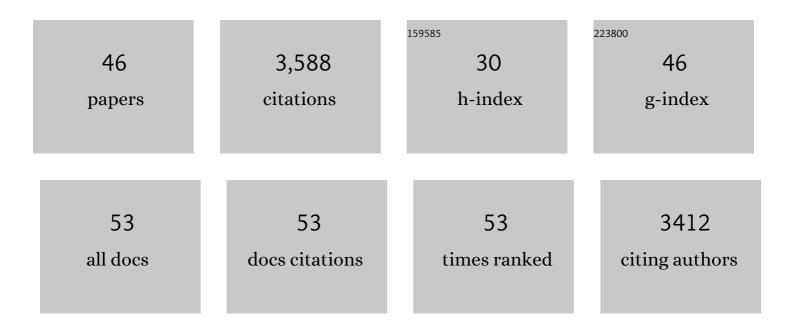
Kim R Hardie

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4737715/publications.pdf Version: 2024-02-01



KIM P HADDIE

#	Article	IF	CITATIONS
1	Fluorescent nanosensors reveal dynamic pH gradients during biofilm formation. Npj Biofilms and Microbiomes, 2021, 7, 50.	6.4	19
2	Disruption of the Pseudomonas aeruginosa Tat system perturbs PQS-dependent quorum sensing and biofilm maturation through lack of the Rieske cytochrome bc1 sub-unit. PLoS Pathogens, 2021, 17, e1009425.	4.7	8
3	Challenges and solutions in polymer drug delivery for bacterial biofilm treatment: A tissue-by-tissue account. Advanced Drug Delivery Reviews, 2021, 178, 113973.	13.7	36
4	JMM Profile: Carbapenems: a broad-spectrum antibiotic. Journal of Medical Microbiology, 2021, 70, .	1.8	29
5	Improving children's and their visitors' hand hygiene compliance. Journal of Infection Prevention, 2020, 21, 60-67.	0.9	11
6	Cryo-OrbiSIMS for 3D Molecular Imaging of a Bacterial Biofilm in Its Native State. Analytical Chemistry, 2020, 92, 9008-9015.	6.5	37
7	A Simple Polymicrobial Biofilm Keratinocyte Colonization Model for Exploring Interactions Between Commensals, Pathogens and Antimicrobials. Frontiers in Microbiology, 2020, 11, 291.	3.5	23
8	New frontiers against antibiotic resistance: A Raman-based approach for rapid detection of bacterial susceptibility and biocide-induced antibiotic cross-tolerance. Sensors and Actuators B: Chemical, 2020, 309, 127774.	7.8	19
9	Antimicrobial resistance: the good, the bad, and the ugly. Emerging Topics in Life Sciences, 2020, 4, 129-136.	2.6	10
10	Editorial Conference Report 6th Midlands Molecular Microbiology Meeting (M4) 2019, 9/10 September, Nottingham University Jubilee Conference Centre, UK. Journal of Medical Microbiology, 2020, 69, 1-2.	1.8	3
11	Dual bioresponsive antibiotic and quorum sensing inhibitor combination nanoparticles for treatment of <i>Pseudomonas aeruginosa</i> biofilms <i>in vitro</i> and <i>ex vivo</i> . Biomaterials Science, 2019, 7, 4099-4111.	5.4	56
12	Microfluidic-based Growth and Imaging of Bacterial Biofilms. Bio-protocol, 2019, 9, .	0.4	4
13	The Role of Flagella in Clostridium difficile Pathogenesis: Comparison between a Non-Epidemic and an Epidemic Strain. PLoS ONE, 2013, 8, e73026.	2.5	117
14	A Novel Virulence Strategy for Pseudomonas aeruginosa Mediated by an Autotransporter with Arginine-Specific Aminopeptidase Activity. PLoS Pathogens, 2012, 8, e1002854.	4.7	45
15	Simultaneous quantitative profiling of N-acyl-l-homoserine lactone and 2-alkyl-4(1H)-quinolone families of quorum-sensing signaling molecules using LC-MS/MS. Analytical and Bioanalytical Chemistry, 2011, 399, 839-850.	3.7	168
16	Comparative genomics and proteomics of Helicobacter mustelae, an ulcerogenic and carcinogenic gastric pathogen. BMC Genomics, 2010, 11, 164.	2.8	40
17	In Helicobacter pylori auto-inducer-2, but not LuxS/MccAB catalysed reverse transsulphuration, regulates motility through modulation of flagellar gene transcription. BMC Microbiology, 2010, 10, 210.	3.3	34
18	Quantitative liquid chromatography–tandem mass spectrometry profiling of activated methyl cycle metabolites involved in LuxS-dependent quorum sensing in Escherichia coli. Analytical Biochemistry, 2010, 403, 20-29.	2.4	56

Kim R Hardie

#	Article	IF	CITATIONS
19	In <i>Helicobacter pylori</i> , LuxS Is a Key Enzyme in Cysteine Provision through a Reverse Transsulfuration Pathway. Journal of Bacteriology, 2010, 192, 1184-1192.	2.2	34
20	Growth Deficiencies of Neisseria meningitidis pfs and luxS Mutants Are Not Due to Inactivation of Quorum Sensing. Journal of Bacteriology, 2009, 191, 1293-1302.	2.2	35
21	Al-2 does not function as a quorum sensing molecule in Campylobacter jejuni during exponential growth in vitro. BMC Microbiology, 2009, 9, 214.	3.3	38
22	Establishing bacterial communities by 'word of mouth': LuxS and autoinducer 2 in biofilm development. Nature Reviews Microbiology, 2008, 6, 635-643.	28.6	206
23	LuxS-independent formation of Al-2 from ribulose-5-phosphate. BMC Microbiology, 2008, 8, 98.	3.3	36
24	Functional association between the Helicobacter pylori virulence factors VacA and CagA. Journal of Medical Microbiology, 2008, 57, 145-150.	1.8	100
25	Helicobacter pylori FlhB Function: the FlhB C-Terminal Homologue HP1575 Acts as a "Spare Part―To Permit Flagellar Export When the HP0770 FlhB CC Domain Is Deleted. Journal of Bacteriology, 2006, 188, 7531-7541.	2.2	17
26	Making 'sense' of metabolism: autoinducer-2, LUXS and pathogenic bacteria. Nature Reviews Microbiology, 2005, 3, 383-396.	28.6	533
27	All subtypes of the cytotoxin VacA adsorb to the surface of Helicobacter pylori post-secretion. Journal of Medical Microbiology, 2005, 54, 621-630.	1.8	7
28	Functional complementation ofE. coli secDandsecGmutants byHelicobacter pylorihomologues. FEMS Microbiology Letters, 2003, 229, 57-63.	1.8	3
29	Electrostatic sensor for identifying interactions between peptides and bacterial membranes. Molecular Immunology, 2003, 40, 407-411.	2.2	14
30	LuxS and Autoinducer-2: Their Contribution to Quorum Sensing and Metabolism in Bacteria. Advances in Applied Microbiology, 2003, 53, 291-396.	2.4	142
31	Autoinducer 2 activity in Escherichia coli culture supernatants can be actively reduced despite maintenance of an active synthase, LuxS. Microbiology (United Kingdom), 2003, 149, 715-728.	1.8	43
32	NapA protects Helicobacter pylori from oxidative stress damage, and its production is influenced by the ferric uptake regulator. Journal of Medical Microbiology, 2003, 52, 461-469.	1.8	99
33	Role of Neisseria meningitidis luxS in Cell-to-Cell Signaling and Bacteremic Infection. Infection and Immunity, 2002, 70, 2245-2248.	2.2	71
34	LuxS: its role in central metabolism and the in vitro synthesis of 4-hydroxy-5-methyl-3(2H)-furanone. Microbiology (United Kingdom), 2002, 148, 909-922.	1.8	314
35	LuxS-dependent quorum sensing in Porphyromonas gingivalis modulates protease and haemagglutinin activities but is not essential for virulence. Microbiology (United Kingdom), 2002, 148, 763-772.	1.8	140
36	Bacterial cell-to-cell communication: sorry, can't talk now — gone to lunch!. Current Opinion in Microbiology, 2002, 5, 216-222.	5.1	301

Kim R Hardie

#	Article	IF	CITATIONS
37	In vitro biosynthesis of the Pseudomonas aeruginosa quorum-sensing signal molecule N-butanoyl-L-homoserine lactone. Molecular Microbiology, 2002, 28, 193-203.	2.5	73
38	Determining the molar mass of a plasma substitute succinylated gelatin by size exclusion chromatography–multi-angle laser light scattering, sedimentation equilibrium and conventional size exclusion chromatography. Journal of Chromatography A, 2002, 957, 139-148.	3.7	15
39	Genetic Dissection of the Outer Membrane Secretin PulD: Are There Distinct Domains for Multimerization and Secretion Specificity?. Journal of Bacteriology, 1999, 181, 7212-7220.	2.2	79
40	Recent progress and future directions in studies of the main terminal branch of the general secretory pathway in Gram-negative bacteria – a review. Gene, 1997, 192, 13-19.	2.2	106
41	The Câ€ŧerminal domain of the secretin PulD contains the binding site for its cognate chaperone, PulS, and confers PulS dependence on plV ^{f1} function. Molecular Microbiology, 1997, 24, 465-475.	2.5	99
42	Analysis of the carbapenem gene cluster of <i>Erwinia carotovora</i> : definition of the antibiotic biosynthetic genes and evidence for a novel βâ€lactam resistance mechanism. Molecular Microbiology, 1997, 26, 545-556.	2.5	91
43	The secretinâ€specific, chaperoneâ€like protein of the general secretory pathway: separation of proteolytic protection and piloting functions . Molecular Microbiology, 1996, 22, 967-976.	2.5	116
44	Independent interaction of the acyltransferase HlyC with two maturation domains of the Escherichia coli toxin HlyA. Molecular Microbiology, 1996, 20, 813-822.	2.5	33
45	Vibrio spp. secrete proaerolysin as a folded dimer without the need for disulphide bond formation. Molecular Microbiology, 1995, 17, 1035-1044.	2.5	78
46	Pore formation in artificial membranes by the secreted hemolysins of Proteus vulgaris and Morganella morganii. FEBS Journal, 1994, 220, 339-347.	0.2	49