

Bernhard Ehlers

List of Publications by Year in descending order

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71
papers

3,544
citations

147566

31
h-index

138251

58
g-index

73
all docs

73
docs citations

73
times ranked

3429
citing authors

#	ARTICLE	IF	CITATIONS
1	Multiple DNA viruses identified in multimammate mouse (<i>Mastomys natalensis</i>) populations from across regions of sub-Saharan Africa. <i>Archives of Virology</i> , 2020, 165, 2291-2299.	0.9	3
2	Search for polyoma-, herpes-, and bornaviruses in squirrels of the family Sciuridae. <i>Virology Journal</i> , 2020, 17, 42.	1.4	11
3	Promoter activity of Merkel cell Polyomavirus variants in human dermal fibroblasts and a Merkel cell carcinoma cell line. <i>Virology Journal</i> , 2020, 17, 54.	1.4	7
4	Cytomegalovirus distribution and evolution in hominines. <i>Virus Evolution</i> , 2019, 5, vez015.	2.2	26
5	Novel Polyomaviruses in Mammals from Multiple Orders and Reassessment of Polyomavirus Evolution and Taxonomy. <i>Viruses</i> , 2019, 11, 930.	1.5	23
6	GENITAL TRACT SCREENING FINDS WIDESPREAD INFECTION WITH MUSTELID GAMMAHERPESVIRUS 1 IN THE EUROPEAN BADGER (<i>MELES MELES</i>). <i>Journal of Wildlife Diseases</i> , 2018, 54, 133.	0.3	12
7	Cytomegaloviruses in a Community of Wild Nonhuman Primates in Taï National Park, Côte d'Ivoire. <i>Viruses</i> , 2018, 10, 11.	1.5	13
8	Survey for zoonotic pathogens in Norway rat populations from Europe. <i>Pest Management Science</i> , 2017, 73, 341-348.	1.7	37
9	Seroprevalence of Cytomegalovirus Infection Among a Rural Population of Côte d'Ivoire. <i>Viral Immunology</i> , 2017, 30, 54-57.	0.6	3
10	Biology, evolution, and medical importance of polyomaviruses: An update. <i>Infection, Genetics and Evolution</i> , 2017, 54, 18-38.	1.0	112
11	Detection and genome characterization of bovine polyomaviruses in beef muscle and ground beef samples from Germany. <i>International Journal of Food Microbiology</i> , 2017, 241, 168-172.	2.1	9
12	ICTV Virus Taxonomy Profile: Polyomaviridae. <i>Journal of General Virology</i> , 2017, 98, 1159-1160.	1.3	107
13	A Role of Sp1 Binding Motifs in Basal and Large T-Antigen-Induced Promoter Activities of Human Polyomavirus HPyV9 and Its Variant UF-1. <i>International Journal of Molecular Sciences</i> , 2017, 18, 2414.	1.8	5
14	Indigenous house mice dominate small mammal communities in northern Afghan military bases. <i>BMC Zoology</i> , 2017, 2, .	0.3	6
15	Large T antigen variants of human polyomaviruses 9 and 12 and seroreactivity against their N terminus. <i>Journal of General Virology</i> , 2017, 98, 704-714.	1.3	7
16	Novel polyomaviruses in shrews (Soricidae) with close similarity to human polyomavirus 12. <i>Journal of General Virology</i> , 2017, 98, 3060-3067.	1.3	20
17	Genome Sequences of Polyomaviruses from the Wild-Living Red Colobus (<i>Ptilocolobus badius</i>) and Western Chimpanzee (<i>Pan troglodytes verus</i>). <i>Genome Announcements</i> , 2016, 4, .	0.8	3
18	Genome Sequences of Murine Pneumotropic Virus (Polyomaviridae) Detected in Wild House Mice (<i>Mus musculus</i>). <i>Genome Announcements</i> , 2016, 4, .	0.8	3

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19	A taxonomy update for the family Polyomaviridae. Archives of Virology, 2016, 161, 1739-1750.	0.9	134
20	Assessing Host-Virus Codivergence for Close Relatives of Merkel Cell Polyomavirus Infecting African Great Apes. Journal of Virology, 2016, 90, 8531-8541.	1.5	21
21	Immunological methods for the detection of porcine lymphotropic herpesviruses (PLHV). Journal of Virological Methods, 2016, 233, 72-77.	1.0	22
22	Genome Sequence of Bovine Polyomavirus 1 Detected in a Salers Cow (Bos taurus) from Catalonia, Spain. Genome Announcements, 2016, 4, .	0.8	3
23	Characterization of the non-coding control region of polyomavirus KI isolated from nasopharyngeal samples from patients with respiratory symptoms or infection and from blood from healthy blood donors in Norway. Journal of General Virology, 2016, 97, 1647-1657.	1.3	10
24	Genome Sequence of a Central Chimpanzee-Associated Polyomavirus Related to BK and JC Polyomaviruses, Pan troglodytes <i>troglydytes</i> Polyomavirus 1. Genome Announcements, 2015, 3, .	0.8	5
25	Genetic identification of cytomegaloviruses in a rural population of CÔte d'Îvoire. Virology Journal, 2015, 12, 155.	1.4	7
26	Genome Sequences of a Rat Polyomavirus Related to Murine Polyomavirus, Rattus norvegicus Polyomavirus 1. Genome Announcements, 2015, 3, .	0.8	20
27	Adenovirus in Rural CÔte D'Ivoire: High Diversity and Cross-Species Detection. EcoHealth, 2015, 12, 441-452.	0.9	16
28	Identification and Functional Comparison of Seven-Transmembrane G-Protein-Coupled BILF1 Receptors in Recently Discovered Nonhuman Primate Lymphocryptoviruses. Journal of Virology, 2015, 89, 2253-2267.	1.5	19
29	Discovery of a polyomavirus in European badgers (Meles meles) and the evolution of host range in the family Polyomaviridae. Journal of General Virology, 2015, 96, 1411-1422.	1.3	12
30	Multiple Cross-Species Transmission Events of Human Adenoviruses (HAdV) during Hominine Evolution. Molecular Biology and Evolution, 2015, 32, 2072-2084.	3.5	54
31	Phylogenomic evidence for recombination of adenoviruses in wild gorillas. Journal of General Virology, 2015, 96, 3090-3098.	1.3	9
32	Early and late promoters of BK polyomavirus, Merkel cell polyomavirus, Trichodysplasia spinulosa-associated polyomavirus and human polyomavirus 12 are among the strongest of all known human polyomaviruses in 10 different cell lines. Journal of General Virology, 2015, 96, 2293-2303.	1.3	18
33	Are human polyomaviruses co-factors for cancers induced by other oncoviruses?. Reviews in Medical Virology, 2014, 24, 343-360.	3.9	26
34	Field immobilization for treatment of an unknown illness in a wild chimpanzee (Pan troglodytes) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 1. 2014, 55, 89-99.	0.7	9
35	High prevalence and diversity of species D adenoviruses (HAdV-D) in human populations of four Sub-Saharan countries. Virology Journal, 2014, 11, 25.	1.4	22
36	Crystallographic and Glycan Microarray Analysis of Human Polyomavirus 9 VP1 Identifies <i>N</i>-Glycolyl Neuraminic Acid as a Receptor Candidate. Journal of Virology, 2014, 88, 6100-6111.	1.5	36

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37	Genome analysis of non-human primate polyomaviruses. <i>Infection, Genetics and Evolution</i> , 2014, 26, 283-294.	1.0	21
38	Serological cross-reactivity between human polyomaviruses. <i>Reviews in Medical Virology</i> , 2013, 23, 250-264.	3.9	45
39	The novel human polyomaviruses <sc>HPyV</sc>6, 7, 9 and beyond. <i>Apmis</i> , 2013, 121, 783-795.	0.9	42
40	Novel Polyomaviruses of Nonhuman Primates: Genetic and Serological Predictors for the Existence of Multiple Unknown Polyomaviruses within the Human Population. <i>PLoS Pathogens</i> , 2013, 9, e1003429.	2.1	35
41	Absence of Frequent Herpesvirus Transmission in a Nonhuman Primate Predator-Prey System in the Wild. <i>Journal of Virology</i> , 2013, 87, 10651-10659.	1.5	23
42	Identification of a Novel Human Polyomavirus in Organs of the Gastrointestinal Tract. <i>PLoS ONE</i> , 2013, 8, e58021.	1.1	131
43	Seroprevalence of human polyomavirus 9 and cross-reactivity to African green monkey-derived lymphotropic polyomavirus. <i>Journal of General Virology</i> , 2012, 93, 698-705.	1.3	49
44	Genome analysis of the new human polyomaviruses. <i>Reviews in Medical Virology</i> , 2012, 22, 354-377.	3.9	48
45	A Novel Human Polyomavirus Closely Related to the African Green Monkey-Derived Lymphotropic Polyomavirus. <i>Journal of Virology</i> , 2011, 85, 4586-4590.	1.5	214
46	African Great Apes Are Naturally Infected with Polyomaviruses Closely Related to Merkel Cell Polyomavirus. <i>Journal of Virology</i> , 2011, 85, 916-924.	1.5	46
47	Novel Adenoviruses in Wild Primates: a High Level of Genetic Diversity and Evidence of Zoonotic Transmissions. <i>Journal of Virology</i> , 2011, 85, 10774-10784.	1.5	96
48	LETTER TO THE EDITOR: Reply to the Comment by Gessain et al. on Mugisha et al. <i>J Med Primatol</i> 2010; 39: 71-76. <i>Journal of Medical Primatology</i> , 2010, 39, 365-366.	0.3	0
49	Lymphocryptovirus phylogeny and the origins of Epstein-Barr virus. <i>Journal of General Virology</i> , 2010, 91, 630-642.	1.3	70
50	A novel adenovirus of Western lowland gorillas (<i>Gorilla gorilla gorilla</i>). <i>Virology Journal</i> , 2010, 7, 303.	1.4	26
51	Novel cytomegaloviruses in free-ranging and captive great apes: phylogenetic evidence for bidirectional horizontal transmission. <i>Journal of General Virology</i> , 2009, 90, 2386-2394.	1.3	42
52	The order Herpesvirales. <i>Archives of Virology</i> , 2009, 154, 171-177.	0.9	790
53	Identification of a novel betaherpesvirus in <i>Mus musculus</i> . <i>Virology Journal</i> , 2009, 6, 225.	1.4	4
54	High genotypic diversity and a novel variant of human cytomegalovirus revealed by combined UL33/UL55 genotyping with broad-range PCR. <i>Virology Journal</i> , 2009, 6, 210.	1.4	14

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55	Development of a recombinant antigen-based ELISA for the sero-detection of porcine lymphotropic herpesviruses. <i>Xenotransplantation</i> , 2008, 15, 357-364.	1.6	19
56	Novel Mammalian Herpesviruses and Lineages within the <i>Gammaherpesvirinae</i> : Cospeciation and Interspecies Transfer. <i>Journal of Virology</i> , 2008, 82, 3509-3516.	1.5	110
57	Identification of Novel Rodent Herpesviruses, Including the First Gammaherpesvirus of <i>Mus musculus</i> . <i>Journal of Virology</i> , 2007, 81, 8091-8100.	1.5	89
58	Discovery of herpesviruses in bats. <i>Journal of General Virology</i> , 2007, 88, 2651-2655.	1.3	85
59	Discovery of herpesviruses in multi-infected primates using locked nucleic acids (LNA) and a bigenic PCR approach. <i>Virology Journal</i> , 2007, 4, 84.	1.4	33
60	Response from authors RE: "Lack of association of herpesviruses with brain tumors". <i>Journal of NeuroVirology</i> , 2007, 13, 86-87.	1.0	2
61	Molecular interactions between porcine and human gammaherpesviruses: implications for xenografts?. <i>Xenotransplantation</i> , 2006, 13, 308-317.	1.6	20
62	Identification of novel alpha- and gammaherpesviruses from cutaneous and mucosal lesions of dolphins and whales. <i>Journal of Virological Methods</i> , 2006, 136, 261-266.	1.0	67
63	Novel herpesviruses of Suidae: indicators for a second genogroup of artiodactyl gammaherpesviruses. <i>Journal of General Virology</i> , 2004, 85, 857-862.	1.3	18
64	A novel porcine gammaherpesvirus. <i>Virology</i> , 2003, 308, 317-329.	1.1	68
65	Approaching virus safety in xenotransplantation: a search for unrecognized herpesviruses in pigs. <i>Xenotransplantation</i> , 2003, 10, 349-356.	1.6	38
66	Novel Simian Homologues of Epstein-Barr Virus. <i>Journal of Virology</i> , 2003, 77, 10695-10699.	1.5	56
67	Identification and Quantification of Ovine Gammaherpesvirus 2 DNA in Fresh and Stored Tissues of Pigs with Symptoms of Porcine Malignant Catarrhal Fever. <i>Journal of Clinical Microbiology</i> , 2003, 41, 900-904.	1.8	36
68	Sequence Analysis of the Genome of Porcine Lymphotropic Herpesvirus 1 and Gene Expression during Posttransplant Lymphoproliferative Disease of Pigs. <i>Virology</i> , 2002, 294, 383-393.	1.1	54
69	Detection and multigenic characterization of a novel gammaherpesvirus in goats. <i>Virus Research</i> , 2001, 75, 87-94.	1.1	76
70	Detection of new DNA polymerase genes of known and potentially novel herpesviruses by PCR with degenerate and deoxyinosine-substituted primers. <i>Virus Genes</i> , 1999, 18, 211-220.	0.7	168
71	Characterization of the DNA polymerase loci of the novel porcine lymphotropic herpesviruses 1 and 2 in domestic and feral pigs. <i>Journal of General Virology</i> , 1999, 80, 3199-3205.	1.3	49