

Simona De Grazia

List of Publications by Year in descending order

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68
papers

1,988
citations

201385

27
h-index

264894

42
g-index

68
all docs

68
docs citations

68
times ranked

1888
citing authors

#	ARTICLE	IF	CITATIONS
1	Review of group A rotavirus strains reported in swine and cattle. <i>Veterinary Microbiology</i> , 2013, 165, 190-199.	0.8	195
2	Relationships among porcine and human P[6] rotaviruses: Evidence that the different human P[6] lineages have originated from multiple interspecies transmission events. <i>Virology</i> , 2006, 344, 509-519.	1.1	119
3	Heterogeneity and Temporal Dynamics of Evolution of G1 Human Rotaviruses in a Settled Population. <i>Journal of Virology</i> , 2006, 80, 10724-10733.	1.5	119
4	Multiple reassortment and interspecies transmission events contribute to the diversity of feline, canine and feline/canine-like human group A rotavirus strains. <i>Infection, Genetics and Evolution</i> , 2011, 11, 1396-1406.	1.0	105
5	Viral gastroenteritis in children hospitalised in Sicily, Italy. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2006, 25, 570-575.	1.3	80
6	Visceral leishmaniasis: host-parasite interactions and clinical presentation in the immunocompetent and in the immunocompromised host. <i>International Journal of Infectious Diseases</i> , 2013, 17, e572-e576.	1.5	71
7	Genetic Variability among Serotype G4 Italian Human Rotaviruses. <i>Journal of Clinical Microbiology</i> , 2005, 43, 1420-1425.	1.8	47
8	A feline rotavirus G3P[9] carries traces of multiple reassortment events and resembles rare human G3P[9] rotaviruses. <i>Journal of General Virology</i> , 2011, 92, 1214-1221.	1.3	47
9	Canine-Origin G3P[3] Rotavirus Strain in Child with Acute Gastroenteritis. <i>Emerging Infectious Diseases</i> , 2007, 13, 1091-1093.	2.0	45
10	Genetic heterogeneity of porcine enteric caliciviruses identified from diarrhoeic piglets. <i>Virus Genes</i> , 2008, 36, 365-373.	0.7	45
11	Genomic characterization of a novel group A lamb rotavirus isolated in Zaragoza, Spain. <i>Virus Genes</i> , 2008, 37, 250-265.	0.7	45
12	Evidence for Recombination between Pandemic GII.4 Norovirus Strains New Orleans 2009 and Sydney 2012: Fig 1. <i>Journal of Clinical Microbiology</i> , 2013, 51, 3855-3857.	1.8	45
13	Evolution of DS-1-like human G2P[4] rotaviruses assessed by complete genome analyses. <i>Journal of General Virology</i> , 2014, 95, 91-109.	1.3	44
14	Norovirus and Gastroenteritis in Hospitalized Children, Italy. <i>Emerging Infectious Diseases</i> , 2007, 13, 1389-1391.	2.0	43
15	Unusual Assortment of Segments in 2 Rare Human Rotavirus Genomes. <i>Emerging Infectious Diseases</i> , 2010, 16, 859-862.	2.0	43
16	Clinically-based determination of safe DNAemia cutoff levels for preemptive therapy or human cytomegalovirus infections in solid organ and hematopoietic stem cell transplant recipients. <i>Journal of Medical Virology</i> , 2004, 73, 412-418.	2.5	36
17	Nationwide surveillance study of human astrovirus infections in an Italian paediatric population. <i>Epidemiology and Infection</i> , 2013, 141, 524-528.	1.0	34
18	Genetic Heterogeneity and Recombination in Human Type 2 Astroviruses. <i>Journal of Clinical Microbiology</i> , 2012, 50, 3760-3764.	1.8	33

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19	Novel recombinant GII.P16_GII.13 and GII.P16_GII.3 norovirus strains in Italy. <i>Virus Research</i> , 2014, 188, 142-145.	1.1	33
20	Surveillance of human astrovirus circulation in Italy 2002-2005: emergence of lineage 2c strains. <i>Clinical Microbiology and Infection</i> , 2011, 17, 97-101.	2.8	32
21	HLA and Killer Cell Immunoglobulin-like Receptors Influence the Natural Course of CMV Infection. <i>Journal of Infectious Diseases</i> , 2014, 210, 1083-1089.	1.9	32
22	Molecular epidemiology of astrovirus infection in Italian children with gastroenteritis. <i>Clinical Microbiology and Infection</i> , 2004, 10, 1025-1029.	2.8	31
23	Lineage diversification and recombination in type-4 human astroviruses. <i>Infection, Genetics and Evolution</i> , 2013, 20, 330-335.	1.0	30
24	Recombinant norovirus GII.g/GII.12 gastroenteritis in children. <i>Infection, Genetics and Evolution</i> , 2012, 12, 169-174.	1.0	29
25	Analysis of the ORF2 of human astroviruses reveals lineage diversification, recombination and rearrangement and provides the basis for a novel sub-classification system. <i>Archives of Virology</i> , 2014, 159, 3185-3196.	0.9	29
26	Molecular characterization of the genotype G9 human rotavirus strains recovered in Palermo, Italy, during the winter of 1999â€“2000. <i>Epidemiology and Infection</i> , 2004, 132, 343-349.	1.0	27
27	G2 rotavirus infections in an infantile population of the South of Italy: Variability of viral strains over time. <i>Journal of Medical Virology</i> , 2005, 77, 587-594.	2.5	27
28	Detection of a Porcine-Like Rotavirus in a Child with Enteritis in Italy. <i>Journal of Clinical Microbiology</i> , 2008, 46, 3501-3507.	1.8	27
29	Molecular characterization of genotype G6 human rotavirus strains detected in Italy from 1986 to 2009. <i>Infection, Genetics and Evolution</i> , 2011, 11, 1449-1455.	1.0	27
30	Antibodies Responses to SARS-CoV-2 in a Large Cohort of Vaccinated Subjects and Seropositive Patients. <i>Vaccines</i> , 2021, 9, 714.	2.1	25
31	Detection of the norovirus variants GII.4 hunter and GIIb/hilversum in Italian children with gastroenteritis. <i>Journal of Medical Virology</i> , 2006, 78, 1656-1662.	2.5	24
32	Norovirus GII.4/Sydney/2012 in Italy, Winter 2012â€“2013. <i>Emerging Infectious Diseases</i> , 2013, 19, 1348-1349.	2.0	23
33	Human cytomegalovirus glycoprotein B genotypes in immunocompetent, immunocompromised, and congenitally infected Italian populations. <i>Archives of Virology</i> , 2003, 148, 547-554.	0.9	22
34	Emerging GII.4 norovirus variants affect children with diarrhea in Palermo, Italy in 2006. <i>Journal of Medical Virology</i> , 2009, 81, 139-145.	2.5	22
35	Investigation and control of a Norovirus outbreak of probable waterborne transmission through a municipal groundwater system. <i>Journal of Water and Health</i> , 2014, 12, 452-464.	1.1	22
36	Diversity of human rotaviruses detected in Sicily, Italy, over a 5-year period (2001â€“2005). <i>Archives of Virology</i> , 2007, 152, 833-837.	0.9	21

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37	Rare AU-1-Like G3P[9] Human Rotaviruses with a Kun-Like NSP4 Gene Detected in Children with Diarrhea in Italy. <i>Journal of Clinical Microbiology</i> , 2008, 46, 357-360.	1.8	20
38	Analysis of early strains of the norovirus pandemic variant GII.4 Sydney 2012 identifies mutations in adaptive sites of the capsid protein. <i>Virology</i> , 2014, 450-451, 355-358.	1.1	20
39	Artificial chromosome libraries of <i>Streptomyces coelicolor</i> A3(2) and <i>Planobispora rosea</i> . <i>FEMS Microbiology Letters</i> , 2003, 218, 181-186.	0.7	19
40	Genotyping of GII.4 and GIIB norovirus RT-PCR amplicons by RFLP analysis. <i>Journal of Virological Methods</i> , 2008, 147, 250-256.	1.0	17
41	Data mining from a 27-years rotavirus surveillance in Palermo, Italy. <i>Infection, Genetics and Evolution</i> , 2014, 28, 377-384.	1.0	17
42	Norovirus GII.17 as Major Epidemic Strain in Italy, Winter 2015-2016. <i>Emerging Infectious Diseases</i> , 2017, 23, 1206-1208.	2.0	15
43	Parvovirus B19 and "cryptogenic"™ chronic hepatitis. <i>Journal of Hepatology</i> , 2003, 38, 375-376.	1.8	13
44	Assessing the burden of viral co-infections in acute gastroenteritis in children: An eleven-year-long investigation. <i>Journal of Clinical Virology</i> , 2020, 129, 104513.	1.6	13
45	Genetic characterization of G3 rotaviruses detected in Italian children in the years 1993-2005. <i>Journal of Medical Virology</i> , 2009, 81, 2089-2095.	2.5	12
46	Assignment of the group A rotavirus NSP4 gene into genotypes using a hemi-nested multiplex PCR assay: a rapid and reproducible assay for strain surveillance studies. <i>Journal of Medical Microbiology</i> , 2009, 58, 303-311.	0.7	11
47	Full-genome sequencing of a Hungarian canine G3P[3] Rotavirus A strain reveals high genetic relatedness with a historic Italian human strain. <i>Virus Genes</i> , 2015, 50, 310-315.	0.7	11
48	Performance analysis of two immunochromatographic assays for the diagnosis of rotavirus infection. <i>Journal of Virological Methods</i> , 2017, 243, 50-54.	1.0	11
49	Analysis of GII.P7 and GII.6 noroviruses circulating in Italy during 2011-2016 reveals a replacement of lineages and complex recombination history. <i>Infection, Genetics and Evolution</i> , 2019, 75, 103991.	1.0	11
50	Emergence in 2017-2019 of novel reassortant equine-like G3 rotavirus strains in Palermo, Sicily. <i>Transboundary and Emerging Diseases</i> , 2022, 69, 813-835.	1.3	11
51	Sentinel hospital-based surveillance for norovirus infection in children with gastroenteritis between 2015 and 2016 in Italy. <i>PLoS ONE</i> , 2018, 13, e0208184.	1.1	10
52	Impact of Vaccination on Rotavirus Genotype Diversity: A Nearly Two-Decade-Long Epidemiological Study before and after Rotavirus Vaccine Introduction in Sicily, Italy. <i>Pathogens</i> , 2022, 11, 424.	1.2	10
53	A case of spotted fever rickettsiosis in a human immunodeficiency virus-positive patient. <i>Journal of Medical Microbiology</i> , 2013, 62, 1363-1364.	0.7	8
54	Epidemiological dynamics of norovirus GII.4 variant New Orleans 2009. <i>Journal of General Virology</i> , 2015, 96, 2919-2927.	1.3	8

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55	Temporal variation in the distribution of type-1 human astrovirus lineages in a settled population over 14 years. <i>Archives of Virology</i> , 2016, 161, 1633-1637.	0.9	8
56	Assessment of SARS-CoV-2 RNA shedding in semen of 36 males with symptomatic, asymptomatic, and convalescent infection during the first and second wave of COVID-19 pandemic in Italy. <i>Asian Journal of Andrology</i> , 2022, 24, 135.	0.8	8
57	Identification of a multi-reassortant G12P[9] rotavirus with novel VP1, VP2, VP3 and NSP2 genotypes in a child with acute gastroenteritis. <i>Infection, Genetics and Evolution</i> , 2015, 35, 34-37.	1.0	7
58	Recombinant GII.P16 genotype challenges RT-PCR-based typing in region A of norovirus genome. <i>Journal of Infection</i> , 2021, 83, 69-75.	1.7	7
59	Molecular evolutionary analysis of type-1 human astroviruses identifies putative sites under selection pressure on the capsid protein. <i>Infection, Genetics and Evolution</i> , 2018, 58, 199-208.	1.0	6
60	Is Italian population protected from Poliovirus? Results of a seroprevalence survey in Florence, Italy. <i>Human Vaccines and Immunotherapeutics</i> , 2018, 14, 2248-2253.	1.4	6
61	Molecular Characterization of Coxsackievirus B5 Isolates from Sewage, Italy 2016-2017. <i>Food and Environmental Virology</i> , 2019, 11, 440-445.	1.5	6
62	Neutralizing Antibodies Response against SARS-CoV-2 Variants of Concern Elicited by Prior Infection or mRNA BNT162b2 Vaccination. <i>Vaccines</i> , 2022, 10, 874.	2.1	5
63	VP7 and VP4 Sequence Analyses of Rotavirus Strains From Italian Children With Viraemia and Acute Diarrhoea. <i>Journal of Pediatric Gastroenterology and Nutrition</i> , 2010, 50, 114-116.	0.9	4
64	Complete genome analysis of contemporary G12P[8] rotaviruses reveals heterogeneity within Wa-like genomic constellation. <i>Infection, Genetics and Evolution</i> , 2016, 44, 85-93.	1.0	4
65	Performance evaluation of gastrointestinal viral ELite panel multiplex RT-PCR assay for the diagnosis of rotavirus, adenovirus and astrovirus infection. <i>Journal of Virological Methods</i> , 2019, 268, 48-52.	1.0	4
66	Differing kinetics of anti-spike protein IgGs and neutralizing antibodies against SARS-CoV-2 after Comirnaty (BNT162b2) immunization. <i>Journal of Applied Microbiology</i> , 2022, , .	1.4	4
67	Evaluation of the diagnostic performances of two commercially available assays for the detection of enteric adenovirus antigens. <i>Diagnostic Microbiology and Infectious Disease</i> , 2021, 101, 115459.	0.8	2
68	Performance evaluation of a newly developed molecular assay for the accurate diagnosis of gastroenteritis associated with norovirus of genogroup II. <i>Archives of Virology</i> , 2018, 163, 3377-3381.	0.9	1