Norbert Nowotny

List of Publications by Year
in descending order

253	12,728	23567	36028
papers	citations	58 h-index	97 g-index

263	263	263	10554
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Maintenance of neutralizing antibodies over ten months in convalescent SARSâ€CoVâ€2 afflicted patients. Transboundary and Emerging Diseases, 2022, 69, 1596-1605.	3.0	11
2	Diversity of West Nile and Usutu virus strains in mosquitoes at an international airport in Austria. Transboundary and Emerging Diseases, 2022, 69, 2096-2109.	3.0	10
3	MERSâ€CoV in sheep, goats, and cattle, United Arab Emirates, 2019: Virological and serological investigations reveal an accidental spillover from dromedaries. Transboundary and Emerging Diseases, 2022, 69, 3066-3072.	3.0	7
4	Crimeanâ€“Congo Hemorrhagic Fever Virus Past Infections Are Associated with Two Innate Immune Response Candidate Genes in Dromedaries. Cells, 2022, 11, 8.	4.1	5
5	Monitoring Urban Zoonotic Virus Activity: Are City Rats a Promising Surveillance Tool for Emerging Viruses?. Viruses, 2022, 14, 1516.	3.3	2
6	Host transcriptomic profiling of COVID-19 patients with mild, moderate, and severe clinical outcomes. Computational and Structural Biotechnology Journal, 2021, 19, 153-160.	4.1	69
7	New World camelids are sentinels for the presence of Borna disease virus. Transboundary and Emerging Diseases, 2021, , .	3.0	7
8	Genotypeâ€phenotype correlation identified a novel SARSâ€CoVâ€2 variant possibly linked to severe disease. Transboundary and Emerging Diseases, 2021, , .	3.0	8
9	Innate and Adaptive Immune Genes Associated with MERS-CoV Infection in Dromedaries. Cells, 2021, 10, 1291.	4.1	6
10	Capillaria hepatica (syn. Calodium hepaticum) as a Cause of Asymptomatic Liver Mass. American Journal of Tropical Medicine and Hygiene, 2021, , .	1.4	0
11	ICTV Virus Taxonomy Profile: Bornaviridae. Journal of General Virology, 2021, 102, .	2.9	24
12	2021 Taxonomic update of phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. Archives of Virology, 2021, 166, 3513-3566.	2.1	62
13	Single-cell transcriptome identifies FCGR3B upregulated subtype of alveolar macrophages in patients with critical COVID-19. IScience, 2021, 24, 103030.	4.1	13
14	Association of Dromedary Camels and Camel Ticks with Reassortant Crimean-Congo Hemorrhagic Fever Virus, United Arab Emirates. Emerging Infectious Diseases, 2021, 27, 2471-2474.	4.3	11
15	African Swine Fever, the forgotten pandemic. Transboundary and Emerging Diseases, 2021, 68, 2637-2639.	3.0	20
16	Crimean-Congo Hemorrhagic Fever Virus in Asia, Africa and Europe. Microorganisms, 2021, 9, 1907.	3.6	54
17	West Nile Virus and Tick-Borne Encephalitis Virus Are Endemic in Equids in Eastern Austria. Viruses, 2021, 13, 1873.	3.3	9
18	The transmission ecology of Tahyna orthobunyavirus in Austria as revealed by longitudinal mosquito sampling and blood meal analysis in floodplain habitats. Parasites and Vectors, 2021, 14, 561.	2.5	9

#	Article	IF	Citations
19	Meclizine Inhibits Pseudorabies Virus Replication by Interfering With Virus Entry and Release. Frontiers in Microbiology, 2021, 12, 795593.	3.5	4
20	Nitrogen Accumulation in Oyster (Crassostrea gigas) Slurry Exposed to Virucidal Cold Atmospheric Plasma Treatment. Life, 2021, 11, 1333.	2.4	4
21	Different dynamics of Usutu virus infections in Austria and Hungary, 2017â€“2018. Transboundary and Emerging Diseases, 2020, 67, 298-307.	3.0	25
22	The knowns and unknowns of West Nile virus in Europe: what did we learn from the 2018 outbreak?. Expert Review of Anti-Infective Therapy, 2020, 18, 145-154.	4.4	54
23	Emergence of West Nile virus lineage 2 in Europe: Characteristics of the first seven cases of West Nile neuroinvasive disease in horses in Austria. Transboundary and Emerging Diseases, 2020, 67, 1189-1197.	3.0	20
24	SARS-CoV-2 Whole Genome Amplification and Sequencing for Effective Population-Based Surveillance and Control of Viral Transmission. Clinical Chemistry, 2020, 66, 1450-1458.	3.2	31
25	Peptidomic Analysis of Skin Secretions of the Caribbean Frogs Leptodactylus insularum and Leptodactylus nesiotus (Leptodactylidae) Identifies an Ocellatin with Broad Spectrum Antimicrobial Activity. Antibiotics, 2020, 9, 718.	3.7	10
26	Multiple early introductions of SARS-CoV-2 into a global travel hub in the Middle East. Scientific Reports, 2020, 10, 17720.	3.3	28
27	2020 taxonomic update for phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. Archives of Virology, 2020, 165, 3023-3072.	2.1	184
28	Modelling West Nile Virus and Usutu Virus Pathogenicity in Human Neural Stem Cells. Viruses, 2020, 12, 882.	3.3	16
29	Autochthonous Transmission of West Nile Virus by a New Vector in Iran, Vector-Host Interaction Modeling and Virulence Gene Determinants. Viruses, 2020, 12, 1449.	3.3	8
30	SARS-CoV-2/COVID-19: Viral Genomics, Epidemiology, Vaccines, and Therapeutic Interventions. Viruses, 2020, 12, 526.	3.3	197
31	Haemosporidioses in wild Eurasian blackbirds (Turdus merula) and song thrushes (T. philomelos): an in situ hybridization study with emphasis on exo-erythrocytic parasite burden. Malaria Journal, 2020, 19, 69.	2.3	20
32	Crimean-Congo Hemorrhagic Fever Virus Endemicity in United Arab Emirates, 2019. Emerging Infectious Diseases, 2020, 26, 1019-1021.	4.3	24
33	Phylogenetic Analysis of Lednice Orthobunyavirus. Microorganisms, 2019, 7, 447.	3.6	5
34	Taxonomy of the order Mononegavirales: second update 2018. Archives of Virology, 2019, 164, 1233-1244.	2.1	70
35	West Nile and Usutu Virus Infections and Challenges to Blood Safety in the European Union. Emerging Infectious Diseases, 2019, 25, 1050-1057.	4.3	42
36	Taxonomy of the order Mononegavirales: update 2019. Archives of Virology, 2019, 164, 1967-1980.	2.1	224

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37	Specific detection and differentiation of tickâ€borne encephalitis and West Nile virus induced IgG antibodies in humans and horses. Transboundary and Emerging Diseases, 2019, 66, 1701-1708.	3.0	15
38	Targeted surveillance reveals native and invasive mosquito species infected with Usutu virus. Parasites and Vectors, 2019, 12, 46.	2.5	36
39	Mosquito biodiversity and mosquito-borne viruses in the United Arab Emirates. Parasites and Vectors, 2019, 12, 153.	2.5	13
40	No Evidence of Mosquito Involvement in the Transmission of Equine Hepacivirus (Flaviviridae) in an Epidemiological Survey of Austrian Horses. Viruses, 2019, 11, 1014.	3.3	11
41	Peptidomic analysis of the host-defense peptides in skin secretions of Rana graeca provides insight into phylogenetic relationships among Eurasian Rana species. Comparative Biochemistry and Physiology Part D: Genomics and Proteomics, 2019, 29, 228-234.	1.0	8
42	Strengthening the Interaction of the Virology Community with the International Committee on Taxonomy of Viruses (ICTV) by Linking Virus Names and Their Abbreviations to Virus Species. Systematic Biology, 2019, 68, 828-839.	5.6	11
43	Taxonomy of the order Mononegavirales: update 2018. Archives of Virology, 2018, 163, 2283-2294.	2.1	153
44	Serological surveillance for Tahyna virus (California encephalitis orthobunyavirus, Peribunyaviridae) neutralizing antibodies in wild ungulates in Austria, Hungary and Romania. Zoonoses and Public Health, 2018, 65, 459-463.	2.2	7
45	Integrated analysis of human-animal-vector surveillance: West Nile virus infections in Austria, 2015â€“2016. Emerging Microbes and Infections, 2018, 7, 1-15.	6.5	22
46	Increase in human West Nile and Usutu virus infections, Austria, 2018. Eurosurveillance, 2018, 23, .	7.0	69
47	Susceptibility and role as competent host of the red-legged partridge after infection with lineage 1 and 2 West Nile virus isolates of Mediterranean and Central European origin. Veterinary Microbiology, 2018, 222, 39-45.	1.9	11
48	Uranotaenia unguiculata Edwards, 1913 are attracted to sound, feed on amphibians, and are infected with multiple viruses. Parasites and Vectors, 2018, 11, 456.	2.5	19
49	Factors Affecting Transmission of Crimean - Congo Hemorrhagic Fever among Slaughterhouse Employees: A Serosurvey in Mashhad, Iran. Jundishapur Journal of Microbiology, 2018, 11, .	0.5	4
50	Spatial and Phylodynamic Survey on Crimean-Congo Hemorrhagic Fever Virus Strains in Northeast of Iran. Jundishapur Journal of Microbiology, 2018, 11, .	0.5	4
51	A novel HRM assay for the simultaneous detection and differentiation of eight poxviruses of medical and veterinary importance. Scientific Reports, 2017, 7, 42892.	3.3	43
52	Taxonomy of the order Mononegavirales: update 2017. Archives of Virology, 2017, 162, 2493-2504.	2.1	173
53	Crimean-Congo hemorrhagic fever cases in the North of Iran have three distinct origins. VirusDisease, 2017, 28, 50-53.	2.0	11
54	Co-circulation of Crimean-Congo Hemorrhagic Fever virus strains Asia 1 and 2 between the border of Iran and Pakistan. Heliyon, 2017, 3, e00439.	3.2	12

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55	Infections of horses and shrews with Bornaviruses in Upper Austria: a novel endemic area of Borna disease. Emerging Microbes and Infections, 2017, 6, 1-9.	6.5	31
56	A Serological Protein Microarray for Detection of Multiple Cross-Reactive Flavivirus Infections in Horses for Veterinary and Public Health Surveillance. Transboundary and Emerging Diseases, 2017, 64, 1801-1812.	3.0	26
57	Usutu virus, Austria and Hungary, 2010â€“2016. Emerging Microbes and Infections, 2017, 6, 1-7.	6.5	34
58	West Nile virus in overwintering mosquitoes, central Europe. Parasites and Vectors, 2017, 10, 452.	2.5	69
59	West Nile virus surveillance in Europe: moving towards an integrated animal-human-vector approach. Eurosurveillance, 2017, 22, .	7.0	71
60	Usutu virus infections among blood donors, Austria, July and August 2017 â€“ Raising awareness for diagnostic challenges. Eurosurveillance, 2017, 22, .	7.0	57
61	Vector prevalence and detection of Crimean-Congo haemorrhagic fever virus in Golestan Province, Iran. Journal of Vector Borne Diseases, 2017, 54, 353.	0.4	22
62	Spread of Aedes japonicus japonicus (Theobald, 1901) in Austria, 2011â€“2015, and first records of the subspecies for Hungary, 2012, and the principality of Liechtenstein, 2015. Parasites and Vectors, 2016, 9, 356.	2.5	36
63	Keratoconjunctivitis in a group of Icelandic horses with suspected Î³-herpesvirus involvement. Equine Veterinary Journal, 2016, 48, 427-429.	1.7	16
64	First record of the Asian bush mosquito, Aedes japonicus japonicus, in Italy: invasion from an established Austrian population. Parasites and Vectors, 2016, 9, 284.	2.5	37
65	Taxonomy of the order Mononegavirales: update 2016. Archives of Virology, 2016, 161, 2351-2360.	2.1	407
66	Rapid detection of European orthobunyaviruses by reverse transcription loop-mediated isothermal amplification assays. Journal of Virological Methods, 2016, 236, 252-257.	2.1	1
67	Phylogeny of tick-derived Crimean-Congo hemorrhagic fever virus strains in Iran. Ticks and Tick-borne Diseases, 2016, 7, 1216-1221.	2.7	22
68	Possibility and Challenges of Conversion of Current Virus Species Names to Linnaean Binomials. Systematic Biology, 2016, 66, syw096.	5.6	17
69	Genetic analysis of imported dengue virus strains by Iranian travelers. Asian Pacific Journal of Tropical Disease, 2016, 6, 850-853.	0.5	8
70	Chronic West Nile virus infection in kea (Nestor notabilis). Veterinary Microbiology, 2016, 183, 135-139.	1.9	12
71	Peptidomic analysis of the extensive array of host-defense peptides in skin secretions of the dodecaploid frog Xenopus ruwenzoriensis (Pipidae). Comparative Biochemistry and Physiology Part D: Genomics and Proteomics, 2016, 19, 18-24.	1.0	4
72	Genetic Diversity of Crimean Congo Hemorrhagic Fever Virus Strains from Iran. Journal of Arthropod-Borne Diseases, 2016, 10, 127-40.	0.9	12

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73	Co-circulation of Usutu virus and West Nile virus in a reed bed ecosystem. Parasites and Vectors, 2015, 8, 520.	2.5	36
74	West Nile Virus Positive Blood Donation and Subsequent Entomological Investigation, Austria, 2014. PLoS ONE, 2015, 10, e0126381.	2.5	24
75	A High-Performance Multiplex Immunoassay for Serodiagnosis of Flavivirus-Associated Neurological Diseases in Horses. BioMed Research International, 2015, 2015, 1-13.	1.9	56
76	First international external quality assessment of molecular diagnostics for Mers-CoV. Journal of Clinical Virology, 2015, 69, 81-85.	3.1	27
77	Taxonomic reorganization of the family Bornaviridae. Archives of Virology, 2015, 160, 621-632.	2.1	97
78	Host-defense and trefoil factor family peptides in skin secretions of the Mawa clawed frog Xenopus boumbaensis (Pipidae). Peptides, 2015, 72, 44-49.	2.4	5
79	Complex Epidemiology of a Zoonotic Disease in a Culturally Diverse Region: Phylogeography of Rabies Virus in the Middle East. PLoS Neglected Tropical Diseases, 2015, 9, e0003569.	3.0	42
80	Phylogenetic characterization of Central/Southern European lineage 2 West Nile virus: analysis of human outbreaks in Italy and Greece, 2013â€“2014. Clinical Microbiology and Infection, 2015, 21, 1122.e1-1122.e10.	6.0	49
81	Close Relationship between West Nile Virus from Turkey and Lineage 1 Strain from Central African Republic. Emerging Infectious Diseases, 2015, 21, 352-355.	4.3	23
82	Variation in interferon sensitivity and induction between Usutu and West Nile (lineages 1 and 2) viruses. Virology, 2015, 485, 189-198.	2.4	24
83	Detection of Diverse Novel Bat Astrovirus Sequences in the Czech Republic. Vector-Borne and Zoonotic Diseases, 2015, 15, 518-521.	1.5	28
84	The challenge of West Nile virus in Europe: knowledge gaps and research priorities. Eurosurveillance, 2015, 20, .	7.0	152
85	Serological evidence of West Nile virus infection in the horse population of northern Serbia. Journal of Infection in Developing Countries, 2014, 8, 914-918.	1.2	14
86	Putative New West Nile Virus Lineage inUranotaenia unguiculataMosquitoes, Austria, 2013. Emerging Infectious Diseases, 2014, 20, 2119-2122.	4.3	72
87	Investigations on California Serogroup Orthobunyaviruses in the Tyrols: First Description of Tahyna Virus in The Alps. Vector-Borne and Zoonotic Diseases, 2014, 14, 272-277.	1.5	15
88	Matrix-Mâ„¢ adjuvanted envelope protein vaccine protects against lethal lineage 1 and 2 West Nile virus infection in mice. Vaccine, 2014, 32, 800-808.	3.8	28
89	Host defense peptides from Lithobates forreri, Hylarana luctuosa, and Hylarana signata (Ranidae): Phylogenetic relationships inferred from primary structures of ranatuerin-2 and brevinin-2 peptides. Comparative Biochemistry and Physiology Part D: Genomics and Proteomics, 2014, 9, 49-57.	1.0	18
90	Arboviruses Pathogenic for Domestic and Wild Animals. Advances in Virus Research, 2014, 89, 201-275.	2.1	146

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91	Seroprevalence of Borrelia burgdorferi sensu lato and tick-borne encephalitis virus in zoo animal species in the Czech Republic. Ticks and Tick-borne Diseases, 2014, 5, 523-527.	2.7	19
92	Host-defense peptides from skin secretions of Fraser's clawed frog Xenopus fraseri (Pipidae): Further insight into the evolutionary history of the Xenopodinae. Comparative Biochemistry and Physiology Part D: Genomics and Proteomics, 2014, 12, 45-52.	1.0	5
93	Comparison of Complete Genome Sequences of Usutu Virus Strains Detected in Spain, Central Europe, and Africa. Vector-Borne and Zoonotic Diseases, 2014, 14, 324-329.	1.5	30
94	Full-length genome analysis of ÄŒalovo strains of Batai orthobunyavirus (Bunyamwera serogroup): Implications to taxonomy. Infection, Genetics and Evolution, 2014, 27, 96-104.	2.3	5
95	Prevalence and distribution patterns of seven different honeybee viruses in diseased colonies: a case study from Croatia. Apidologie, 2014, 45, 701-706.	2.0	20
96	Prevalence of asinine herpesvirus type 5 (AsHV-5) infection in clinically normal Lipizzaner horses. Veterinary Journal, 2014, 200, 200-203.	1.7	9
97	Usutu Virus in Blackbirds (Turdus merula), Czech Republic, 2011-2012. Transboundary and Emerging Diseases, 2014, 61, 273-276.	3.0	41
98	Experimental infection of house sparrows (Passer domesticus) with West Nile virus strains of lineages 1 and 2. Veterinary Microbiology, 2014, 172, 542-547.	1.9	23
99	Middle East respiratory syndrome coronavirus (MERS-CoV) in dromedary camels, Oman, 2013. Eurosurveillance, 2014, 19, 20781.	7.0	125
100	The Bicolored White-Toothed Shrew Crocidura leucodon (HERMANN 1780) Is an Indigenous Host of Mammalian Borna Disease Virus. PLoS ONE, 2014, 9, e93659.	2.5	57
101	The Complete Sequence of a West Nile Virus Lineage 2 Strain Detected in a Hyalomma marginatum marginatum Tick Collected from a Song Thrush (Turdus philomelos) in Eastern Romania in 2013 Revealed Closest Genetic Relationship to Strain Volgograd 2007. PLoS ONE, 2014, 9, e109905.	2.5	50
102	Authorsâ€™ reply: Middle East respiratory syndrome coronavirus (MERS-CoV) in dromedary camels: are dromedary camels a reservoir for MERS-CoV?. Eurosurveillance, 2014, 19, .	7.0	1
103	West Nile virus lineage 2 isolated from Culex modestus mosquitoes in the Czech Republic, 2013: expansion of the European WNV endemic area to the North?. Eurosurveillance, 2014, 19, 2-5.	7.0	43
104	Partial genetic characterization of Sedlec virus (Orthobunyavirus, Bunyaviridae). Infection, Genetics and Evolution, 2013, 19, 244-249.	2.3	4
105	Middle East respiratory syndrome coronavirus neutralising serum antibodies in dromedary camels: a comparative serological study. Lancet Infectious Diseases, The, 2013, 13, 859-866.	9.1	616
106	Detection of Plasmodium sp.-infested Anopheles hyrcanus (Pallas 1771) (Diptera: Culicidae) in Austria, 2012. Wiener Klinische Wochenschrift, 2013, 125, 139-143.	1.9	6
107	Molecular characterization of the African orthobunyavirus Ilesha virus. Infection, Genetics and Evolution, 2013, 20, 124-130.	2.3	9
108	Progestin treatment does not affect expression of cytokines, steroid receptors, oxytocin receptor, and cyclooxygenase 2 in fetal membranes and endometrium from pony mares at parturition. Theriogenology, 2013, 79, 59-68.	2.1	5

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109	Low Usutu virusseroprevalence in four zoological gardens in central Europe. BMC Veterinary Research, 2013, 9, 153.	1.9	27
110	Barkedji virus, a novel mosquito-borne flavivirus identified in Culex perexiguus mosquitoes, Israel, 2011. Journal of General Virology, 2013, 94, 2449-2457.	2.9	35
111	Explosive spread of a neuroinvasive lineage 2 West Nile virus in Central Europe, 2008/2009. Veterinary Microbiology, 2013, 165, 61-70.	1.9	192
112	Nyamiviridae: Proposal for a new family in the order Mononegavirales. Archives of Virology, 2013, 158, 2209-2226.	2.1	29
113	Pathogenesis of West Nile virus lineage 1 and 2 in experimentally infected large falcons. Veterinary Microbiology, 2013, 161, 263-273.	1.9	61
114	Detection of equid herpesviruses 2 and 5 in a herd of 266 Lipizzaners in association with ocular findings. Veterinary Microbiology, 2013, 164, 139-144.	1.9	33
115	Usutu Virus, Italy, 1996. Emerging Infectious Diseases, 2013, 19, 274-277.	4.3	186
116	Prevalence of linear keratopathy in a herd of Lipizzaners over an 18â€month period. Veterinary Record, 2013, 173, 192-192.	0.3	3
117	Usutu virus growth in human cell lines: induction of and sensitivity to type I and III interferons. Journal of General Virology, 2013, 94, 789-795.	2.9	16
118	Tick-borne Encephalitis Virus in Horses, Austria, 2011. Emerging Infectious Diseases, 2013, 19, 635-637.	4.3	55
119	Comparison of the Neuropathology Induced by Two West Nile Virus Strains. PLoS ONE, 2013, 8, e84473.	2.5	20
120	Flaviviruses in Europe: Complex Circulation Patterns and Their Consequences for the Diagnosis and Control of West Nile Disease. International Journal of Environmental Research and Public Health, 2013, 10, 6049-6083.	2.6	140
121	Clinical Course of Ophthalmic Findings and Potential Influence Factors of Herpesvirus Infections: 18 Month Follow-Up of a Closed Herd of Lipizzaners. PLoS ONE, 2013, 8, e79888.	2.5	10
122	Monitoring of West Nile Virus Infections in Germany. Zoonoses and Public Health, 2012, 59, 95-101.	2.2	33
123	Identification of Mixed Infections with Different Genotypes of Avian Bornaviruses in Psittacine Birds with Proventricular Dilatation Disease. Avian Diseases, 2012, 56, 414-417.	1.0	13
124	Cowpox virus isolate virulent in humans shows attenuated phenotype in mice. Research in Veterinary Science, 2012, 92, 333-337.	1.9	2
125	Immature and Mature Human Astrovirus: Structure, Conformational Changes, and Similarities to Hepatitis E Virus. Journal of Molecular Biology, 2012, 422, 650-658.	4.2	60
126	Detection and molecular characterization of Suid herpesvirus type 1 in Austrian wild boar and hunting dogs. Veterinary Microbiology, 2012, 157, 276-284.	1.9	47

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127	Cannabinoids lead to enhanced virulence of the smallpox vaccine (vaccinia) virus. Immunobiology, 2011, 216, 670-677.	1.9	17
128	Protection provided by a recombinant ALVACÂ®-WNV vaccine expressing the prM/E genes of a lineage 1 strain of WNV against a virulent challenge with a lineage 2 strain. Vaccine, 2011, 29, 4608-4612.	3.8	41
129	Characterization of antimicrobial peptides in skin secretions from discrete populations of Lithobates chiricahuensis (Ranidae) from central and southern Arizona. Peptides, 2011, 32, 664-669.	2.4	25
130	Emergence of canine distemper in Bavarian wildlife associated with a specific amino acid exchange in the haemagglutinin protein. Veterinary Journal, 2011, 187, 399-401.	1.7	47
131	Emergence and establishment of Usutu virus infection in wild and captive avian species in and around Zurich, Switzerlandâ€"Genomic and pathologic comparison to other central European outbreaks. Veterinary Microbiology, 2011, 148, 207-212.	1.9	105
132	Detection and molecular analysis of West Nile virus infections in birds of prey in the eastern part of Austria in 2008 and 2009. Veterinary Microbiology, 2011, 149, 358-366.	1.9	107
133	Genetic Characterization of West Nile Virus Lineage 2, Greece, 2010. Emerging Infectious Diseases, 2011, 17, 920-922.	4.3	172
134	Chasing Jenner's Vaccine: Revisiting Cowpox Virus Classification. PLoS ONE, 2011, 6, e23086.	2.5	95
135	Iflavirus. , 2011, , 2011-2017.		0
136	Partial sequencing of recent Portuguese myxoma virus field isolates exhibits a high degree of genetic stability. Veterinary Microbiology, 2010, 140, 161-166.	1.9	10
137	Usutu virus in wild birds in northern Italy. Veterinary Microbiology, 2010, 141, 159-163.	1.9	90
138	Zoonotic mosquito-borne flaviviruses: Worldwide presence of agents with proven pathogenicity and potential candidates of future emerging diseases. Veterinary Microbiology, 2010, 140, 271-280.	1.9	158
139	Detection of Hepatitis E virus in samples of animal origin collected in Hungary. Veterinary Microbiology, 2010, 143, 106-116.	1.9	75
140	Localization of avian bornavirus RNA by in situ hybridization in tissues of psittacine birds with proventricular dilatation disease. Veterinary Microbiology, 2010, 145, 9-16.	1.9	13
141	Anchored Pan Dengue RTâ€PCR and Fast Sanger sequencing for detection of Dengue RNA in human serum. Journal of Medical Virology, 2010, 82, 1701-1710.	5.0	4
142	Natural zoonotic infections of two marmosets and one domestic rabbit with herpes simplex virus type 1 did not reveal a correlation with a certain gG-, gI- or gE genotype. Clinical Microbiology and Infection, 2010, 16, 1669-1672.	6.0	10
143	Distribution of Borna Disease Virus Antigen and RNA in Tissues of Naturally Infected Bicolored White-Toothed Shrews, Crocidura leucodon, Supporting Their Role as Reservoir Host Species. Veterinary Pathology, 2010, 47, 236-244.	1.7	47
144	Mosquito (Diptera: Culicidae) Surveillance for Arboviruses in an Area Endemic for West Nile (Lineage) Tj ETQq0 0 0 rgBT /Overlock 10 Tf	1.8	43

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145	Antimicrobial peptides from the skin secretions of the South-East Asian frog Hylarana erythraea (Ranidae). Peptides, 2010, 31, 548-554.	2.4	31
146	Diazepam leads to enhanced severity of orthopoxvirus infection and immune suppression. Vaccine, 2010, 28, 6152-6158.	3.8	28
147	West Nile Virus Monitoring of Migratory and Resident Birds in Germany. Vector-Borne and Zoonotic Diseases, 2010, 10, 639-647.	1.5	61
148	Novel Avian Bornavirus in a Nonpsittacine Species (Canary; Serinus canaria) with Enteric Ganglioneuritis and Encephalitis. Journal of Virology, 2009, 83, 11367-11371.	3.4	65
149	Avian Bornaviruses in Psittacine Birds from Europe and Australia with Proventricular Dilatation Disease. Emerging Infectious Diseases, 2009, 15, 1453-1459.	4.3	87
150	Evolution of rabbit haemorrhagic disease virus (RHDV) in the European rabbit (Oryctolagus) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 542 T	1.9	50
151	Genetic analysis and phylogenetic comparison of Black queen cell virus genotypes. Veterinary Microbiology, 2009, 139, 227-234.	1.9	28
152	Altered gene expression may underlie prolonged duration of the QT interval and ventricular action potential in streptozotocin-induced diabetic rat heart. Molecular and Cellular Biochemistry, 2009, 328, 57-65.	3.1	20
153	Antimicrobial peptides from the skins of North American frogs. Biochimica Et Biophysica Acta - Biomembranes, 2009, 1788, 1556-1563.	2.6	107
154	Antimicrobial peptides from the skin secretions of the New World frogs Lithobates capito and Lithobates warszewitschii (Ranidae). Peptides, 2009, 30, 1775-1781.	2.4	20
155	Ecthyma contagiosum (orf) â€“ report of a human case from the United Arab Emirates and review of the literature. Journal of Cutaneous Pathology, 2008, 35, 603-607.	1.3	40
156	Serological evidence of continuing high Usutu virus (Flaviviridae) activity and establishment of herd immunity in wild birds in Austria. Veterinary Microbiology, 2008, 127, 237-248.	1.9	68
157	Explaining Usutu virus dynamics in Austria: Model development and calibration. Preventive Veterinary Medicine, 2008, 85, 166-186.	1.9	68
158	Prevalence of pathogenic bee viruses in Hungarian apiaries: Situation before joining the European Union. Journal of Invertebrate Pathology, 2008, 98, 235-238.	3.2	46
159	Characterization of antimicrobial peptides from the skin secretions of the Malaysian frogs, Odorrana hosii and Hylarana picturata (Anura:Ranidae). Toxicon, 2008, 52, 465-473.	1.6	49
160	Influence of different semen extenders and seminal plasma on PMN migration and on expression of IL-1Î², IL-6, TNF-Î± and COX-2 mRNA in the equine endometrium. Theriogenology, 2008, 70, 843-851.	2.1	66
161	Embryo transfer induces a subclinical endometritis in recipient mares which can be prevented by treatment with non-steroid anti-inflammatory drugs. Theriogenology, 2008, 70, 1147-1158.	2.1	43
162	Uterine involution and endometrial function in postpartum pony mares. American Journal of Veterinary Research, 2008, 69, 1525-1534.	0.6	27

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