

# Nicholas J Davis-Poynter

## List of Publications by Year in descending order

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67  
papers

3,841  
citations

136940

32  
h-index

128286

60  
g-index

68  
all docs

68  
docs citations

68  
times ranked

2801  
citing authors

#	ARTICLE	IF	CITATIONS
1	Construction and properties of a mutant of herpes simplex virus type 1 with glycoprotein H coding sequences deleted. <i>Journal of Virology</i> , 1992, 66, 341-348.	3.4	411
2	Inhibition of natural killer cells by a cytomegalovirus MHC class I homologue in vivo. <i>Nature</i> , 1997, 386, 510-514.	27.8	290
3	Analysis of Equid Herpesvirus 1 Strain Variation Reveals a Point Mutation of the DNA Polymerase Strongly Associated with Neuropathogenic versus Nonneuropathogenic Disease Outbreaks. <i>Journal of Virology</i> , 2006, 80, 4047-4060.	3.4	244
4	Equine Herpesvirus-1 Consensus Statement. <i>Journal of Veterinary Internal Medicine</i> , 2009, 23, 450-461.	1.6	241
5	Identification and characterization of a G protein-coupled receptor homolog encoded by murine cytomegalovirus. <i>Journal of Virology</i> , 1997, 71, 1521-1529.	3.4	206
6	A Point Mutation in a Herpesvirus Polymerase Determines Neuropathogenicity. <i>PLoS Pathogens</i> , 2007, 3, e160.	4.7	176
7	The Murine Cytomegalovirus Chemokine Homolog, m131/129, Is a Determinant of Viral Pathogenicity. <i>Journal of Virology</i> , 1999, 73, 6800-6809.	3.4	123
8	Analysis of the contributions of herpes simplex virus type 1 membrane proteins to the induction of cell-cell fusion. <i>Journal of Virology</i> , 1994, 68, 7586-7590.	3.4	114
9	Glycoprotein G isoforms from some alphaherpesviruses function as broad-spectrum chemokine binding proteins. <i>EMBO Journal</i> , 2003, 22, 833-846.	7.8	111
10	Sequence Variation of the SeM Gene of <i>Streptococcus equi</i> Allows Discrimination of the Source of Strangles Outbreaks. <i>Journal of Clinical Microbiology</i> , 2006, 44, 480-486.	3.9	95
11	Distinct Brainstem and Forebrain Circuits Receiving Tracheal Sensory Neuron Inputs Revealed Using a Novel Conditional Anterograde Transsynaptic Viral Tracing System. <i>Journal of Neuroscience</i> , 2015, 35, 7041-7055.	3.6	94
12	Transneuronal tracing of airways-related sensory circuitry using herpes simplex virus 1, strain H129. <i>Neuroscience</i> , 2012, 207, 148-166.	2.3	77
13	Mutations in the cytoplasmic tail of herpes simplex virus glycoprotein H suppress cell fusion by a syncytial strain. <i>Journal of Virology</i> , 1994, 68, 6985-6993.	3.4	77
14	M144, a Murine Cytomegalovirus (Mcmv)-Encoded Major Histocompatibility Complex Class I Homologue, Confers Tumor Resistance to Natural Killer Cell-Mediated Rejection. <i>Journal of Experimental Medicine</i> , 1999, 190, 435-444.	8.5	74
15	Evidence for multiple sensory circuits in the brain arising from the respiratory system: an anterograde viral tract tracing study in rodents. <i>Brain Structure and Function</i> , 2015, 220, 3683-3699.	2.3	66
16	Anterograde neuronal circuit tracing using a genetically modified herpes simplex virus expressing EGFP. <i>Journal of Neuroscience Methods</i> , 2012, 209, 158-167.	2.5	62
17	Murine Cytomegalovirus Exploits Olfaction To Enter New Hosts. <i>MBio</i> , 2016, 7, e00251-16.	4.1	62
18	Masters of deception: A review of herpesvirus immune evasion strategies. <i>Immunology and Cell Biology</i> , 1996, 74, 513-522.	2.3	61

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19	Lymph Node Macrophages Restrict Murine Cytomegalovirus Dissemination. <i>Journal of Virology</i> , 2015, 89, 7147-7158.	3.4	61
20	The M33 Chemokine Receptor Homolog of Murine Cytomegalovirus Exhibits a Differential Tissue-Specific Role during In Vivo Replication and Latency. <i>Journal of Virology</i> , 2009, 83, 7590-7601.	3.4	57
21	Mutation of the Maturase Lipoprotein Attenuates the Virulence of <i>Streptococcus equi</i> to a Greater Extent than Does Loss of General Lipoprotein Lipidation. <i>Infection and Immunity</i> , 2006, 74, 6907-6919.	2.2	55
22	A novel streptococcal integrative conjugative element involved in iron acquisition. <i>Molecular Microbiology</i> , 2008, 70, 1274-1292.	2.5	55
23	Murine Cytomegalovirus Spreads by Dendritic Cell Recirculation. <i>MBio</i> , 2017, 8, .	4.1	52
24	Evidence supporting the inclusion of strains from each of the two co-circulating lineages of H3N8 equine influenza virus in vaccines. <i>Vaccine</i> , 2004, 22, 4101-4109.	3.8	50
25	Functional Analysis of the Murine Cytomegalovirus Chemokine Receptor Homologue M33: Ablation of Constitutive Signaling Is Associated with an Attenuated Phenotype In Vivo. <i>Journal of Virology</i> , 2008, 82, 1884-1898.	3.4	49
26	Inhibition of NK Cells by Murine CMV-Encoded Class I MHC Homologue m144. <i>Cellular Immunology</i> , 1999, 191, 145-151.	3.0	42
27	Cytomegalovirus evasion of natural killer cell responses. <i>Immunological Reviews</i> , 1999, 168, 187-197.	6.0	41
28	Partial Functional Complementation between Human and Mouse Cytomegalovirus Chemokine Receptor Homologues. <i>Journal of Virology</i> , 2011, 85, 6091-6095.	3.4	40
29	GAG mimetic functionalised solid and mesoporous silica nanoparticles as viral entry inhibitors of herpes simplex type 1 and type 2 viruses. <i>Nanoscale</i> , 2016, 8, 16192-16196.	5.6	40
30	Comparison of hamster and pony challenge models for evaluation of effect of antigenic drift on cross protection afforded by equine influenza vaccines. <i>Equine Veterinary Journal</i> , 2010, 35, 458-462.	1.7	38
31	Induction of Antibody Responses to African Horse Sickness Virus (AHSV) in Ponies after Vaccination with Recombinant Modified Vaccinia Ankara (MVA). <i>PLoS ONE</i> , 2009, 4, e5997.	2.5	37
32	Equine herpesvirus-1 abortion: atypical cases with lesions largely or wholly restricted to the placenta. <i>Equine Veterinary Journal</i> , 2010, 36, 79-82.	1.7	37
33	Development and evaluation of ELISA procedures to detect antibodies against the major envelope protein (GL) of equine arteritis virus. <i>Journal of Virological Methods</i> , 2000, 90, 167-183.	2.1	35
34	In vitro characterisation of high and low virulence isolates of equine herpesvirus-1 and -4. <i>Research in Veterinary Science</i> , 2003, 75, 83-86.	1.9	33
35	From sabotage to camouflage: viral evasion of cytotoxic T lymphocyte and natural killer cell-mediated immunity. <i>Seminars in Cell and Developmental Biology</i> , 1998, 9, 369-378.	5.0	30
36	Generation of a Candidate Live Marker Vaccine for Equine Arteritis Virus by Deletion of the Major Virus Neutralization Domain. <i>Journal of Virology</i> , 2003, 77, 8470-8480.	3.4	30

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37	Alveolar Macrophages Are a Prominent but Nonessential Target for Murine Cytomegalovirus Infecting the Lungs. <i>Journal of Virology</i> , 2016, 90, 2756-2766.	3.4	29
38	The Equine Herpesvirus 2 E1 Open Reading Frame Encodes a Functional Chemokine Receptor. <i>Journal of Virology</i> , 1999, 73, 9843-9848.	3.4	29
39	Report of the equine herpesvirus-1 Havermeier Workshop, San Gimignano, Tuscany, June 2004. <i>Veterinary Immunology and Immunopathology</i> , 2006, 111, 3-13.	1.2	28
40	Frequency and phenotype of EHV-1 specific, IFN- $\gamma$ synthesising lymphocytes in ponies: The effects of age, pregnancy and infection. <i>Developmental and Comparative Immunology</i> , 2007, 31, 202-214.	2.3	28
41	Evaluation of microporous polycaprolactone matrices for controlled delivery of antiviral microbicides to the female genital tract. <i>Journal of Materials Science: Materials in Medicine</i> , 2013, 24, 2719-2727.	3.6	27
42	Sequence analysis of the equid herpesvirus 2 chemokine receptor homologues E1, ORF74 and E6 demonstrates high sequence divergence between field isolates. <i>Journal of General Virology</i> , 2007, 88, 2450-2462.	2.9	27
43	Murine cytomegalovirus degrades MHC class II to colonize the salivary glands. <i>PLoS Pathogens</i> , 2018, 14, e1006905.	4.7	24
44	Type 1 Interferons and NK Cells Limit Murine Cytomegalovirus Escape from the Lymph Node Subcapsular Sinus. <i>PLoS Pathogens</i> , 2016, 12, e1006069.	4.7	23
45	Detection of equine arteritis virus (EAV)-specific cytotoxic CD8+ T lymphocyte precursors from EAV-infected ponies. <i>Journal of General Virology</i> , 2003, 84, 2745-2753.	2.9	22
46	Identification of Common Mechanisms by Which Human and Mouse Cytomegalovirus Seven-Transmembrane Receptor Homologues Contribute to <i>In Vivo</i> Phenotypes in a Mouse Model. <i>Journal of Virology</i> , 2013, 87, 4112-4117.	3.4	21
47	Cytomegalovirus MHC class I homologues and natural killer cells: an overview. <i>Microbes and Infection</i> , 2000, 2, 521-532.	1.9	20
48	Evaluation of a prototype sub-unit vaccine against equine arteritis virus comprising the entire ectodomain of the virus large envelope glycoprotein (GL): induction of virus-neutralizing antibody and assessment of protection in ponies. <i>Journal of General Virology</i> , 2001, 82, 2425-2435.	2.9	20
49	Structure-Activity Relationships of GAG Mimetic-Functionalized Mesoporous Silica Nanoparticles and Evaluation of Acyclovir-Loaded Antiviral Nanoparticles with Dual Mechanisms of Action. <i>ACS Omega</i> , 2018, 3, 1689-1699.	3.5	17
50	A molecular approach to the identification of cytotoxic T-lymphocyte epitopes within equine herpesvirus 1. <i>Journal of General Virology</i> , 2006, 87, 2507-2515.	2.9	16
51	Human cytomegalovirus US28 allows dendritic cell exit from lymph nodes. <i>Journal of General Virology</i> , 2018, 99, 1509-1514.	2.9	16
52	Analysis of the subcellular trafficking properties of murine cytomegalovirus M78, a 7 transmembrane receptor homologue. <i>Journal of General Virology</i> , 2009, 90, 59-68.	2.9	14
53	Structural Diversity in Conserved Regions Like the DRY-Motif among Viral 7TM Receptors: A Consequence of Evolutionary Pressure?. <i>Advances in Virology</i> , 2012, 2012, 1-15.	1.1	14
54	Murine Cytomegalovirus Homologues of Cellular Immunomodulatory Genes. <i>Intervirology</i> , 1999, 42, 331-341.	2.8	11

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55	Human and Murine Cytomegalovirus Evasion of Cytotoxic T Lymphocyte and Natural Killer Cell-Mediated Immune Responses. <i>Seminars in Virology</i> , 1998, 8, 369-376.	3.9	10
56	Utilisation of bacteriophage display libraries to identify peptide sequences recognised by Equine herpesvirus type 1 specific equine sera. <i>Journal of Virological Methods</i> , 2000, 88, 89-104.	2.1	10
57	Virus-Encoded 7 Transmembrane Receptors. <i>Progress in Molecular Biology and Translational Science</i> , 2015, 129, 353-393.	1.7	10
58	Murine Cytomegalovirus Glycoprotein O Promotes Epithelial Cell Infection <i>In Vivo</i> . <i>Journal of Virology</i> , 2019, 93, .	3.4	10
59	Investigation and management of an outbreak of abortion related to equine herpesvirus type 1 in unvaccinated ponies. <i>Veterinary Record</i> , 2007, 160, 378-380.	0.3	9
60	Luciferase-tagged wild-type and tropism-deficient mouse cytomegaloviruses reveal early dynamics of host colonization following peripheral challenge. <i>Journal of General Virology</i> , 2016, 97, 3379-3391.	2.9	9
61	Use of polarised equine endothelial cell cultures and an in vitro thrombosis model for potential characterisation of EHV-1 strain variation. <i>Veterinary Microbiology</i> , 2006, 113, 243-249.	1.9	7
62	Synergistic activity of tenofovir and nevirapine combinations released from polycaprolactone matrices for potential enhanced prevention of HIV infection through the vaginal route. <i>European Journal of Pharmaceutics and Biopharmaceutics</i> , 2014, 88, 406-414.	4.3	7
63	Analysis of ORFs 2b, 3, 4, and partial ORF5 of sequential isolates of equine arteritis virus shows genetic variation following experimental infection of horses. <i>Veterinary Microbiology</i> , 2008, 129, 262-268.	1.9	4
64	The Cytoplasmic C-Tail of the Mouse Cytomegalovirus 7 Transmembrane Receptor Homologue, M78, Regulates Endocytosis of the Receptor and Modulates Virus Replication in Different Cell Types. <i>PLoS ONE</i> , 2016, 11, e0165066.	2.5	4
65	Constitutive Signaling by the Human Cytomegalovirus G Protein Coupled Receptor Homologs US28 and UL33 Enables Trophoblast Migration In Vitro. <i>Viruses</i> , 2022, 14, 391.	3.3	4
66	The Mouse Cytomegalovirus G Protein-Coupled Receptor Homolog, M33, Coordinates Key Features of <i>In Vivo</i> Infection via Distinct Components of Its Signaling Repertoire. <i>Journal of Virology</i> , 2022, 96, JVI0186721.	3.4	3
67	Introduction: Virus stealth strategies. <i>Seminars in Cell and Developmental Biology</i> , 1998, 9, 319.	5.0	0