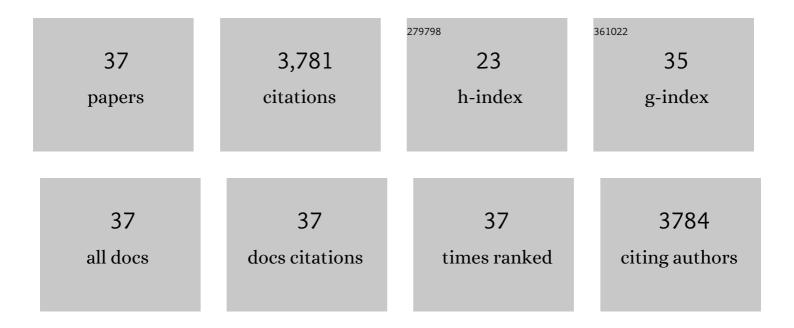
Yechezkel Kashi

List of Publications by Year in descending order

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VECHEZKEL KASHL

#	Article	IF	CITATIONS
1	Residues in chaperonin GroEL required for polypeptide binding and release. Nature, 1994, 371, 614-619.	27.8	653
2	Simple sequence repeats as advantageous mutators in evolution. Trends in Genetics, 2006, 22, 253-259.	6.7	471
3	Mechanism of GroEL action: Productive release of polypeptide from a sequestered position under groes. Cell, 1995, 83, 577-587.	28.9	431
4	Simple sequence repeats as a source of quantitative genetic variation. Trends in Genetics, 1997, 13, 74-78.	6.7	404
5	GroEL-mediated protein folding proceeds by multiple rounds of binding and release of nonnative forms. Cell, 1994, 78, 693-702.	28.9	375
6	Radiation induces proinflammatory dysbiosis: transmission of inflammatory susceptibility by host cytokine induction. Gut, 2018, 67, 97-107.	12.1	229
7	Evolutionary tuning knobs. Endeavour, 1997, 21, 36-40.	0.4	136
8	Active food packaging films with synergistic antimicrobial activity. Food Control, 2017, 76, 117-126.	5.5	120
9	Resistance to Adjuvant Arthritis Is Due to Protective Antibodies Against Heat Shock Protein Surface Epitopes and the Induction of IL-10 Secretion. Journal of Immunology, 2002, 168, 6463-6469.	0.8	82
10	Finding Approximate Tandem Repeats in Genomic Sequences. Journal of Computational Biology, 2005, 12, 928-942.	1.6	82
11	Vibrio cholerae Hemagglutinin/Protease Degrades Chironomid Egg Masses. Applied and Environmental Microbiology, 2003, 69, 4200-4204.	3.1	78
12	Vibrio cholerae Strain Typing and Phylogeny Study Based on Simple Sequence Repeats. Journal of Clinical Microbiology, 2007, 45, 736-746.	3.9	77
13	Adult non-biting midges: possible windborne carriers of Vibrio cholerae non-O1 non-O139. Environmental Microbiology, 2005, 7, 576-585.	3.8	70
14	Antibacterial and antifungal LDPE films for active packaging. Polymers for Advanced Technologies, 2015, 26, 110-116.	3.2	59
15	A PCR Method Based on 16S rRNA Sequence for Simultaneous Detection of the Genus Listeria and the Species Listeria monocytogenes in Food Products. Journal of Food Protection, 2003, 66, 1658-1665.	1.7	55
16	Predominant Effect of Host Genetics on Levels of Lactobacillus johnsonii Bacteria in the Mouse Gut. Applied and Environmental Microbiology, 2011, 77, 6531-6538.	3.1	39
17	Characterization of the 5' flanking region of the growth hormone gene of the marine teleost, gilthead sea bream Sparus aurata: analysis of a polymorphic microsatellite in the proximal promoter. Fisheries Science, 2005, 71, 479-490.	1.6	38
18	Murine Genetic Background Has a Stronger Impact on the Composition of the Gut Microbiota than Maternal Inoculation or Exposure to Unlike Exogenous Microbiota. Applied and Environmental Microbiology, 2019, 85, .	3.1	37

YECHEZKEL KASHI

#	Article	IF	CITATIONS
19	Indication for Co-evolution of Lactobacillus johnsonii with its hosts. BMC Microbiology, 2012, 12, 149.	3.3	31
20	ICEVchInd5 is prevalent in epidemic Vibrio cholerae O1 El Tor strains isolated in India. International Journal of Medical Microbiology, 2011, 301, 318-324.	3.6	27
21	Array-Based Binary Analysis for Bacterial Typing. Analytical Chemistry, 2005, 77, 319-326.	6.5	26
22	Epidemiologic Study of <i>Vibrio vulnificus</i> Infections by Using Variable Number Tandem Repeats. Emerging Infectious Diseases, 2009, 15, 1282-1285.	4.3	26
23	Phylogeny and Strain Typing of Escherichiacoli , Inferred from Variation at Mononucleotide RepeatLoci. Applied and Environmental Microbiology, 2004, 70, 2464-2473.	3.1	25
24	The Dimeric Structure of the Cpn60.2 Chaperonin of Mycobacterium tuberculosis at 2.8ÂÃ Reveals Possible Modes of Function. Journal of Molecular Biology, 2011, 412, 192-203.	4.2	25
25	Intestinal Dysbiosis in Carriers of Carbapenem-Resistant <i>Enterobacteriaceae</i> . MSphere, 2020, 5, .	2.9	25
26	The association between nonâ€biting midges and <i>Vibrio cholerae</i> . Environmental Microbiology, 2008, 10, 3193-3200.	3.8	24
27	Genetic diversity of the human pathogen Vibrio vulnificus: A new phylogroup. International Journal of Food Microbiology, 2012, 153, 436-443.	4.7	23
28	<i>Vibrio vulnificus</i> Typing Based on Simple Sequence Repeats: Insights into the Biotype 3 Group. Journal of Clinical Microbiology, 2007, 45, 2951-2959.	3.9	22
29	Finding approximate tandem repeats in genomic sequences. , 2004, , .		19
30	Biodiversity of Enterococcus faecalis based on genomic typing. International Journal of Food Microbiology, 2013, 165, 27-34.	4.7	15
31	Towards the definition of pathogenic microbe. International Journal of Food Microbiology, 2006, 112, 236-243.	4.7	11
32	<i>Vibrio cholerae</i> Autoinducer CAI-1 Interferes with <i>Pseudomonas aeruginosa</i> Quorum Sensing and Inhibits its Growth. ACS Chemical Biology, 2012, 7, 659-665.	3.4	10
33	Mono-nucleotide repeats (MNRs): a neglected polymorphism for generating high density genetic maps in silico. Human Genetics, 2004, 115, 213-20.	3.8	9
34	Amplified Intergenic Locus Polymorphism as a Basis for Bacterial Typing of Listeria spp. and Escherichia coli. Applied and Environmental Microbiology, 2005, 71, 3144-3152.	3.1	9
35	Environmental monitoring of <i>Vibrio cholerae</i> using chironomids in India. Environmental Microbiology Reports, 2010, 2, 96-103.	2.4	9
36	Adhesion of Vibrio cholerae to Granular Starches. Applied and Environmental Microbiology, 2005, 71, 4850-4855.	3.1	8

#	Article	IF	CITATIONS
37	Host Genetics and Gut Microbiota. , 2012, , 281-295.		1