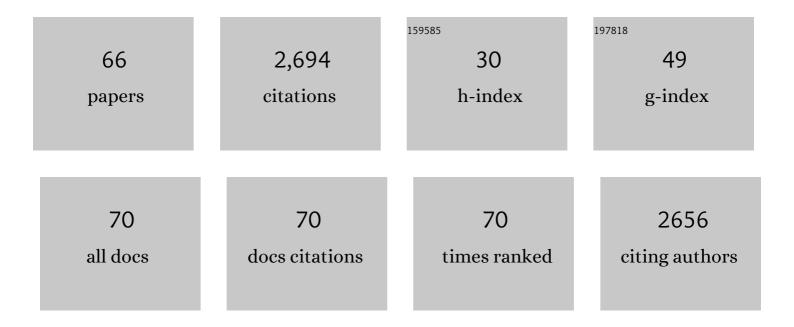
Joachim Frey

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Galactocerebroside biosynthesis pathways of <i>Mycoplasma</i> species: an antigen triggering Guillain–Barré–Stohl syndrome. Microbial Biotechnology, 2021, 14, 1201-1211.	4.2	5
2	Production of neutralizing antibodies against the secreted Clostridium chauvoei toxin A (CctA) upon blackleg vaccination. Anaerobe, 2019, 56, 78-87.	2.1	11
3	Systemic infection in European perch with thermoadapted virulent Aeromonas salmonicida (Perca) Tj ETQq1 1	0.784314 1.9	rgBT /Overloc
4	Reproduction of contagious caprine pleuropneumonia reveals the ability of convalescent sera to reduce hydrogen peroxide production in vitro. Veterinary Research, 2019, 50, 10.	3.0	24
5	Transposon-associated lincosamide resistance lnu (C) gene identified in Brachyspira hyodysenteriae ST83. Veterinary Microbiology, 2018, 214, 51-55.	1.9	30
6	Genetic Separation of Listeria monocytogenes Causing Central Nervous System Infections in Animals. Frontiers in Cellular and Infection Microbiology, 2018, 8, 20.	3.9	22
7	Blackleg in cattle: current understanding and future research needs. Ciencia Rural, 2018, 48, .	0.5	10
8	Remote Sensing of Potential Biosignatures from Rocky, Liquid, or Icy (Exo)Planetary Surfaces. Astrobiology, 2017, 17, 231-252.	3.0	29
9	Postepizootic Persistence of Asymptomatic Mycoplasma conjunctivae Infection in Iberian Ibex. Applied and Environmental Microbiology, 2017, 83, .	3.1	13
10	Infectious keratoconjunctivitis and occurrence of <i>Mycoplasma conjunctivae</i> and Chlamydiaceae in small domestic ruminants from Central Karakoram, Pakistan. Veterinary Record, 2017, 181, 237-237.	0.3	10
11	A review of methods used for studying the molecular epidemiology of Brachyspira hyodysenteriae. Veterinary Microbiology, 2017, 207, 181-194.	1.9	2
12	Clostridium chauvoei, an Evolutionary Dead-End Pathogen. Frontiers in Microbiology, 2017, 8, 1054.	3.5	33
13	Hyperinvasiveness and increased intercellular spread of Listeria monocytogenes sequence type 1 are independent of listeriolysin S, internalin F and internalin J1. Journal of Medical Microbiology, 2017, 66, 1053-1062.	1.8	33
14	Assessing Fifty Years of General Health Surveillance of Roe Deer in Switzerland: A Retrospective Analysis of Necropsy Reports. PLoS ONE, 2017, 12, e0170338.	2.5	33
15	Long-term dynamics of Mycoplasma conjunctivae at the wildlife-livestock interface in the Pyrenees. PLoS ONE, 2017, 12, e0186069.	2.5	13
16	Whole genome SNP analysis of bovine B. anthracis strains from Switzerland reflects strict regional separation of Simmental and Swiss Brown breeds in the past. Veterinary Microbiology, 2016, 196, 1-8.	1.9	14
17	Galactofuranose in <scp><i>M</i></scp> <i>ycoplasma mycoides</i> is important for membrane integrity and conceals adhesins but does not contribute to serum resistance. Molecular Microbiology, 2016, 99, 55-70.	2.5	34
18	Design of an Immersion Vaccine Against Aeromonad Septicemia in Perch (Perca fluviatilis L.). Methods in Molecular Biology, 2016, 1404, 203-209.	0.9	0

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19	Comparative genomics of Bacillus anthracis from the wool industry highlights polymorphisms of lineage A.Br.Vollum. Infection, Genetics and Evolution, 2016, 46, 50-58.	2.3	18
20	Listeria monocytogenes sequence type 1 is predominant in ruminant rhombencephalitis. Scientific Reports, 2016, 6, 36419.	3.3	105
21	Infectious keratoconjunctivitis in wild Caprinae: merging field observations and molecular analyses sheds light on factors shaping outbreak dynamics. BMC Veterinary Research, 2016, 13, 67.	1.9	12
22	Insect pathogenicity in plant-beneficial pseudomonads: phylogenetic distribution and comparative genomics. ISME Journal, 2016, 10, 2527-2542.	9.8	127
23	High quality draft genomes of the Mycoplasma mycoides subsp. mycoides challenge strains Afadé and B237. Standards in Genomic Sciences, 2015, 10, 89.	1.5	21
24	Listeria monocytogenes Spreads within the Brain by Actin-Based Intra-Axonal Migration. Infection and Immunity, 2015, 83, 2409-2419.	2.2	56
25	Invasion and persistence of Mycoplasma bovis in embryonic calf turbinate cells. Veterinary Research, 2015, 46, 53.	3.0	46
26	<i>AsaGEI2b</i> : a new variant of a genomic island identified in the <i>Aeromonas salmonicida</i> subsp. <i>salmonicida</i> JF3224 strain isolated from a wild fish in Switzerland. FEMS Microbiology Letters, 2015, 362, fnv093.	1.8	22
27	Vaccination of Cattle with the N Terminus of LppQ of Mycoplasma mycoides subsp. mycoides Results in Type III Immune Complex Disease upon Experimental Infection. Infection and Immunity, 2015, 83, 1992-2000.	2.2	22
28	Cyto-adherence of Mycoplasma mycoides subsp. mycoides to bovine lung epithelial cells. BMC Veterinary Research, 2015, 11, 27.	1.9	11
29	The immune response of bovine mammary epithelial cells to live or heat-inactivated Mycoplasma bovis. Veterinary Microbiology, 2015, 179, 336-340.	1.9	38
30	Field-Applicable Recombinase Polymerase Amplification Assay for Rapid Detection of Mycoplasma capricolum subsp. capripneumoniae. Journal of Clinical Microbiology, 2015, 53, 2810-2815.	3.9	55
31	A naturally occurring prfA truncation in a Listeria monocytogenes field strain contributes to reduced replication and cell-to-cell spread. Veterinary Microbiology, 2015, 179, 91-101.	1.9	37
32	Virulence, persistence and dissemination of Mycoplasma bovis. Veterinary Microbiology, 2015, 179, 15-22.	1.9	101
33	One-Step Identification of Five Prominent Chicken Salmonella Serovars and Biotypes. Journal of Clinical Microbiology, 2015, 53, 3881-3883.	3.9	44
34	Increased spread and replication efficiency of Listeria monocytogenes in organotypic brain-slices is related to multilocus variable number of tandem repeat analysis (MLVA) complex. BMC Microbiology, 2015, 15, 134.	3.3	20
35	Pyogranulomatous Pneumonia in Goats Caused by an Undescribed Porphyromonas Species, "Porphyromonas katsikii― Journal of Clinical Microbiology, 2015, 53, 795-798.	3.9	3
36	Patho-genetics of Clostridium chauvoei. Research in Microbiology, 2015, 166, 384-392.	2.1	37

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37	Identification of Animal Pasteurellaceae by MALDI-TOF Mass Spectrometry. Methods in Molecular Biology, 2015, 1247, 235-243.	0.9	10
38	Complete Genome Sequences of Virulent Mycoplasma capricolum subsp. <i>capripneumoniae</i> Strains F38 and ILRI181. Genome Announcements, 2014, 2, .	0.8	17
39	Molecular genetic analysis of Dichelobacter nodosus proteases AprV2/B2, AprV5/B5 and BprV/B in clinical material from European sheep flocks. Veterinary Microbiology, 2014, 168, 177-184.	1.9	34
40	Crystal ball – 2013. Microbial Biotechnology, 2013, 6, 3-16.	4.2	6
41	The Aeromonas salmonicida subsp. salmonicida exoproteome: global analysis, moonlighting proteins and putative antigens for vaccination against furunculosis. Proteome Science, 2013, 11, 44.	1.7	41
42	Cytotoxin CctA, a major virulence factor of Clostridium chauvoei conferring protective immunity against myonecrosis. Vaccine, 2012, 30, 5500-5505.	3.8	39
43	The role of RTX toxins in host specificity of animal pathogenic Pasteurellaceae. Veterinary Microbiology, 2011, 153, 51-58.	1.9	48
44	Genetic and functional characterization of the NanA sialidase from Clostridium chauvoei. Veterinary Research, 2011, 42, 2.	3.0	28
45	Alpine ibex (Capra i. ibex) is not a reservoir for chlamydial infections of domestic ruminants and humans. European Journal of Wildlife Research, 2011, 57, 233-240.	1.4	10
46	Pseudomonas chlororaphis subsp. piscium subsp. nov., isolated from freshwater fish. International Journal of Systematic and Evolutionary Microbiology, 2010, 60, 2753-2757.	1.7	34
47	Basfia succiniciproducens gen. nov., sp. nov., a new member of the family Pasteurellaceae isolated from bovine rumen. International Journal of Systematic and Evolutionary Microbiology, 2010, 60, 44-50.	1.7	87
48	Unmarked insertional mutagenesis in the bovine pathogen Mycoplasma mycoides subsp. mycoides SC: characterization of a lppQ mutant. Microbiology (United Kingdom), 2008, 154, 2427-2436.	1.8	20
49	Aeromonas Exoenzyme T of Aeromonas salmonicida Is a Bifunctional Protein That Targets the Host Cytoskeleton. Journal of Biological Chemistry, 2007, 282, 28843-28852.	3.4	56
50	AopP, a type III effector protein of Aeromonas salmonicida, inhibits the NF-κB signalling pathway. Microbiology (United Kingdom), 2006, 152, 2809-2818.	1.8	83
51	Genetic relatedness within the genus Campylobacter inferred from rpoB sequences. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 937-945.	1.7	72
52	Attenuated virulence of an Aeromonas salmonicida subsp. salmonicida type III secretion mutant in a rainbow trout model. Microbiology (United Kingdom), 2005, 151, 2111-2118.	1.8	87
53	Phylogeny of the family Pasteurellaceae based on rpoB sequences. International Journal of Systematic and Evolutionary Microbiology, 2004, 54, 1393-1399.	1.7	194
54	Detection, Identification, and Subtyping of Actinobacillus pleuropneumoniae. , 2003, 216, 87-96.		32

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#	Article	IF	CITATIONS
55	RTX toxins in Pasteurellaceae. International Journal of Medical Microbiology, 2002, 292, 149-158.	3.6	92
56	Genetic diversity amongMycoplasma species bovine group 7: Clonal isolates from an outbreak of polyarthritis, mastitis, and abortion in dairy cattle. Electrophoresis, 2001, 22, 3551-3561.	2.4	28
57	Target genes for virulence assessment ofEscherichia coliisolates from water, food and the environment. FEMS Microbiology Reviews, 2000, 24, 107-117.	8.6	125
58	Characterization of a Predominant Immunogenic Outer Membrane Protein of Riemerella anatipestifer. Vaccine Journal, 2000, 7, 168-174.	2.6	72
59	Target genes for virulence assessment of Escherichia coli isolates from water, food and the environment. FEMS Microbiology Reviews, 2000, 24, 107-117. Genomic and antigenic differences between the European and African/Australian clusters of	8.6	4
60	Mycoplasma mycoides subsp. mycoides SC The GenBank accession numbers for the nucleotide sequences determined in this work are: AF165134 for the 3·4Âkb HindIII fragment from M. mycoides subsp. mycoides SC strain L2; AF165135 for the analogous locus in strain Afadé (containing lppB and IS1634); and AF1651136 for the DNA segment containing lppB[MmymyLC] and ORF6[MmymyLC] from M. mycoides	1.8	73
61	subsp. mycoides LC strain Y-goat Microbiology (United Kingdom), 2000, 146, 477-486. Characterization of apxIVA, a new RTX determinant of Actinobacillus pleuropneumoniae. Microbiology (United Kingdom), 1999, 145, 2105-2116.	1.8	196
62	IS <i>1634</i> , a Novel Insertion Element Creating Long, Variable-Length Direct Repeats Which Is Specific for <i>Mycoplasma mycoides</i> subsp. <i>mycoides</i> Small-Colony Type. Journal of Bacteriology, 1999, 181, 1319-1323.	2.2	49
63	Identification of Clostridium chauvoei in cultures and clinical material from blackleg using PCR. Veterinary Microbiology, 1997, 57, 291-298.	1.9	25
64	Identification of a locus involved in the utilization of iron by Actinobacillus pleuropneumoniae. FEMS Microbiology Letters, 1996, 143, 1-6.	1.8	1
65	Association of the CAMP phenomenon in Actinobacillus pleuropneumoniae with the RTX toxins ApxI, ApxII and ApxIII. FEMS Microbiology Letters, 1994, 124, 245-251.	1.8	27
66	Use of a Microchip to Detect Antibiotic Resistance Genes inBacillus anthracis. , 0, , 147-152.		1