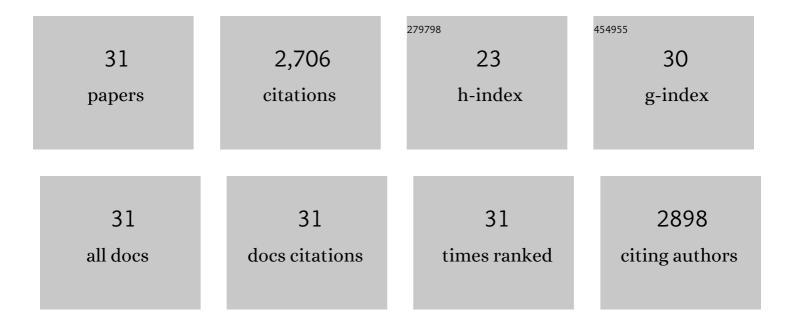
## Jan Lw Rademaker

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Predicting enteric methane emission of dairy cows with milk Fourier-transform infrared spectra and gas chromatography–based milk fatty acid profiles. Journal of Dairy Science, 2018, 101, 5582-5598.	3.4	30
2	Concurrence of spores of Clostridium tyrobutyricum, Clostridium beijerinckii and Paenibacillus polymyxa in silage, dairy cow faeces and raw milk. International Dairy Journal, 2016, 63, 70-77.	3.0	28
3	Verification of fresh grass feeding, pasture grazing and organic farming by cows farm milk fatty acid profile. Food Chemistry, 2014, 164, 234-241.	8.2	67
4	Verification of fresh grass feeding, pasture grazing and organic farming by FTIR spectroscopy analysis of bovine milk. Food Research International, 2014, 60, 59-65.	6.2	37
5	Competitive Selection of Lactic Acid Bacteria That Persist in the Human Oral Cavity. Applied and Environmental Microbiology, 2011, 77, 8445-8450.	3.1	12
6	High-Resolution Amplified Fragment Length Polymorphism Typing of Lactococcus lactis Strains Enables Identification of Genetic Markers for Subspecies-Related Phenotypes. Applied and Environmental Microbiology, 2011, 77, 5192-5198.	3.1	21
7	Phenotypic and genomic diversity of <i>Lactobacillus plantarum</i> strains isolated from various environmental niches. Environmental Microbiology, 2010, 12, 758-773.	3.8	262
8	Regulatory Phenotyping Reveals Important Diversity within the Species <i>Lactococcus lactis</i> . Applied and Environmental Microbiology, 2009, 75, 5687-5694.	3.1	26
9	An inter-laboratory ring trial for the detection and isolation of Mycobacterium avium subsp. paratuberculosis from raw milk artificially contaminated with naturally infected faeces. Food Microbiology, 2008, 25, 128-135.	4.2	29
10	Probiotic Lactobacillus plantarum 299v Does Not Counteract Unfavorable Phytohemagglutinin-Induced Changes in the Rat Intestinal Microbiota. Applied and Environmental Microbiology, 2008, 74, 5244-5249.	3.1	7
11	Diversity Analysis of Dairy and Nondairy <i>Lactococcus lactis</i> Isolates, Using a Novel Multilocus Sequence Analysis Scheme and (GTG) <sub>5</sub> -PCR Fingerprinting. Applied and Environmental Microbiology, 2007, 73, 7128-7137.	3.1	95
12	Effective Heat Inactivation of Mycobacterium avium subsp. paratuberculosis in Raw Milk Contaminated with Naturally Infected Feces. Applied and Environmental Microbiology, 2007, 73, 4185-4190.	3.1	49
13	Assessment of microbial population dynamics during yoghurt and hard cheese fermentation and ripening by DNA population fingerprinting. International Dairy Journal, 2006, 16, 457-466.	3.0	32
14	Classification and Identification of Xanthomonas translucens Isolates, Including Those Pathogenic to Ornamental Asparagus. Phytopathology, 2006, 96, 876-884.	2.2	41
15	Genetic diversity and biogeography of haloalkaliphilic sulphur-oxidizing bacteria belonging to the genus Thioalkalivibrio. FEMS Microbiology Ecology, 2006, 56, 95-101.	2.7	65
16	Growth stimulation of Brevibacterium sp. by siderophores. Journal of Applied Microbiology, 2006, 101, 637-646.	3.1	52
17	Natural diversity and adaptive responses of Lactococcus lactis. Current Opinion in Biotechnology, 2006, 17, 183-190.	6.6	97
18	A Comprehensive Species to Strain Taxonomic Framework for Xanthomonas. Phytopathology, 2005, 95, 1098-1111.	2.2	138

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#	Article	IF	CITATIONS
19	The surface microflora dynamics of bacterial smear-ripened Tilsit cheese determined by T-RFLP DNA population fingerprint analysis. International Dairy Journal, 2005, 15, 785-794.	3.0	50
20	Experimental taxonomic studies in Psilocybe sect. Psilocybe. Mycological Research, 2002, 106, 1251-1261.	2.5	2
21	The beneficial plant growth-promoting association of Rhizobium leguminosarum bv. trifolii with rice roots. Functional Plant Biology, 2001, 28, 845.	2.1	116
22	Comparison of AFLP and rep-PCR genomic fingerprinting with DNA-DNA homology studies: Xanthomonas as a model system International Journal of Systematic and Evolutionary Microbiology, 2000, 50, 665-677.	1.7	316
23	Synopsis on the Taxonomy of the Genus Xanthomonas. Phytopathology, 2000, 90, 677-682.	2.2	121
24	Amplified-Fragment Length Polymorphism Analysis: the State of an Art. Journal of Clinical Microbiology, 1999, 37, 3083-3091.	3.9	416
25	Identification of bacterial rep-PCR genomic fingerprints using a backpropagation neural network. FEMS Microbiology Letters, 1999, 177, 249-256.	1.8	20
26	THETHREEDSOFPCR-BASEDGENOMICANALYSISOFPHYTOBACTERIA: Diversity, Detection, and Disease Diagnosis. Annual Review of Phytopathology, 1999, 37, 81-125.	7.8	255
27	Multiphasic Analysis of Xanthomonads Causing Bacterial Spot Disease on Tomato and Pepper in the Caribbean and Central America: Evidence for Common Lineages Within and Between Countries. Phytopathology, 1999, 89, 328-335.	2.2	84
28	Identification of bacterial rep-PCR genomic fingerprints using a backpropagation neural network. FEMS Microbiology Letters, 1999, 177, 249-256.	1.8	0
29	Genotypic Characterization of <1>Bradyrhizobium 1 Strains Nodulating Endemic Woody Legumes of the Canary Islands by PCR-Restriction Fragment Length Polymorphism Analysis of Genes Encoding 16S rRNA (16S rDNA) and 16S-23S rDNA Intergenic Spacers, Repetitive Extragenic Palindromic PCR Genomic Fingerprinting, and Partial 16S rDNA Sequencing. Applied and Environmental Microbiology, 1998, 64,	3.1	200
30	Detection and identification of <i>Clavibacter michiganensis</i> subsp. <i>sepedonicus</i> and <i>Clavibacter michiganensis</i> subsp. <i>michiganensis</i> by nonradioactive hybridization, polymerase chain reaction, and restriction enzyme analysis. Canadian Journal of Microbiology, 1994, 40, 1007-1018.	1.7	22
31	Specific 16S ribosomal RNA targeted oligonucleotide probe against <i>Clavibacter michiganensis</i> subsp. <i>sepedonicus</i> . Canadian Journal of Microbiology, 1993, 39, 1029-1034.	1.7	16