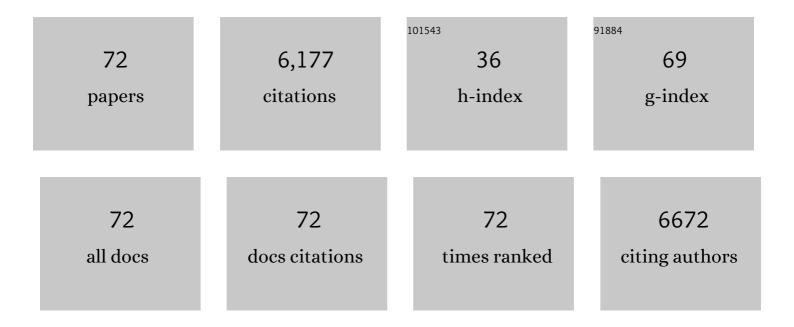
## Mika O Salminen

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Identification of Breakpoints in Intergenotypic Recombinants of HIV Type 1 by Bootscanning. AIDS Research and Human Retroviruses, 1995, 11, 1423-1425.	1.1	827
2	An automated genotyping system for analysis of HIV-1 and other microbial sequences. Bioinformatics, 2005, 21, 3797-3800.	4.1	468
3	Prevalence of Drugâ€Resistant HIVâ€1 Variants in Untreated Individuals in Europe: Implications for Clinical Management. Journal of Infectious Diseases, 2005, 192, 958-966.	4.0	385
4	Global and regional molecular epidemiology of HIV-1, 1990–2015: a systematic review, global survey, and trend analysis. Lancet Infectious Diseases, The, 2019, 19, 143-155.	9.1	255
5	Interventions to prevent HIV and Hepatitis C in people who inject drugs: A review of reviews to assess evidence of effectiveness. International Journal of Drug Policy, 2014, 25, 34-52.	3.3	238
6	Serological and molecular findings during SARS-CoV-2 infection: the first case study in Finland, January to February 2020. Eurosurveillance, 2020, 25, .	7.0	226
7	Full Genome Sequences of Human Immunodeficiency Virus Type 1 Subtypes G and A/G Intersubtype Recombinants. Virology, 1998, 247, 22-31.	2.4	224
8	Evidence of Recombination among Enteroviruses. Journal of Virology, 1999, 73, 8741-8749.	3.4	214
9	Transmission of Drugâ€Resistant HIVâ€1 Is Stabilizing in Europe. Journal of Infectious Diseases, 2009, 200, 1503-1508.	4.0	213
10	Infection with hepatitis B and C virus in Europe: a systematic review of prevalence and cost-effectiveness of screening. BMC Infectious Diseases, 2013, 13, 181.	2.9	190
11	Recovery of Virtually Full-Length HIV-1 Provirus of Diverse Subtypes from Primary Virus Cultures Using the Polymerase Chain Reaction. Virology, 1995, 213, 80-86.	2.4	189
12	HIV-1 genetic subtype A/B recombinant strain causing an explosive epidemic in injecting drug users in Kaliningrad. Aids, 1998, 12, 1907-1919.	2.2	187
13	Near Full-Length Clones and Reference Sequences for Subtype C Isolates of HIV Type 1 from Three Different Continents. AIDS Research and Human Retroviruses, 2001, 17, 161-168.	1.1	182
14	Simplified Strategy for Detection of Recombinant Human Immunodeficiency Virus Type 1 Group M Isolates by <i>gag/env</i> Heteroduplex Mobility Assay. Journal of Virology, 2000, 74, 363-370.	3.4	152
15	Tracing the HIV-1 subtype B mobility in Europe: a phylogeographic approach. Retrovirology, 2009, 6, 49.	2.0	114
16	Detection of HIV Type 1 <i>env</i> Subtypes A, B, C, and E in Asia Using Dried Blood Spots: A New Surveillance Tool for Molecular Epidemiology. AIDS Research and Human Retroviruses, 1996, 12, 1435-1441.	1.1	113
17	Transmission of drug-resistant HIV-1 in Europe remains limited to single classes. Aids, 2008, 22, 625-635.	2.2	109
18	Understanding coronavirus disease (COVID-19) risk perceptions among the public to enhance risk communication efforts: a practical approach for outbreaks, Finland, February 2020. Eurosurveillance, 2020, 25, .	7.0	107

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19	The Calculated Genetic Barrier for Antiretroviral Drug Resistance Substitutions Is Largely Similar for Different HIV-1 Subtypes. Journal of Acquired Immune Deficiency Syndromes (1999), 2006, 41, 352-360.	2.1	90
20	An Isolate of Human Immunodeficiency Virus Type 1 Originally Classified as Subtype I Represents a Complex Mosaic Comprising Three Different Group M Subtypes (A, G, and I). Journal of Virology, 1998, 72, 10234-10241.	3.4	90
21	Characterization of subtype A HIV-1 from Africa by full genome sequencing. Aids, 1999, 13, 1819-1826.	2.2	87
22	New HIV diagnoses among adults aged 50 years or older in 31 European countries, 2004–15: an analysis of surveillance data. Lancet HIV,the, 2017, 4, e514-e521.	4.7	87
23	Full-Length Sequence of an Ethiopian Human Immunodeficiency Virus Type 1 (HIV-1) Isolate of Genetic Subtype C. AIDS Research and Human Retroviruses, 1996, 12, 1329-1339.	1.1	86
24	Evidence of Two Distinct Subsubtypes within the HIV-1 Subtype A Radiation. AIDS Research and Human Retroviruses, 2001, 17, 675-688.	1.1	82
25	Human Immunodeficiency Virus Type 1 Subtypes Have a Distinct Long Terminal Repeat That Determines the Replication Rate in a Host-Cell-Specific Manner. Journal of Virology, 2004, 78, 3675-3683.	3.4	82
26	Molecular epidemiology of coxsackievirus B4 and disclosure of the correct VP1/2Apro cleavage site: evidence for high genomic diversity and long-term endemicity of distinct genotypes. Journal of General Virology, 2000, 81, 803-812.	2.9	65
27	Infectious Molecular Clones with the Nonhomologous Dimer Initiation Sequences Found in Different Subtypes of Human Immunodeficiency Virus Type 1 Can Recombine and Initiate a Spreading Infection In Vitro. Journal of Virology, 1998, 72, 3991-3998.	3.4	65
28	An AB Recombinant and Its Parental HIV Type 1 Strains in the Area of the Former Soviet Union: Low Requirements for Sequence Identity in Recombination. AIDS Research and Human Retroviruses, 2000, 16, 1047-1053.	1.1	63
29	Molecular Epidemiology of HIV-1 Based on Phylogenetic Analysis of in Vivo gag p7/p9 Direct Sequences. Virology, 1993, 195, 185-194.	2.4	56
30	<b>Short Communication</b> : Contribution of the CCR5 and MBL Genes to Susceptibility to HIV Type 1 Infection in the Finnish Population. AIDS Research and Human Retroviruses, 1998, 14, 695-698.	1.1	56
31	Global and regional epidemiology of HIV-1 recombinants in 1990–2015: a systematic review and global survey. Lancet HIV,the, 2020, 7, e772-e781.	4.7	51
32	Two Viral Strains and a Possible Novel Recombinant Are Responsible for the Explosive Injecting Drug Use-Associated HIV Type 1 Epidemic in Estonia. AIDS Research and Human Retroviruses, 2004, 20, 1148-1156.	1.1	50
33	Possible Occurrence of a Genetic Bottleneck in Dengue Serotype 2 Viruses between the 1980 and 1987 Epidemic Seasons in Bangkok, Thailand. American Journal of Tropical Medicine and Hygiene, 1997, 57, 100-108.	1.4	47
34	Virtually Full-Length Subtype F and F/D Recombinant HIV-1 from Africa and South America. Virology, 2000, 269, 95-104.	2.4	44
35	Genotypic and Phenotypic Analysis of HIV Type 1 Primary Isolates from Western Cameroon. AIDS Research and Human Retroviruses, 2002, 18, 39-48.	1.1	44
36	Analysis of results from the Joint External Evaluation: examining its strength and assessing for trends among participating countries. Journal of Global Health, 2018, 8, 020416.	2.7	43

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37	HIV in hiding: methods and data requirements for the estimation of the number of people living with undiagnosed HIV. Aids, 2011, 25, 1017-1023.	2.2	37
38	Virtually Full-Length Sequences of HIV Type 1 Subtype J Reference Strains. AIDS Research and Human Retroviruses, 1999, 15, 293-297.	1.1	31
39	HIV Epidemiology in the Northwestern Federal District of Russia: Dominance of HIV Type 1 Subtype A. AIDS Research and Human Retroviruses, 2006, 22, 1074-1080.	1.1	31
40	An outbreak of the circulating recombinant form AECM240 HIV-1 in the Finnish injection drug user population. Aids, 2000, 14, 2613-2615.	2.2	31
41	A study on HIV and hepatitis C virus among commercial sex workers in Tallinn. Sexually Transmitted Infections, 2008, 84, 189-191.	1.9	30
42	Analysis of HIV-1 Genetic Subtypes in Finland Reveals Good Correlation Between Molecular and Epidemiological Data. Scandinavian Journal of Infectious Diseases, 2000, 32, 475-480.	1.5	26
43	DNA Sequence of the Long Terminal Repeat of Human Immunodeficiency Virus Type 1 Subtype A through G. AIDS Research and Human Retroviruses, 1999, 15, 485-488.	1.1	25
44	Construction and Biological Characterization of Infectious Molecular Clones of HIV-1 Subtypes B and E (CRF01_AE) Generated by the Polymerase Chain Reaction. Virology, 2000, 278, 103-110.	2.4	25
45	Tick-borne encephalitis virus infections in Lithuanian domestic animals and ticks. Scandinavian Journal of Infectious Diseases, 2005, 37, 742-746.	1.5	25
46	Diversity of the Human Immunodeficiency Virus Type 1 Envelope Glycoprotein in San Francisco Men's Health Study Participants. AIDS Research and Human Retroviruses, 1998, 14, 329-337.	1.1	23
47	Genetic Characterization of HIV-1 Strains in the Baltic Countries and Russia. Scandinavian Journal of Infectious Diseases, 1996, 28, 537-541.	1.5	21
48	HIV-1 in Ethiopia: Phylogenetic divergence from other HIV-1 strains. Virus Genes, 1991, 5, 359-66.	1.6	20
49	Should Europe have a universal hepatitis B vaccination programme?. BMJ, The, 2013, 347, f4057-f4057.	6.0	20
50	Respiratory diphtheria in an asylum seeker from Afghanistan arriving to Finland via Sweden, December 2015. Eurosurveillance, 2016, 21, .	7.0	20
51	Near Full-Length Genome Analysis of HIV Type 1 CRF02.AG, Subtype C and CRF02.AG Subtype G Recombinants. AIDS Research and Human Retroviruses, 2000, 16, 1183-1189.	1.1	18
52	Determinants of late HIV diagnosis among different transmission groups in Finland from 1985 to 2005. HIV Medicine, 2010, 11, 360-367.	2.2	18
53	New subtype of HIV-1 in Ethiopia. Lancet, The, 1990, 336, 942-942.	13.7	17
54	European monitoring of notifications of hepatitis C virus infection in the general population and among injecting drug users (IDUs) – the need to improve quality and comparability. Eurosurveillance, 2008, 13, .	7.0	17

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55	Morphological differentiation of human SH-SY5Y neuroblastoma cells inhibits human immunodeficiency virus type 1 infection. Journal of General Virology, 1994, 75, 201-206.	2.9	16
56	Rapid and Simple Characterization ofIn VivoHIV-1 Sequences Using Solid-Phase Direct Sequencing. AIDS Research and Human Retroviruses, 1992, 8, 1733-1742.	1.1	15
57	High plasma HIV load in the CRF01-AE outbreak among injecting drug users in Finland. Scandinavian Journal of Infectious Diseases, 2005, 37, 276-283.	1.5	15
58	Activation of integrated human immunodeficiency virus type 1 in human neuroblastoma cells by the cytokines tumour necrosis factor alpha and interleukin-6. Journal of General Virology, 1992, 73, 1753-1760.	2.9	13
59	HIV-1 subtype H near-full length genome reference strains and analysis of subtype-H-containing inter-subtype recombinants. Aids, 2000, 14, 1533-1543.	2.2	13
60	Characterization of novel recombinant HIV-1 genomes using the branching index. Virology, 2003, 316, 116-125.	2.4	13
61	Initial surveillance of 2009 influenza A(H1N1) pandemic in the European Union and European Economic Area, April – September 2009. Eurosurveillance, 2010, 15, .	7.0	13
62	Sequence Note: Complex Mosaic Structure of the Partial Envelope Sequence from a Gambian HIV Type 1 Isolate. AIDS Research and Human Retroviruses, 1996, 12, 169-171.	1.1	12
63	HIV Type 1 <i>env</i> Gene Diversity Detected in Taiwan. AIDS Research and Human Retroviruses, 1997, 13, 201-204.	1.1	11
64	HIV-1 Subtypes in Denmark. Scandinavian Journal of Infectious Diseases, 2001, 33, 697-701.	1.5	10
65	<b>Sequence Note</b> : Full-Length Sequence of Human Immunodeficiency Virus Type 1 Subtype A, Recombined with Subtype C in the <i>env</i> V3 Domain. AIDS Research and Human Retroviruses, 1998, 14, 289-292.	1.1	9
66	High plasma HIV load in the CRF01-AE outbreak among injecting drug users in Finland. Scandinavian Journal of Infectious Diseases, 2005, 37, 276-83.	1.5	9
67	Detecting and characterizing individual recombination events. , 2009, , 519-548.		4
68	HIV-infection in Estonia. Clinical and Diagnostic Virology, 1993, 1, 11-16.	1.7	3
69	HIV-1 D subtype viruses in Finland. Clinical and Diagnostic Virology, 1996, 5, 205-210.	1.7	3
70	Diversity and Distribution of <i>gag</i> and <i>env</i> Subtypes among 146 HIV Type 1 Isolates in Taiwan. AIDS Research and Human Retroviruses, 1997, 13, 1539-1543.	1.1	2
71	Characterization of novel recombinant HIV-1 genomes using the branching index. Virology, 2003, 316, 116-116.	2.4	0
72	Recombination and Molecular Epidemiology of HIV-1 and Enteroviruses. , 2002, , 97-119.		0

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