

# Julian Ian Rood

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

256  
papers

12,766  
citations

61  
h-index

103  
g-index

268  
ext. papers

14,152  
ext. citations

5.2  
avg. IF

6.06  
L-index

#	Paper	IF	Citations
256	Reevaluation of whether a Functional Agr-like Quorum-Sensing System Is Necessary for Production of Wild-Type Levels of Epsilon-Toxin by Clostridium perfringens Type D Strains.. <i>MBio</i> , <b>2022</b> , e0049622	7.8	
255	Retraction for Chen et al., "Epsilon-Toxin Production by Clostridium perfringens Type D Strain CN3718 Is Dependent upon the Operon but Not the VirS/VirR Two-Component Regulatory System".. <i>MBio</i> , <b>2022</b> , e0049522	7.8	
254	The ever-expanding tcp conjugation locus of pCW3 from Clostridium perfringens. <i>Plasmid</i> , <b>2021</b> , 113, 102516	3.3	2
253	Cardiopulmonary Lesions in Sheep Produced by Experimental Acute Type D Enterotoxemia. <i>Veterinary Pathology</i> , <b>2021</b> , 58, 103-113	2.8	1
252	A pasture-based experimental infection model for footrot in sheep. <i>Small Ruminant Research</i> , <b>2021</b> , 195, 106305	1.7	2
251	Classifying mobile genetic elements and their interactions from sequence data: The importance of existing biological knowledge. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2021</b> , 118,	11.5	1
250	The EngCP endo $\beta$ -N-acetylgalactosaminidase is a virulence factor involved in Clostridium perfringens gas gangrene infections. <i>International Journal of Medical Microbiology</i> , <b>2020</b> , 310, 151398	3.7	1
249	Two putative zinc metalloproteases contribute to the virulence of strains that cause avian necrotic enteritis. <i>Journal of Veterinary Diagnostic Investigation</i> , <b>2020</b> , 32, 259-267	1.5	8
248	The Tcp plasmids of Clostridium perfringens require the resP gene to ensure stable inheritance. <i>Plasmid</i> , <b>2020</b> , 107, 102461	3.3	1
247	Virulence Plasmids of the Pathogenic Clostridia. <i>Microbiology Spectrum</i> , <b>2019</b> , 7,	8.9	7
246	pCP13, a representative of a new family of conjugative toxin plasmids in Clostridium perfringens. <i>Plasmid</i> , <b>2019</b> , 102, 37-45	3.3	9
245	Histotoxic Clostridial Infections. <i>Microbiology Spectrum</i> , <b>2019</b> , 7,	8.9	8
244	Virulence Plasmids of the Pathogenic Clostridia <b>2019</b> , 954-976		
243	Sporulation and Germination in Clostridial Pathogens <b>2019</b> , 903-926		0
242	Enterotoxic Clostridia: Clostridium perfringens Enteric Diseases <b>2019</b> , 977-990		2
241	Staphylococcal Plasmids, Transposable and Integrative Elements <b>2019</b> , 499-520		
240	Enterotoxic Clostridia: Clostridioides difficile Infections <b>2019</b> , 991-1011		

239 Enterococcal Genetics **2019**, 398-425

238 Expansion of the Clostridium perfringens toxin-based typing scheme. *Anaerobe*, **2018**, 53, 5-10 2.8 219

237 Concurrent Host-Pathogen Transcriptional Responses in a Murine Myonecrosis Infection. *MBio*, **2018**, 9, 7.8 17

236 Pathogenicity Locus Plasmid pCS1-1 Encodes a Novel Clostridial Conjugation Locus. *MBio*, **2018**, 9, 7.8 14

235 The incidence of Clostridioides difficile and Clostridium perfringens netF-positive strains in diarrheic dogs. *Anaerobe*, **2018**, 49, 58-62 2.8 22

234 Whole genome analysis reveals the diversity and evolutionary relationships between necrotic enteritis-causing strains of Clostridium perfringens. *BMC Genomics*, **2018**, 19, 379 4.5 29

233 Antibiotic resistance plasmids and mobile genetic elements of Clostridium perfringens. *Plasmid*, **2018**, 99, 32-39 3.3 11

232 Vibrational spectroscopy combined with transcriptomic analysis for investigation of bacterial responses towards acid stress. *Applied Microbiology and Biotechnology*, **2018**, 102, 333-343 5.7 9

231 Crystal structure of TcpK in complex with oriT DNA of the antibiotic resistance plasmid pCW3. *Nature Communications*, **2018**, 9, 3732 17.4 13

230 The Tcp conjugation system of Clostridium perfringens. *Plasmid*, **2017**, 91, 28-36 3.3 27

229 Bovine antibodies targeting primary and recurrent Clostridium difficile disease are a potent antibiotic alternative. *Scientific Reports*, **2017**, 7, 3665 4.9 25

228 Evidence that compatibility of closely related replicons in Clostridium perfringens depends on linkage to parMRC-like partitioning systems of different subfamilies. *Plasmid*, **2017**, 91, 68-75 3.3 14

227 General Physiological and Virulence Properties of the Pathogenic Clostridia **2016**, 7-12 3

226 The adherent abilities of Clostridium perfringens strains are critical for the pathogenesis of avian necrotic enteritis. *Veterinary Microbiology*, **2016**, 197, 53-61 3.3 33

225 Tohru Shimizu Memorial. *Anaerobe*, **2016**, 41, 3-4 2.8

224 Functional analysis of an feoB mutant in Clostridium perfringens strain 13. *Anaerobe*, **2016**, 41, 10-17 2.8 12

223 Analysis of the virulence-associated RevSR two-component signal transduction system of Clostridium perfringens. *International Journal of Medical Microbiology*, **2016**, 306, 429-42 3.7 2

222 NetB and necrotic enteritis: the hole movable story. *Avian Pathology*, **2016**, 45, 295-301 2.4 40

221	The NEAT Domain-Containing Proteins of <i>Clostridium perfringens</i> Bind Heme. <i>PLoS ONE</i> , <b>2016</b> , 11, e0162981	8	
220	CdtR Regulates TcdA and TcdB Production in <i>Clostridium difficile</i> . <i>PLoS Pathogens</i> , <b>2016</b> , 12, e1005758	7.6	37
219	TcpM: a novel relaxase that mediates transfer of large conjugative plasmids from <i>Clostridium perfringens</i> . <i>Molecular Microbiology</i> , <b>2016</b> , 99, 884-96	4.1	24
218	RNA-seq analysis of virR and revR mutants of <i>Clostridium perfringens</i> . <i>BMC Genomics</i> , <b>2016</b> , 17, 391	4.5	7
217	Animal models to study the pathogenesis of human and animal <i>Clostridium perfringens</i> infections. <i>Veterinary Microbiology</i> , <b>2015</b> , 179, 23-33	3.3	43
216	Plasmid partitioning systems of conjugative plasmids from <i>Clostridium perfringens</i> . <i>Plasmid</i> , <b>2015</b> , 80, 90-6	3.3	17
215	Molecular characterization and antimicrobial susceptibilities of <i>Clostridium difficile</i> clinical isolates from Victoria, Australia. <i>Anaerobe</i> , <b>2015</b> , 34, 80-3	2.8	6
214	Binding of <i>Clostridium perfringens</i> to collagen correlates with the ability to cause necrotic enteritis in chickens. <i>Veterinary Microbiology</i> , <b>2015</b> , 180, 299-303	3.3	46
213	Functional analysis of a bacitracin resistance determinant located on ICECp1, a novel Tn916-like element from a conjugative plasmid in <i>Clostridium perfringens</i> . <i>Antimicrobial Agents and Chemotherapy</i> , <b>2015</b> , 59, 6855-65	5.9	26
212	Genomic analyses of <i>Clostridium perfringens</i> isolates from five toxinotypes. <i>Research in Microbiology</i> , <b>2015</b> , 166, 255-63	4	45
211	Two novel membrane proteins, TcpD and TcpE, are essential for conjugative transfer of pCW3 in <i>Clostridium perfringens</i> . <i>Journal of Bacteriology</i> , <b>2015</b> , 197, 774-81	3.5	20
210	<i>Clostridium perfringens</i> type A-E toxin plasmids. <i>Research in Microbiology</i> , <b>2015</b> , 166, 264-79	4	38
209	Comparative neuropathology of ovine enterotoxemia produced by <i>Clostridium perfringens</i> type D wild-type strain CN1020 and its genetically modified derivatives. <i>Veterinary Pathology</i> , <b>2015</b> , 52, 465-75	2.8	18
208	<i>Clostridium perfringens</i> extracellular toxins and enzymes: 20 and counting. <i>Microbiology Australia</i> , <b>2015</b> , 36, 114	0.8	44
207	<i>Dichelobacter</i> <b>2015</b> , 1-7		
206	Solution structure and DNA binding of the catalytic domain of the large serine resolvase TnpX. <i>Journal of Molecular Recognition</i> , <b>2015</b> , 28, 316-24	2.6	1
205	NanR, a Transcriptional Regulator That Binds to the Promoters of Genes Involved in Sialic Acid Metabolism in the Anaerobic Pathogen <i>Clostridium perfringens</i> . <i>PLoS ONE</i> , <b>2015</b> , 10, e0133217	3.7	14
204	A two-component regulatory system modulates twitching motility in <i>Dichelobacter nodosus</i> . <i>Veterinary Microbiology</i> , <b>2015</b> , 179, 34-41	3.3	2

203	The pore-forming Etoxin from clostridium septicum activates the MAPK pathway in a Ras-c-Raf-dependent and independent manner. <i>Toxins</i> , <b>2015</b> , 7, 516-34	4.9	17
202	Defining the Roles of TcdA and TcdB in Localized Gastrointestinal Disease, Systemic Organ Damage, and the Host Response during Clostridium difficile Infections. <i>MBio</i> , <b>2015</b> , 6, e00551	7.8	161
201	Opioid analgesics stop the development of clostridial gas gangrene. <i>Journal of Infectious Diseases</i> , <b>2014</b> , 210, 483-92	7	6
200	Towards an understanding of the role of Clostridium perfringens toxins in human and animal disease. <i>Future Microbiology</i> , <b>2014</b> , 9, 361-77	2.9	231
199	Virulence Plasmids of Spore-Forming Bacteria. <i>Microbiology Spectrum</i> , <b>2014</b> , 2,	8.9	22
198	Detection of an en masse and reversible B- to A-DNA conformational transition in prokaryotes in response to desiccation. <i>Journal of the Royal Society Interface</i> , <b>2014</b> , 11, 20140454	4.1	42
197	Identification of a two-component signal transduction system that regulates maltose genes in Clostridium perfringens. <i>Anaerobe</i> , <b>2014</b> , 30, 199-204	2.8	2
196	Genomic evidence for a globally distributed, bimodal population in the ovine footrot pathogen Dichelobacter nodosus. <i>MBio</i> , <b>2014</b> , 5, e01821-14	7.8	24
195	Utility of the clostridial site-specific recombinase TnpX to clone toxic-product-encoding genes and selectively remove genomic DNA fragments. <i>Applied and Environmental Microbiology</i> , <b>2014</b> , 80, 3597-3603	4.8	8
194	Vaccination with recombinant NetB toxin partially protects broiler chickens from necrotic enteritis. <i>Veterinary Research</i> , <b>2013</b> , 44, 54	3.8	49
193	Structural and functional analysis of the pore-forming toxin NetB from Clostridium perfringens. <i>MBio</i> , <b>2013</b> , 4, e00019-13	7.8	56
192	Maternal immunization with vaccines containing recombinant NetB toxin partially protects progeny chickens from necrotic enteritis. <i>Veterinary Research</i> , <b>2013</b> , 44, 108	3.8	30
191	Epsilon toxin is essential for the virulence of Clostridium perfringens type D infection in sheep, goats, and mice. <i>Infection and Immunity</i> , <b>2013</b> , 81, 2405-14	3.7	69
190	Toxin plasmids of Clostridium perfringens. <i>Microbiology and Molecular Biology Reviews</i> , <b>2013</b> , 77, 208-33	13.2	166
189	Regulation of sialidase production in Clostridium perfringens by the orphan sensor histidine kinase ReeS. <i>PLoS ONE</i> , <b>2013</b> , 8, e73525	3.7	13
188	Spo0A differentially regulates toxin production in evolutionarily diverse strains of Clostridium difficile. <i>PLoS ONE</i> , <b>2013</b> , 8, e79666	3.7	55
187	Serine proteases and ovine footrot. <i>Microbiology Australia</i> , <b>2013</b> , 34, 37	0.8	
186	The peptidoglycan hydrolase TcpG is required for efficient conjugative transfer of pCW3 in Clostridium perfringens. <i>Plasmid</i> , <b>2012</b> , 67, 139-47	3.3	40

185	The role of toxin A and toxin B in the virulence of <i>Clostridium difficile</i> . <i>Trends in Microbiology</i> , <b>2012</b> , 20, 21-9	12.4	112
184	The conjugation protein TcpC from <i>Clostridium perfringens</i> is structurally related to the type IV secretion system protein VirB8 from Gram-negative bacteria. <i>Molecular Microbiology</i> , <b>2012</b> , 83, 275-88	4.1	63
183	The AprV5 subtilase is required for the optimal processing of all three extracellular serine proteases from <i>Dichelobacter nodosus</i> . <i>PLoS ONE</i> , <b>2012</b> , 7, e47932	3.7	11
182	Detection and diversity of a putative novel heterogeneous polymorphic proline-glycine repeat (Pgr) protein in the footrot pathogen <i>Dichelobacter nodosus</i> . <i>Veterinary Microbiology</i> , <b>2011</b> , 147, 358-66	3.3	13
181	The pathogenesis of ovine footrot. <i>Veterinary Microbiology</i> , <b>2011</b> , 153, 59-66	3.3	32
180	Necrotic enteritis-derived <i>Clostridium perfringens</i> strain with three closely related independently conjugative toxin and antibiotic resistance plasmids. <i>MBio</i> , <b>2011</b> , 2,	7.8	71
179	Epsilon-toxin production by <i>Clostridium perfringens</i> type D strain CN3718 is dependent upon the agr operon but not the VirS/VirR two-component regulatory system. <i>MBio</i> , <b>2011</b> , 2,	7.8	39
178	S1 pocket of a bacterially derived subtilisin-like protease underpins effective tissue destruction. <i>Journal of Biological Chemistry</i> , <b>2011</b> , 286, 42180-42187	5.4	16
177	TcsL is an essential virulence factor in <i>Clostridium sordellii</i> ATCC 9714. <i>Infection and Immunity</i> , <b>2011</b> , 79, 1025-32	3.7	41
176	Regulation of virulence by the RevR response regulator in <i>Clostridium perfringens</i> . <i>Infection and Immunity</i> , <b>2011</b> , 79, 2145-53	3.7	28
175	The anti-sigma factor TcdC modulates hypervirulence in an epidemic BI/NAP1/027 clinical isolate of <i>Clostridium difficile</i> . <i>PLoS Pathogens</i> , <b>2011</b> , 7, e1002317	7.6	109
174	The cysteine protease Eclostripain is not essential for the pathogenesis of <i>Clostridium perfringens</i> -mediated myonecrosis. <i>PLoS ONE</i> , <b>2011</b> , 6, e22762	3.7	11
173	Association between avian necrotic enteritis and <i>Clostridium perfringens</i> strains expressing NetB toxin. <i>Veterinary Research</i> , <b>2010</b> , 41, 21	3.8	106
172	The role of toxin A and toxin B in <i>Clostridium difficile</i> -associated disease: Past and present perspectives. <i>Gut Microbes</i> , <b>2010</b> , 1, 58-64	8.8	72
171	Novel use of tryptose sulfite cycloserine egg yolk agar for isolation of <i>Clostridium perfringens</i> during an outbreak of necrotizing enterocolitis in a neonatal unit. <i>Journal of Clinical Microbiology</i> , <b>2010</b> , 48, 4263-5	9.7	11
170	The VirSR two-component signal transduction system regulates NetB toxin production in <i>Clostridium perfringens</i> . <i>Infection and Immunity</i> , <b>2010</b> , 78, 3064-72	3.7	79
169	The subtilisin-like protease AprV2 is required for virulence and uses a novel disulphide-tethered exosite to bind substrates. <i>PLoS Pathogens</i> , <b>2010</b> , 6, e1001210	7.6	66
168	NetB, a pore-forming toxin from necrotic enteritis strains of <i>Clostridium perfringens</i> . <i>Toxins</i> , <b>2010</b> , 2, 1913-27	4.9	80

167	Crystallization of the virulent and benign subtilisin-like proteases from the ovine footrot pathogen <i>Dichelobacter nodosus</i> . <i>Acta Crystallographica Section F: Structural Biology Communications</i> , <b>2010</b> , 66, 289-93		3
166	Methods for gene cloning and targeted mutagenesis. <i>Methods in Molecular Biology</i> , <b>2010</b> , 646, 183-201	1.4	1
165	Functional analysis of the VirSR phosphorelay from <i>Clostridium perfringens</i> . <i>PLoS ONE</i> , <b>2009</b> , 4, e5849	3.7	28
164	The NanI and NanJ sialidases of <i>Clostridium perfringens</i> are not essential for virulence. <i>Infection and Immunity</i> , <b>2009</b> , 77, 4421-8	3.7	40
163	Development and application of new mouse models to study the pathogenesis of <i>Clostridium perfringens</i> type C Enterotoxemias. <i>Infection and Immunity</i> , <b>2009</b> , 77, 5291-9	3.7	47
162	The putative coupling protein TcpA interacts with other pCW3-encoded proteins to form an essential part of the conjugation complex. <i>Journal of Bacteriology</i> , <b>2009</b> , 191, 2926-33	3.5	38
161	Programmed cellular necrosis mediated by the pore-forming alpha-toxin from <i>Clostridium septicum</i> . <i>PLoS Pathogens</i> , <b>2009</b> , 5, e1000516	7.6	91
160	tISCpe8, an IS1595-family lincomycin resistance element located on a conjugative plasmid in <i>Clostridium perfringens</i> . <i>Journal of Bacteriology</i> , <b>2009</b> , 191, 6345-51	3.5	22
159	Pore-forming activity of alpha-toxin is essential for <i>clostridium septicum</i> -mediated myonecrosis. <i>Infection and Immunity</i> , <b>2009</b> , 77, 943-51	3.7	35
158	Identification of a leukotoxin sequence from <i>Fusobacterium equinum</i> . <i>Veterinary Microbiology</i> , <b>2009</b> , 133, 394-5	3.3	3
157	Toxin B is essential for virulence of <i>Clostridium difficile</i> . <i>Nature</i> , <b>2009</b> , 458, 1176-9	50.4	542
156	Cross-complementation of <i>Clostridium perfringens</i> PLC and <i>Clostridium septicum</i> alpha-toxin mutants reveals PLC is sufficient to mediate gas gangrene. <i>Microbes and Infection</i> , <b>2009</b> , 11, 413-8	9.3	19
155	Rethinking our understanding of the pathogenesis of necrotic enteritis in chickens. <i>Trends in Microbiology</i> , <b>2009</b> , 17, 32-6	12.4	213
154	Revised nomenclature for transposable genetic elements. <i>Plasmid</i> , <b>2008</b> , 60, 167-73	3.3	143
153	Influence of gastric acid on susceptibility to infection with ingested bacterial pathogens. <i>Infection and Immunity</i> , <b>2008</b> , 76, 639-45	3.7	115
152	NetB, a new toxin that is associated with avian necrotic enteritis caused by <i>Clostridium perfringens</i> . <i>PLoS Pathogens</i> , <b>2008</b> , 4, e26	7.6	405
151	Molecular and cellular basis of microvascular perfusion deficits induced by <i>Clostridium perfringens</i> and <i>Clostridium septicum</i> . <i>PLoS Pathogens</i> , <b>2008</b> , 4, e1000045	7.6	62
150	Functional characterization and localization of the TcpH conjugation protein from <i>Clostridium perfringens</i> . <i>Journal of Bacteriology</i> , <b>2008</b> , 190, 5075-86	3.5	38

149	Twitching motility is essential for virulence in <i>Dichelobacter nodosus</i> . <i>Journal of Bacteriology</i> , <b>2008</b> , 190, 3323-35	3.5	45
148	Genome sequence of the saprophyte <i>Leptospira biflexa</i> provides insights into the evolution of <i>Leptospira</i> and the pathogenesis of leptospirosis. <i>PLoS ONE</i> , <b>2008</b> , 3, e1607	3.7	225
147	Beta toxin is essential for the intestinal virulence of <i>Clostridium perfringens</i> type C disease isolate CN3685 in a rabbit ileal loop model. <i>Molecular Microbiology</i> , <b>2008</b> , 67, 15-30	4.1	139
146	Isolation of the Bacteriophage DinoHI from <i>Dichelobacter nodosus</i> and its Interactions with other Integrated Genetic Elements. <i>Open Microbiology Journal</i> , <b>2008</b> , 2, 1-9	0.8	7
145	Genome sequence and identification of candidate vaccine antigens from the animal pathogen <i>Dichelobacter nodosus</i> . <i>Nature Biotechnology</i> , <b>2007</b> , 25, 569-75	44.5	58
144	Type IV fimbrial biogenesis is required for protease secretion and natural transformation in <i>Dichelobacter nodosus</i> . <i>Journal of Bacteriology</i> , <b>2007</b> , 189, 5022-33	3.5	52
143	Epsilon-toxin plasmids of <i>Clostridium perfringens</i> type D are conjugative. <i>Journal of Bacteriology</i> , <b>2007</b> , 189, 7531-8	3.5	62
142	Binary toxin production in <i>Clostridium difficile</i> is regulated by CdtR, a LytTR family response regulator. <i>Journal of Bacteriology</i> , <b>2007</b> , 189, 7290-301	3.5	101
141	TcpA, an FtsK/SpoIIIE homolog, is essential for transfer of the conjugative plasmid pCW3 in <i>Clostridium perfringens</i> . <i>Journal of Bacteriology</i> , <b>2007</b> , 189, 7782-90	3.5	52
140	Both epsilon-toxin and beta-toxin are important for the lethal properties of <i>Clostridium perfringens</i> type B isolates in the mouse intravenous injection model. <i>Infection and Immunity</i> , <b>2007</b> , 75, 1443-52	3.7	50
139	Development and application of an oral challenge mouse model for studying <i>Clostridium perfringens</i> type D infection. <i>Infection and Immunity</i> , <b>2007</b> , 75, 4282-8	3.7	32
138	Comparison of the RpoH-dependent regulon and general stress response in <i>Neisseria gonorrhoeae</i> . <i>Journal of Bacteriology</i> , <b>2006</b> , 188, 4769-76	3.5	25
137	Functional identification of conjugation and replication regions of the tetracycline resistance plasmid pCW3 from <i>Clostridium perfringens</i> . <i>Journal of Bacteriology</i> , <b>2006</b> , 188, 4942-51	3.5	89
136	Dissecting the contributions of <i>Clostridium perfringens</i> type C toxins to lethality in the mouse intravenous injection model. <i>Infection and Immunity</i> , <b>2006</b> , 74, 5200-10	3.7	81
135	Regulation of type IV fimbrial biogenesis in <i>Dichelobacter nodosus</i> . <i>Journal of Bacteriology</i> , <b>2006</b> , 188, 4801-11	3.5	22
134	Ecf, an alternative sigma factor from <i>Neisseria gonorrhoeae</i> , controls expression of msrAB, which encodes methionine sulfoxide reductase. <i>Journal of Bacteriology</i> , <b>2006</b> , 188, 3463-9	3.5	32
133	Genome reduction in <i>Leptospira borgpetersenii</i> reflects limited transmission potential. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2006</b> , 103, 14560-5	11.5	250
132	Skewed genomic variability in strains of the toxigenic bacterial pathogen, <i>Clostridium perfringens</i> . <i>Genome Research</i> , <b>2006</b> , 16, 1031-40	9.7	250

131	Alpha-toxin of <i>Clostridium perfringens</i> is not an essential virulence factor in necrotic enteritis in chickens. <i>Infection and Immunity</i> , <b>2006</b> , 74, 6496-500	3.7	185
130	Two distinct regions of the large serine recombinase TnpX are required for DNA binding and biological function. <i>Molecular Microbiology</i> , <b>2006</b> , 60, 591-601	4.1	12
129	Construction and analysis of chromosomal <i>Clostridium difficile</i> mutants. <i>Molecular Microbiology</i> , <b>2006</b> , 61, 1335-51	4.1	134
128	Regulation systems of toxin expression <b>2006</b> , 64-82		
127	<i>Clostridium perfringens</i> and Histotoxic Disease <b>2006</b> , 753-770		12
126	Construction of an alpha toxin gene knockout mutant of <i>Clostridium perfringens</i> type A by use of a mobile group II intron. <i>Applied and Environmental Microbiology</i> , <b>2005</b> , 71, 7542-7	4.8	116
125	Epsilon-toxin is required for most <i>Clostridium perfringens</i> type D vegetative culture supernatants to cause lethality in the mouse intravenous injection model. <i>Infection and Immunity</i> , <b>2005</b> , 73, 7413-21	3.7	56
124	Clostridial Toxins Involved in Human Enteric and Histotoxic Infections <b>2005</b> , 169-209		6
123	The alpha-toxin of <i>Clostridium septicum</i> is essential for virulence. <i>Molecular Microbiology</i> , <b>2005</b> , 57, 1357-66	4.6	94
122	Identification of a <i>Dichelobacter nodosus</i> ferric uptake regulator and determination of its regulatory targets. <i>Journal of Bacteriology</i> , <b>2005</b> , 187, 366-75	3.5	17
121	Identification of the structural and functional domains of the large serine recombinase TnpX from <i>Clostridium perfringens</i> . <i>Journal of Biological Chemistry</i> , <b>2005</b> , 280, 2503-11	5.4	18
120	Revised nomenclature of <i>Clostridium difficile</i> toxins and associated genes. <i>Journal of Medical Microbiology</i> , <b>2005</b> , 54, 113-117	3.2	76
119	Highly conserved alpha-toxin sequences of avian isolates of <i>Clostridium perfringens</i> . <i>Journal of Clinical Microbiology</i> , <b>2004</b> , 42, 1345-7	9.7	39
118	The spatial organization of the VirR boxes is critical for VirR-mediated expression of the perfringolysin O gene, pfoA, from <i>Clostridium perfringens</i> . <i>Journal of Bacteriology</i> , <b>2004</b> , 186, 3321-30	3.5	35
117	The <i>Clostridium perfringens</i> TetA(P) efflux protein contains a functional variant of the Motif A region found in major facilitator superfamily transport proteins. <i>Microbiology (United Kingdom)</i> , <b>2004</b> , 150, 127-134	2.9	28
116	The large resolvase TnpX is the only transposon-encoded protein required for transposition of the Tn4451/3 family of integrative mobilizable elements. <i>Molecular Microbiology</i> , <b>2004</b> , 51, 1787-800	4.1	36
115	DNA binding properties of TnpX indicate that different synapses are formed in the excision and integration of the Tn4451 family. <i>Molecular Microbiology</i> , <b>2004</b> , 53, 1195-207	4.1	23
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108	Identification of essential residues in the Erm(B) rRNA methyltransferase of <i>Clostridium perfringens</i> . <i>Antimicrobial Agents and Chemotherapy</i> , <b>2002</b> , 46, 1253-61	5.9	20
107	Environmental response and autoregulation of <i>Clostridium difficile</i> TxeR, a sigma factor for toxin gene expression. <i>Journal of Bacteriology</i> , <b>2002</b> , 184, 5971-8	3.5	104
106	The FxRxHrS motif: a conserved region essential for DNA binding of the VirR response regulator from <i>Clostridium perfringens</i> . <i>Journal of Molecular Biology</i> , <b>2002</b> , 322, 997-1011	6.5	21
105	Induction of pCW3-encoded tetracycline resistance in <i>Clostridium perfringens</i> involves a host-encoded factor. <i>Plasmid</i> , <b>2001</b> , 46, 229-32	3.3	8
104	Enterotoxin plasmid from <i>Clostridium perfringens</i> is conjugative. <i>Infection and Immunity</i> , <b>2001</b> , 69, 3483-37		96
103	The type IV fimbrial subunit gene (fimA) of <i>Dichelobacter nodosus</i> is essential for virulence, protease secretion, and natural competence. <i>Journal of Bacteriology</i> , <b>2001</b> , 183, 4451-8	3.5	85
102	Transcriptional analysis of the tet(P) operon from <i>Clostridium perfringens</i> . <i>Journal of Bacteriology</i> , <b>2001</b> , 183, 7110-9	3.5	28
101	Synergistic effects of alpha-toxin and perfringolysin O in <i>Clostridium perfringens</i> -mediated gas gangrene. <i>Infection and Immunity</i> , <b>2001</b> , 69, 7904-10	3.7	141
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99	Comparison of Tn5397 from <i>Clostridium difficile</i> , Tn916 from <i>Enterococcus faecalis</i> and the CW459tet(M) element from <i>Clostridium perfringens</i> shows that they have similar conjugation regions but different insertion and excision modules. <i>Microbiology (United Kingdom)</i> , <b>2001</b> , 147, 1243-1251	2.9	80
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92	Glutamate residues in the putative transmembrane region are required for the function of the VirS sensor histidine kinase from <i>Clostridium perfringens</i> . <i>Microbiology (United Kingdom)</i> , <b>2000</b> , 146 ( Pt 2), 517-525	2.9	13
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