical Virology</i> , 2010 , 82, 1515-26	19.7	13
54	The 2C putative helicase of echovirus 30 adopts a hexameric ring-shaped structure. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2010 , 66, 1116-20		12
53	Under construction: classification of hepatitis B virus genotypes and subgenotypes. <i>Intervirology</i> , 2009 , 52, 323-5	2.5	41
52	CTL escape mutations of core protein are more frequent in strains of HBeAg negative patients with low levels of HBV DNA. <i>Journal of Clinical Virology</i> , 2009 , 46, 259-64	14.5	21
51	D2: major subgenotype of hepatitis B virus in Russia and the Baltic region. <i>Journal of General Virology</i> , 2008 , 89, 1829-1839	4.9	54
50	Molecular characterization and phylogenetic analysis of the complete genome of a hepatitis E virus from European swine. <i>Virus Genes</i> , 2008 , 37, 39-48	2.3	20
49	Molecular epidemiological studies show that hepatitis A virus is endemic among active homosexual men in Europe. <i>Journal of Medical Virology</i> , 2007 , 79, 356-65	19.7	55
48	New enteroviruses, EV-93 and EV-94, associated with acute flaccid paralysis in the Democratic Republic of the Congo. <i>Journal of Medical Virology</i> , 2007 , 79, 393-400	19.7	64
47	Genetic characterization of hepatitis C virus strains in Estonia: fluctuations in the predominating subtype with time. <i>Journal of Medical Virology</i> , 2007 , 79, 374-82	19.7	55
46	Rapid diagnosis of acute hemorrhagic conjunctivitis due to coxsackievirus A24 variant by real-time one-step RT-PCR. <i>Journal of Virological Methods</i> , 2007 , 142, 89-94	2.6	6
45	Universal detection of hepatitis E virus by two real-time PCR assays: TaqMan and Primer-Probe Energy Transfer. <i>Journal of Virological Methods</i> , 2007 , 146, 226-35	2.6	85
44	Enterovirus 94, a proposed new serotype in human enterovirus species D. <i>Journal of General Virology</i> , 2007 , 88, 849-858	4.9	54
43	Echovirus 6 strains derived from a clinical isolate show differences in haemagglutination ability and cell entry pathway. <i>Virus Research</i> , 2007 , 130, 1-9	6.4	10
42	Characterization of HCV strains in an oncohematological pediatric department reveals little horizontal transmission but multiple introductions by un-screened blood products in the past. <i>Journal of Medical Virology</i> , 2006 , 78, 1411-22	19.7	4
41	Removal of viruses, parasitic protozoa and microbial indicators in conventional and membrane processes in a wastewater pilot plant. <i>Water Research</i> , 2006 , 40, 1449-57	12.5	183
40	T1764G1766 core promoter double mutants are restricted to Hepatitis B virus strains with an A1757 and are common in genotype D. <i>Journal of General Virology</i> , 2005 , 86, 2451-2458	4.9	31

39	Divergent genetic evolution of hemagglutinin in influenza A H1N1 and A H1N2 subtypes isolated in the south-France since the winter of 2001-2002. <i>Journal of Clinical Virology</i> , 2005 , 33, 230-6	14.5	11
38	Outbreak of acute hemorrhagic conjunctivitis in French Guiana and West Indies caused by coxsackievirus A24 variant: phylogenetic analysis reveals Asian import. <i>Journal of Medical Virology</i> , 2005 , 75, 559-65	19.7	35
37	Full genome analysis of hepatitis B virus genotype E strains from South-Western Africa and Madagascar reveals low genetic variability. <i>Journal of Medical Virology</i> , 2005 , 77, 47-52	19.7	56
36	Full-length open reading frame of a recombinant hepatitis C virus strain from St Petersburg: proposed mechanism for its formation. <i>Journal of General Virology</i> , 2004 , 85, 1853-1857	4.9	68
35	Molecular identification and characterization of two proposed new enterovirus serotypes, EV74 and EV75. <i>Journal of General Virology</i> , 2004 , 85, 3205-3212	4.9	92
34	Hepatitis B virus genotype D strains from Estonia share sequence similarity with strains from Siberia and may specify ayw4. <i>Journal of Medical Virology</i> , 2004 , 74, 221-7	19.7	31
33	Molecular epidemiology of hepatitis C virus in Iran as reflected by phylogenetic analysis of the NS5B region. <i>Journal of Medical Virology</i> , 2004 , 74, 246-52	19.7	56
32	Genetic diversity of hepatitis B virus strains derived worldwide: genotypes, subgenotypes, and HBsAg subtypes. <i>Intervirology</i> , 2004 , 47, 289-309	2.5	676
31	Sequential changes in hepatitis A virus genotype distribution in Estonia during 1994 to 2001. <i>Journal of Medical Virology</i> , 2003 , 70, 187-93	19.7	42
30	The T(1858) variant predisposing to the precore stop mutation correlates with one of two major genotype F hepatitis B virus clades. <i>Journal of General Virology</i> , 2003 , 84, 2083-2087	4.9	42
29	Sequencing of Untypable Enteroviruses reveals two new types, EV-77 and EV-78, within human enterovirus type B and substitutions in the BC loop of the VP1 protein for known types. <i>Journal of General Virology</i> , 2003 , 84, 827-836	4.9	114
28	A natural intergenotypic recombinant of hepatitis C virus identified in St. Petersburg. <i>Journal of Virology</i> , 2002 , 76, 4034-43	6.6	261
27	Open reading frame sequence of an Asian enterovirus 73 strain reveals that the prototype from California is recombinant. <i>Journal of General Virology</i> , 2002 , 83, 1721-1728	4.9	36
26	Genotype H: a new Amerindian genotype of hepatitis B virus revealed in Central America. <i>Journal of General Virology</i> , 2002 , 83, 2059-2073	4.9	556
25	Hepatitis B virus genotypes and HBsAg subtypes in refugees and injection drug users in the United States determined by LiPA and monoclonal EIA. <i>Journal of Medical Virology</i> , 2001 , 64, 305-11	19.7	35
24	Presumed common source outbreaks of hepatitis A in an endemic area confirmed by limited sequencing within the VP1 region. <i>Journal of Medical Virology</i> , 2001 , 65, 449-456	19.7	32
23	Homotypic echoviruses share aminoterminal VP1 sequence homology applicable for typing. <i>Journal of Medical Virology</i> , 2001 , 63, 35-44	19.7	77
22	Prevalence and incidence of bloodborne viral infections among Danish prisoners. <i>European Journal of Epidemiology</i> , 2000 , 16, 1043-9	12.1	60

21	An aberrant genotype revealed in recombinant hepatitis B virus strains from Vietnam. <i>Journal of General Virology</i> , 2000 , 81, 2267-2272	4.9	104
20	Antigenic diversity of hepatitis B virus strains of genotype F in Amerindians and other population groups from Venezuela. <i>Journal of Clinical Microbiology</i> , 1998 , 36, 648-51	9.7	71
19	Molecular epidemiology of hepatitis B virus in Central America reflected in the genetic variability of the small S gene. <i>Journal of Infectious Diseases</i> , 1997 , 176, 851-8	7	70
18	Genotype F prevails in HBV infected patients of Hispanic origin in Central America and may carry the precore stop mutant 1997 , 51, 305-312		93
17	High prevalence of GB virus C strains genetically related to strains with Asian origin in Nicaraguan hemophiliacs 1997 , 52, 149-155		21
16	Complete sequencing of a gibbon hepatitis B virus genome reveals a unique genotype distantly related to the chimpanzee hepatitis B virus. <i>Virology</i> , 1996 , 218, 214-23	3.6	89
15	Genetic relatedness of Sindbis virus strains from Europe, Middle East, and Africa. <i>Virology</i> , 1996 , 222, 440-5	3.6	46
14	Subtypes, genotypes and molecular epidemiology of the hepatitis B virus as reflected by sequence variability of the S-gene. <i>Intervirology</i> , 1995 , 38, 24-34	2.5	323
13	Complete genomes, phylogenetic relatedness, and structural proteins of six strains of the hepatitis B virus, four of which represent two new genotypes. <i>Virology</i> , 1994 , 198, 489-503	3.6	657
12	Contrasting Roles of Rivers and Wells as Sources of Drinking Water on Attack and Fatality Rates in a Hepatitis E Epidemic in Somalia. <i>American Journal of Tropical Medicine and Hygiene</i> , 1994 , 51, 466-474	3.2	74
11	Important role of hepatitis C virus infection as a cause of chronic liver disease in Somalia. <i>Scandinavian Journal of Infectious Diseases</i> , 1993 , 25, 559-64		17
10	Interferon Alpha-2b Treatment in an HIV-infected Patient with Hepatitis B Virus Induced Nephrotic Syndrome. <i>Scandinavian Journal of Infectious Diseases</i> , 1993 , 25, 133-135		7
9	Complete nucleotide sequences of six hepatitis B viral genomes encoding the surface antigen subtypes ayw4, adw4q-, and adrq- and their phylogenetic classification. <i>Archives of Virology Supplementum</i> , 1993 , 8, 189-99		13
8	Detection of HBV DNA by PCR in serum from an HBsAg negative blood donor implicated in cases of post-transfusion hepatitis B. <i>Archives of Virology Supplementum</i> , 1992 , 4, 116-8		7
7	Minor role of hepatitis B virus in the causation of chronic liver disease in Somalia indicated by a case-control study. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 1991 , 85, 104-8	2	4
6	Comparison of class and subclass distribution of antibodies to the hepatitis B core and B e antigens in chronic hepatitis B. <i>Journal of Medical Virology</i> , 1990 , 30, 1-6	19.7	26
5	Typing of hepatitis B virus genomes by a simplified polymerase chain reaction. <i>Journal of Medical Virology</i> , 1990 , 31, 215-21	19.7	38
4	High frequency of hepatitis B virus DNA in anti-HBe positive sera on longitudinal follow-up of patients with renal transplants and chronic hepatitis B. <i>Journal of Medical Virology</i> , 1989 , 27, 322-8	19.7	18

- 3 A solid-phase reverse immunosorbent test for the detection of enterovirus IgM. *Journal of Virological Methods*, **1988**, 20, 73-82 2.6 26
- 2 Epidemiology of hepatitis B in Somalia: inference from a cross-sectional survey of serological markers. *Transactions of the Royal Society of Tropical Medicine and Hygiene*, **1987**, 81, 824-8 2 6
- 1 Reactivity of sera from patients with liver disease with glutaraldehyde-treated erythrocytes: role for false positive results in hemagglutination tests. *Journal of Immunological Methods*, **1984**, 71, 163-73 2.5