

Susan A Nadin-Davis

List of Publications by Year in descending order

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Version: 2024-02-01

42
papers

1,401
citations

361413

20
h-index

330143

37
g-index

42
all docs

42
docs citations

42
times ranked

1050
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Relationships between fox populations and rabies virus spread in northern Canada. PLoS ONE, 2021, 16, e0246508. | 2.5 | 18 |
| 2 | Molecular epidemiology. , 2020, , 143-193. | | 6 |
| 3 | A long-distance translocation initiated an outbreak of raccoon rabies in Hamilton, Ontario, Canada. PLoS Neglected Tropical Diseases, 2020, 14, e0008113. | 3.0 | 9 |
| 4 | Population- and Variant-Based Genome Analyses of Viruses from Vaccine-Derived Rabies Cases Demonstrate Product Specific Clusters and Unique Patterns. Viruses, 2020, 12, 115. | 3.3 | 8 |
| 5 | Rabies virus infection in a 21-year-old male presenting with ascending paralysis after a bat scratch. Jammi, 2020, 5, 201-208. | 0.5 | 1 |
| 6 | ONRABÂ® oral rabies vaccine is shed from, but does not persist in, captive mammals. Vaccine, 2019, 37, 4310-4317. | 3.8 | 4 |
| 7 | Rapid identification of the raccoon rabies virus variant using a real-time reverse-transcriptase polymerase chain reaction. Journal of Virological Methods, 2019, 273, 113713. | 2.1 | 4 |
| 8 | Origins of the arctic fox variant rabies viruses responsible for recent cases of the disease in southern Ontario. PLoS Neglected Tropical Diseases, 2019, 13, e0007699. | 3.0 | 19 |
| 9 | In-depth genome analyses of viruses from vaccine-derived rabies cases and corresponding live-attenuated oral rabies vaccines. Vaccine, 2019, 37, 4758-4765. | 3.8 | 14 |
| 10 | Geography but not alternative host species explain the spread of raccoon rabies virus in Vermont. Epidemiology and Infection, 2018, 146, 1977-1986. | 2.1 | 10 |
| 11 | Application of high-throughput sequencing to whole rabies viral genome characterisation and its use for phylogenetic re-evaluation of a raccoon strain incursion into the province of Ontario. Virus Research, 2017, 232, 123-133. | 2.2 | 23 |
| 12 | Rabies. Nature Reviews Disease Primers, 2017, 3, 17091. | 30.5 | 239 |
| 13 | Processes Underlying Rabies Virus Incursions across USâ€“Canada Border as Revealed by Whole-Genome Phylogeography. Emerging Infectious Diseases, 2017, 23, 1454-1461. | 4.3 | 34 |
| 14 | The phylogeography of Myotis bat-associated rabies viruses across Canada. PLoS Neglected Tropical Diseases, 2017, 11, e0005541. | 3.0 | 19 |
| 15 | Characterization of epitopes on the rabies virus glycoprotein by selection and analysis of escape mutants. Virus Research, 2016, 220, 161-171. | 2.2 | 7 |
| 16 | Spatio-temporal Analysis of the Genetic Diversity of Arctic Rabies Viruses and Their Reservoir Hosts in Greenland. PLoS Neglected Tropical Diseases, 2016, 10, e0004779. | 3.0 | 34 |
| 17 | Comparison of culture versus quantitative real-time polymerase chain reaction for the detection of Taylorella equigenitalis in field samples from naturally infected horses in Canada and Germany. Canadian Journal of Veterinary Research, 2015, 79, 161-9. | 0.2 | 2 |
| 18 | Molecular Epidemiology. , 2013, , 123-177. | | 5 |

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|----|--|-----|-----------|
| 19 | Genetic characterization and phylogenetic analysis of skunk-associated rabies viruses in North America with special emphasis on the central plains. <i>Virus Research</i> , 2013, 174, 27-36. | 2.2 | 28 |
| 20 | Microarray for Identification of the Chiropteran Host Species of Rabies Virus in Canada. <i>Microarrays</i> (Basel, Switzerland), 2013, 2, 153-169. | 1.4 | 5 |
| 21 | Recent emergence of the Arctic rabies virus lineage. <i>Virus Research</i> , 2012, 163, 352-362. | 2.2 | 33 |
| 22 | DNA Barcoding Facilitates Bat Species Identification for Improved Surveillance of Bat-associated Rabies across Canada. <i>The Open Zoology Journal</i> , 2012, 5, 27-37. | 0.4 | 12 |
| 23 | Genetic Tracking of the Raccoon Variant of Rabies Virus in Eastern North America. <i>Epidemics</i> , 2011, 3, 76-87. | 3.0 | 21 |
| 24 | Molecular Phylogenetics of the Lyssaviruses—Insights from a Coalescent Approach. <i>Advances in Virus Research</i> , 2011, 79, 203-238. | 2.1 | 19 |
| 25 | Generation and characterization of a panel of anti-phosphoprotein monoclonal antibodies directed against Mokola virus. <i>Virus Research</i> , 2011, 160, 238-245. | 2.2 | 3 |
| 26 | Spatial and temporal dynamics of rabies virus variants in big brown bat populations across Canada: footprints of an emerging zoonosis. <i>Molecular Ecology</i> , 2010, 19, 2120-2136. | 3.9 | 36 |
| 27 | Characterization of a panel of anti-phosphoprotein monoclonal antibodies generated against the raccoon strain of rabies virus. <i>Virus Research</i> , 2010, 152, 126-136. | 2.2 | 10 |
| 28 | Development of real-time reverse transcriptase polymerase chain reaction methods for human rabies diagnosis. <i>Journal of Medical Virology</i> , 2009, 81, 1484-1497. | 5.0 | 70 |
| 29 | ERA VACCINE-DERIVED CASES OF RABIES IN WILDLIFE AND DOMESTIC ANIMALS IN ONTARIO, CANADA, 1989–2004. <i>Journal of Wildlife Diseases</i> , 2008, 44, 71-85. | 0.8 | 75 |
| 30 | ORIGINS OF THE RABIES VIRUSES ASSOCIATED WITH AN OUTBREAK IN NEWFOUNDLAND DURING 2002–2003. <i>Journal of Wildlife Diseases</i> , 2008, 44, 86-98. | 0.8 | 10 |
| 31 | Laboratory diagnosis of rabies in Canada for calendar year 2006. <i>Canadian Veterinary Journal</i> , 2008, 49, 359-61. | 0.0 | 8 |
| 32 | Emergence of Arctic-like Rabies Lineage in India. <i>Emerging Infectious Diseases</i> , 2007, 13, 111-116. | 4.3 | 53 |
| 33 | A molecular epidemiological analysis of the incursion of the raccoon strain of rabies virus into Canada. <i>Epidemiology and Infection</i> , 2006, 134, 534-547. | 2.1 | 22 |
| 34 | Persistence of genetic variants of the arctic fox strain of Rabies virus in southern Ontario. <i>Canadian Journal of Veterinary Research</i> , 2006, 70, 11-9. | 1.1 | 26 |
| 35 | Lyssavirus P Gene Characterisation Provides Insights into the Phylogeny of the Genus and Identifies Structural Similarities and Diversity within the Encoded Phosphoprotein. <i>Virology</i> , 2002, 298, 286-305. | 2.4 | 85 |
| 36 | Antigenic and genetic divergence of rabies viruses from bat species indigenous to Canada. <i>Virus Research</i> , 2001, 74, 139-156. | 2.2 | 117 |

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|----|--|-----|-----------|
| 37 | Phylogeographic patterns exhibited by Ontario rabies virus variants. <i>Epidemiology and Infection</i> , 1999, 123, 325-336. | 2.1 | 48 |
| 38 | Polymerase chain reaction protocols for rabies virus discrimination. <i>Journal of Virological Methods</i> , 1998, 75, 1-8. | 2.1 | 79 |
| 39 | Polymorphism of rabies viruses within the phosphoprotein and matrix protein genes. <i>Archives of Virology</i> , 1997, 142, 979-992. | 2.1 | 29 |
| 40 | Bovine herpesvirus 1 isolates contain variable copy numbers of GR-rich tandem repeats in the gl non-coding regions of their genomes. <i>Virus Genes</i> , 1996, 13, 263-268. | 1.6 | 5 |
| 41 | A molecular epidemiological study of rabies virus in central Ontario and western Quebec. <i>Journal of General Virology</i> , 1994, 75, 2575-2583. | 2.9 | 81 |
| 42 | Identification of Regional Variants of the Rabies Virus within the Canadian Province of Ontario. <i>Journal of General Virology</i> , 1993, 74, 829-837. | 2.9 | 70 |