

# Jukka Niemimaa

## List of Publications by Year in descending order

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33  
papers

1,124  
citations

394421

19  
h-index

434195

31  
g-index

34  
all docs

34  
docs citations

34  
times ranked

1230  
citing authors

#	ARTICLE	IF	CITATIONS
1	Geographical Distribution of Ljungan Virus in Small Mammals in Europe. <i>Vector-Borne and Zoonotic Diseases</i> , 2020, 20, 692-702.	1.5	5
2	Microevolution of bank voles ( <i>Myodes glareolus</i> ) at neutral and immune-related genes during multiannual dynamic cycles: Consequences for Puumala hantavirus epidemiology. <i>Infection, Genetics and Evolution</i> , 2017, 49, 318-329.	2.3	37
3	DISTRIBUTION AND SEASONAL VARIATION OF LJUNGAN VIRUS IN BANK VOLES ( <i>MYODES GLAREOLUS</i> ) IN FENNOSCANDIA. <i>Journal of Wildlife Diseases</i> , 2017, 53, 552.	0.8	5
4	Run to the hills: gene flow among mountain areas leads to low genetic differentiation in the Norwegian lemming. <i>Biological Journal of the Linnean Society</i> , 2017, 121, 1-14.	1.6	10
5	Harmonizing circumpolar monitoring of Arctic fox: benefits, opportunities, challenges and recommendations. <i>Polar Research</i> , 2017, 36, 2.	1.6	35
6	Homage to Hersteinsson and Macdonald: climate warming and resource subsidies cause red fox range expansion and Arctic fox decline. <i>Polar Research</i> , 2017, 36, 3.	1.6	72
7	Metagenomic Evaluation of Bacteria from Voles. <i>Vector-Borne and Zoonotic Diseases</i> , 2017, 17, 123-133.	1.5	9
8	Temporal dynamics of Puumala hantavirus infection in cyclic populations of bank voles. <i>Scientific Reports</i> , 2016, 6, 21323.	3.3	38
9	Severe Ocular Cowpox in a Human, Finland. <i>Emerging Infectious Diseases</i> , 2015, 21, 2261-2263.	4.3	31
10	Monitoring biothreat agents ( <i>Francisella tularensis</i> , <i>Bacillus anthracis</i> and <i>Yersinia pestis</i> ) with a portable real-time PCR instrument. <i>Journal of Microbiological Methods</i> , 2015, 115, 89-93.	1.6	19
11	Life-long shedding of Puumala hantavirus in wild bank voles ( <i>Myodes glareolus</i> ). <i>Journal of General Virology</i> , 2015, 96, 1238-1247.	2.9	77
12	Detection of <i>Francisella tularensis</i> in Voles in Finland. <i>Vector-Borne and Zoonotic Diseases</i> , 2014, 14, 193-198.	1.5	27
13	Hantaviruses in Finnish soricomorphs: Evidence for two distinct hantaviruses carried by <i>Sorex araneus</i> suggesting ancient host-switch. <i>Infection, Genetics and Evolution</i> , 2014, 27, 51-61.	2.3	22
14	Carnivore conservation in practice: replicated management actions on a large spatial scale. <i>Journal of Applied Ecology</i> , 2013, 50, 59-67.	4.0	93
15	Experimental investigation of a hantavirus host-switch between arvicoline rodents <i>Lemmus lemmus</i> and <i>Myodes glareolus</i> . <i>Journal of Vector Ecology</i> , 2013, 38, 408-410.	1.0	4
16	Cocirculation of two Puumala hantavirus lineages in Latvia: A russian lineage described previously and a novel Latvian lineage. <i>Journal of Medical Virology</i> , 2012, 84, 314-318.	5.0	22
17	Genetic evidence for the presence of two distinct hantaviruses associated with <i>Apodemus</i> mice in Croatia and analysis of local strains. <i>Journal of Medical Virology</i> , 2011, 83, 108-114.	5.0	23
18	Orthopox Virus Infections in Eurasian Wild Rodents. <i>Vector-Borne and Zoonotic Diseases</i> , 2011, 11, 1133-1140.	1.5	53

#	ARTICLE	IF	CITATIONS
19	Systematic relationships of hymenolepidid cestodes of rodents and shrews inferred from sequences of 28S ribosomal RNA. <i>Zoologica Scripta</i> , 2010, 39, 631-641.	1.7	66
20	Molecular systematics and morphometrics of <i>Anoplocephaloides dentata</i> (Cestoda, Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 702 T	1.7	20
21	Genetic analysis of hantaviruses carried by <i>Myodes</i> and <i>Microtus</i> rodents in Buryatia. <i>Virology Journal</i> , 2008, 5, 4.	3.4	21
22	New Genetic Lineage of Tula Hantavirus in <i>Microtus arvalis obscurus</i> in Eastern Kazakhstan. <i>The Open Virology Journal</i> , 2008, 2, 32-36.	1.8	17
23	Serological evidence for Borna disease virus infection in humans, wild rodents and other vertebrates in Finland. <i>Journal of Clinical Virology</i> , 2007, 38, 64-69.	3.1	45
24	Taxonomy and genetic divergence of <i>Paranoplocephala kalelai</i> (Tenora, Haukisalmi et Henttonen, 1985) (Cestoda, Anoplocephalidae) in the grey-sided vole <i>Myodes rufocanus</i> in northern Fennoscandia. <i>Acta Parasitologica</i> , 2007, 52, 335.	1.1	6
25	Morphological and molecular characterisation of <i>Paranoplocephala buryatiensis</i> n. sp. and <i>P. longivaginata</i> Chechulin & Gulyaev, 1998 (Cestoda: Anoplocephalidae) in voles of the genus <i>Clethrionomys</i> . <i>Systematic Parasitology</i> , 2006, 66, 55-71.	1.1	18
26	Distribution of Puumala Hantavirus in Denmark: Analysis of Bank Voles ( <i>Clethrionomys glareolus</i> ) from Fyn and Jutland. <i>Vector-Borne and Zoonotic Diseases</i> , 2002, 2, 37-45.	1.5	16
27	Parasite diversity of Norwegian lemmings ( <i>Lemmus lemmus</i> ). <i>Journal of Zoology</i> , 2001, 253, 549-553.	1.7	3
28	Isolation of Dobrava Virus from <i>Apodemus flavicollis</i> in Greece. <i>Journal of Clinical Microbiology</i> , 2001, 39, 2291-2293.	3.9	27
29	Molecular evolution of Puumala hantavirus in Fennoscandia: phylogenetic analysis of strains from two recolonization routes, Karelia and Denmark. <i>Journal of General Virology</i> , 2000, 81, 2833-2841.	2.9	44
30	Analysis of Puumala hantavirus genome in patients with nephropathia epidemica and rodent carriers from the sites of infection. , 1999, 59, 397-405.		37
31	Isolation and Characterization of a Hantavirus from <i>Lemmus sibiricus</i> : Evidence for Host Switch during Hantavirus Evolution. <i>Journal of Virology</i> , 1999, 73, 5586-5592.	3.4	128
32	Genetic variation of wild Puumala viruses within the serotype, local rodent populations and individual animal. <i>Virus Research</i> , 1995, 38, 25-41.	2.2	82
33	Review of tapeworms of rodents in the Republic of Buryatia, with emphasis on anoplocephalid cestodes. <i>ZooKeys</i> , 0, 8, 1-18.	1.1	12