Steven James Norris

List of Publications by Citations

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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.

131 papers

7,469 citations

49 h-index

84 g-index

140 ext. papers

8,406 ext. citations

6.6 avg, IF

5.75 L-index

#	Paper	IF	Citations
131	Complete genome sequence of Treponema pallidum, the syphilis spirochete. <i>Science</i> , 1998 , 281, 375-88	33.3	756
130	Antigenic variation in Lyme disease borreliae by promiscuous recombination of VMP-like sequence cassettes. <i>Cell</i> , 1997 , 89, 275-85	56.2	530
129	Correlation between plasmid content and infectivity in Borrelia burgdorferi. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000 , 97, 13865-70	11.5	376
128	A plasmid-encoded nicotinamidase (PncA) is essential for infectivity of Borrelia burgdorferi in a mammalian host. <i>Molecular Microbiology</i> , 2003 , 48, 753-64	4.1	218
127	Comparison of the genome of the oral pathogen Treponema denticola with other spirochete genomes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004 , 101, 5646-51	11.5	215
126	Genetic variation of the Borrelia burgdorferi gene vlsE involves cassette-specific, segmental gene conversion. <i>Infection and Immunity</i> , 1998 , 66, 3698-704	3.7	185
125	Disruption of the genes encoding antigen 85A and antigen 85B of Mycobacterium tuberculosis H37Rv: effect on growth in culture and in macrophages. <i>Infection and Immunity</i> , 2000 , 68, 767-78	3.7	175
124	Adherence of Borrelia burgdorferi to the proteoglycan decorin. <i>Infection and Immunity</i> , 1995 , 63, 3467-	73 .7	175
123	Low-passage-associated proteins of Borrelia burgdorferi B31: characterization and molecular cloning of OspD, a surface-exposed, plasmid-encoded lipoprotein. <i>Infection and Immunity</i> , 1992 , 60, 466	5 2-7 2	173
122	Kinetics and in vivo induction of genetic variation of vlsE in Borrelia burgdorferi. <i>Infection and Immunity</i> , 1998 , 66, 3689-97	3.7	146
121	Intact flagellar motor of Borrelia burgdorferi revealed by cryo-electron tomography: evidence for stator ring curvature and rotor/C-ring assembly flexion. <i>Journal of Bacteriology</i> , 2009 , 191, 5026-36	3.5	132
120	BBE02 disruption mutants of Borrelia burgdorferi B31 have a highly transformable, infectious phenotype. <i>Infection and Immunity</i> , 2004 , 72, 7147-54	3.7	132
119	Human antibody responses to VlsE antigenic variation protein of Borrelia burgdorferi. <i>Journal of Clinical Microbiology</i> , 1999 , 37, 3997-4004	9.7	106
118	Crystal structure of Lyme disease variable surface antigen VlsE of Borrelia burgdorferi. <i>Journal of Biological Chemistry</i> , 2002 , 277, 21691-6	5.4	104
117	High- and low-infectivity phenotypes of clonal populations of in vitro-cultured Borrelia burgdorferi. <i>Infection and Immunity</i> , 1995 , 63, 2206-12	3.7	103
116	Molecular architecture of the bacterial flagellar motor in cells. <i>Biochemistry</i> , 2014 , 53, 4323-33	3.2	93
115	Characterization of a manganese-dependent regulatory protein, TroR, from Treponema pallidum. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1999 , 96, 10887-92	11.5	93

114	Analysis of an ordered, comprehensive STM mutant library in infectious Borrelia burgdorferi: insights into the genes required for mouse infectivity. <i>PLoS ONE</i> , 2012 , 7, e47532	3.7	91
113	Cellular architecture of Treponema pallidum: novel flagellum, periplasmic cone, and cell envelope as revealed by cryo electron tomography. <i>Journal of Molecular Biology</i> , 2010 , 403, 546-61	6.5	90
112	Purification of Treponema pallidum, Nichols strain, by Percoll density gradient centrifugation. <i>Sexually Transmitted Diseases</i> , 1984 , 11, 275-86	2.4	87
111	Detailed analysis of sequence changes occurring during vlsE antigenic variation in the mouse model of Borrelia burgdorferi infection. <i>PLoS Pathogens</i> , 2009 , 5, e1000293	7.6	86
110	Whole genome sequences of three Treponema pallidum ssp. pertenue strains: yaws and syphilis treponemes differ in less than 0.2% of the genome sequence. <i>PLoS Neglected Tropical Diseases</i> , 2012 , 6, e1471	4.8	86
109	A novel Treponema pallidum antigen, TP0136, is an outer membrane protein that binds human fibronectin. <i>Infection and Immunity</i> , 2008 , 76, 1848-57	3.7	83
108	Origin of modern syphilis and emergence of a pandemic Treponema pallidum cluster. <i>Nature Microbiology</i> , 2016 , 2, 16245	26.6	81
107	Antigenic variation with a twistthe Borrelia story. <i>Molecular Microbiology</i> , 2006 , 60, 1319-22	4.1	80
106	Decreased electroporation efficiency in Borrelia burgdorferi containing linear plasmids lp25 and lp56: impact on transformation of infectious B. burgdorferi. <i>Infection and Immunity</i> , 2002 , 70, 4798-804	3.7	79
105	Long-Term Culture of the Syphilis Spirochete subsp <i>MBio</i> , 2018 , 9,	7.8	79
104	Cryoelectron tomography reveals the sequential assembly of bacterial flagella in Borrelia		77
	burgdorferi. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 14390-5	11.5	//
103		11.5 4·5	76
103	110, 14390-5 Genetic diversity in Treponema pallidum: implications for pathogenesis, evolution and molecular		
	Genetic diversity in Treponema pallidum: implications for pathogenesis, evolution and molecular diagnostics of syphilis and yaws. <i>Infection, Genetics and Evolution</i> , 2012 , 12, 191-202 A family of surface-exposed proteins of 20 kilodaltons in the genus Borrelia. <i>Infection and Immunity</i> , 1994 , 62, 2792-9 Role of acetyl-phosphate in activation of the Rrp2-RpoN-RpoS pathway in Borrelia burgdorferi	4.5	76
102	Genetic diversity in Treponema pallidum: implications for pathogenesis, evolution and molecular diagnostics of syphilis and yaws. <i>Infection, Genetics and Evolution</i> , 2012 , 12, 191-202 A family of surface-exposed proteins of 20 kilodaltons in the genus Borrelia. <i>Infection and Immunity</i> , 1994 , 62, 2792-9 Role of acetyl-phosphate in activation of the Rrp2-RpoN-RpoS pathway in Borrelia burgdorferi.	4·5 3·7	76 76
102	Genetic diversity in Treponema pallidum: implications for pathogenesis, evolution and molecular diagnostics of syphilis and yaws. <i>Infection, Genetics and Evolution</i> , 2012 , 12, 191-202 A family of surface-exposed proteins of 20 kilodaltons in the genus Borrelia. <i>Infection and Immunity</i> , 1994 , 62, 2792-9 Role of acetyl-phosphate in activation of the Rrp2-RpoN-RpoS pathway in Borrelia burgdorferi. <i>PLoS Pathogens</i> , 2010 , 6, e1001104	4·5 3·7 7.6	76 76 75
102	Genetic diversity in Treponema pallidum: implications for pathogenesis, evolution and molecular diagnostics of syphilis and yaws. <i>Infection, Genetics and Evolution</i> , 2012 , 12, 191-202 A family of surface-exposed proteins of 20 kilodaltons in the genus Borrelia. <i>Infection and Immunity</i> , 1994 , 62, 2792-9 Role of acetyl-phosphate in activation of the Rrp2-RpoN-RpoS pathway in Borrelia burgdorferi. <i>PLoS Pathogens</i> , 2010 , 6, e1001104 Toxin synthesis by Clostridium difficile is regulated through quorum signaling. <i>MBio</i> , 2015 , 6, e02569 Analysis of Borrelia burgdorferi vlsE gene expression and recombination in the tick vector. <i>Infection</i>	4·5 3·7 7·6 7.8	76 76 75 74

96	Identity of Treponema pallidum subsp. pallidum polypeptides: Correlation of sodium dodecyl sulfate-polyacrylamide gel electrophoresis results from different laboratories. <i>Electrophoresis</i> , 1987 , 8, 77-92	3.6	66
95	Linear and circular plasmid content in Borrelia burgdorferi clinical isolates. <i>Infection and Immunity</i> , 2003 , 71, 3699-706	3.7	65
94	vls Antigenic Variation Systems of Lyme Disease Borrelia: Eluding Host Immunity through both Random, Segmental Gene Conversion and Framework Heterogeneity. <i>Microbiology Spectrum</i> , 2014 , 2,	8.9	63
93	Central role of the Holliday junction helicase RuvAB in vlsE recombination and infectivity of Borrelia burgdorferi. <i>PLoS Pathogens</i> , 2009 , 5, e1000679	7.6	63
92	Effects of vlsE complementation on the infectivity of Borrelia burgdorferi lacking the linear plasmid lp28-1. <i>Infection and Immunity</i> , 2004 , 72, 6577-85	3.7	63
91	Identification and transcriptional analysis of a Treponema pallidum operon encoding a putative ABC transport system, an iron-activated repressor protein homolog, and a glycolytic pathway enzyme homolog. <i>Gene</i> , 1997 , 197, 47-64	3.8	62
90	Relationship of Treponema denticola periplasmic flagella to irregular cell morphology. <i>Journal of Bacteriology</i> , 1997 , 179, 1628-35	3.5	61
89	Transcriptome of Treponema pallidum: gene expression profile during experimental rabbit infection. <i>Journal of Bacteriology</i> , 2005 , 187, 1866-74	3.5	57
88	Conversion of a linear to a circular plasmid in the relapsing fever agent Borrelia hermsii. <i>Journal of Bacteriology</i> , 1996 , 178, 793-800	3.5	57
87	Complete genome sequence of Treponema pallidum ssp. pallidum strain SS14 determined with oligonucleotide arrays. <i>BMC Microbiology</i> , 2008 , 8, 76	4.5	55
86	Understanding barriers to Borrelia burgdorferi dissemination during infection using massively parallel sequencing. <i>Infection and Immunity</i> , 2013 , 81, 2347-57	3.7	52
85	Genome scale identification of Treponema pallidum antigens. <i>Infection and Immunity</i> , 2005 , 73, 4445-50	3.7	52
84	Complete genome sequence of Treponema paraluiscuniculi, strain Cuniculi A: the loss of infectivity to humans is associated with genome decay. <i>PLoS ONE</i> , 2011 , 6, e20415	3.7	51
83	Reactivity of antibodies from syphilis patients to a protein array representing the Treponema pallidum proteome. <i>Journal of Clinical Microbiology</i> , 2006 , 44, 888-91	9.7	50
82	Genome analysis of Treponema pallidum subsp. pallidum and subsp. pertenue strains: most of the genetic differences are localized in six regions. <i>PLoS ONE</i> , 2010 , 5, e15713	3.7	48
81	Biology of Treponema pallidum: correlation of functional activities with genome sequence data. <i>Journal of Molecular Microbiology and Biotechnology</i> , 2001 , 3, 37-62	0.9	48
80	A mutant of Mycobacterium tuberculosis H37Rv that lacks expression of antigen 85A is attenuated in mice but retains vaccinogenic potential. <i>Infection and Immunity</i> , 2004 , 72, 7084-95	3.7	46
79	Effect of complement component C3 deficiency on experimental Lyme borreliosis in mice. <i>Infection and Immunity</i> , 2003 , 71, 4432-40	3.7	46

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78	The genome of Treponema pallidum: new light on the agent of syphilis. <i>FEMS Microbiology Reviews</i> , 1998 , 22, 323-32	15.1	43	
77	Mycobacterial protein HbhA binds human complement component C3. <i>Infection and Immunity</i> , 2001 , 69, 7501-11	3.7	43	
76	In vitro cultivation of Treponema pallidum: independent confirmation. <i>Infection and Immunity</i> , 1982 , 36, 437-9	3.7	42	
75	Transcriptional regulation of the Borrelia burgdorferi antigenically variable VlsE surface protein. Journal of Bacteriology, 2006 , 188, 4879-89	3.5	41	
74	Comparative reactivity of human sera to recombinant VlsE and other Borrelia burgdorferi antigens in class-specific enzyme-linked immunosorbent assays for Lyme borreliosis. <i>Journal of Medical Microbiology</i> , 2002 , 51, 649-655	3.2	40	
73	Characterization and serologic analysis of the Treponema pallidum proteome. <i>Infection and Immunity</i> , 2010 , 78, 2631-43	3.7	39	
72	Identification of potential virulence determinants by Himar1 transposition of infectious Borrelia burgdorferi B31. <i>Infection and Immunity</i> , 2006 , 74, 6690-9	3.7	39	
71	Systematic cloning of Treponema pallidum open reading frames for protein expression and antigen discovery. <i>Genome Research</i> , 2003 , 13, 1665-74	9.7	38	
70	Genome differences between Treponema pallidum subsp. pallidum strain Nichols and T. paraluiscuniculi strain Cuniculi A. <i>Infection and Immunity</i> , 2007 , 75, 5859-66	3.7	36	
69	Extensive interplasmidic duplications change the virulence phenotype of the relapsing fever agent Borrelia turicatae. <i>Molecular Microbiology</i> , 1999 , 34, 1120-32	4.1	35	
68	Molecular mechanism for rotational switching of the bacterial flagellar motor. <i>Nature Structural and Molecular Biology</i> , 2020 , 27, 1041-1047	17.6	34	
67	Global Tn-seq analysis of carbohydrate utilization and vertebrate infectivity of Borrelia burgdorferi. <i>Molecular Microbiology</i> , 2016 , 101, 1003-23	4.1	33	
66	Characterization of the vls antigenic variation loci of the Lyme disease spirochaetes Borrelia garinii Ip90 and Borrelia afzelii ACAI. <i>Molecular Microbiology</i> , 2003 , 47, 1407-17	4.1	31	
65	Conservation and heterogeneity of vlsE among human and tick isolates of Borrelia burgdorferi. <i>Infection and Immunity</i> , 2000 , 68, 1714-8	3.7	29	
64	Physical map of the genome of Treponema pallidum subsp. pallidum (Nichols). <i>Journal of Bacteriology</i> , 1995 , 177, 1797-804	3.5	29	
63	Molecular studies in Treponema pallidum evolution: toward clarity?. <i>PLoS Neglected Tropical Diseases</i> , 2008 , 2, e184	4.8	28	
62	Mutations in the Borrelia burgdorferi Flagellar Type III Secretion System Genes fliH and flil Profoundly Affect Spirochete Flagellar Assembly, Morphology, Motility, Structure, and Cell Division. <i>MBio</i> , 2015 , 6, e00579-15	7.8	27	
61	Transposon mutagenesis as an approach to improved understanding of Borrelia pathogenesis and biology. <i>Frontiers in Cellular and Infection Microbiology</i> , 2014 , 4, 63	5.9	27	

60	Structural insights into flagellar stator-rotor interactions. <i>ELife</i> , 2019 , 8,	8.9	27
59	High-throughput plasmid content analysis of Borrelia burgdorferi B31 by using Luminex multiplex technology. <i>Applied and Environmental Microbiology</i> , 2011 , 77, 1483-92	4.8	26
58	Decreased infectivity despite unaltered C3 binding by a DeltahbhA mutant of Mycobacterium tuberculosis. <i>Infection and Immunity</i> , 2002 , 70, 6751-60	3.7	26
57	From microbial genome sequence to applications. <i>Research in Microbiology</i> , 2000 , 151, 151-8	4	26
56	Phosphoenolpyruvate Phosphotransferase System Components Modulate Gene Transcription and Virulence of Borrelia burgdorferi. <i>Infection and Immunity</i> , 2015 , 84, 754-64	3.7	24
55	Characterization of the cytoplasmic filament protein gene (cfpA) of Treponema pallidum subsp. pallidum. <i>Journal of Bacteriology</i> , 1996 , 178, 3177-87	3.5	24
54	Genome structure of spirochetes. <i>Research in Microbiology</i> , 1992 , 143, 615-21	4	24
53	In vitro culture system to determine MICs and MBCs of antimicrobial agents against Treponema pallidum subsp. pallidum (Nichols strain). <i>Antimicrobial Agents and Chemotherapy</i> , 1988 , 32, 68-74	5.9	24
52	Influence of oxygen tension, sulfhydryl compounds, and serum on the motility and virulence of Treponema pallidum (Nichols strain) in a cell-free system. <i>Infection and Immunity</i> , 1978 , 22, 689-97	3.7	23
51	A high-throughput genetic screen identifies previously uncharacterized Borrelia burgdorferi genes important for resistance against reactive oxygen and nitrogen species. <i>PLoS Pathogens</i> , 2017 , 13, e100	6225	23
50	BAC library of T. pallidum DNA in E. coli. <i>Genome Research</i> , 2002 , 12, 515-22	9.7	21
49	Function of the Borrelia burgdorferi FtsH Homolog Is Essential for Viability both In Vitro and In Vivo and Independent of HflK/C. <i>MBio</i> , 2016 , 7, e00404-16	7.8	21
48	Antigenicity and recombination of VlsE, the antigenic variation protein of Borrelia burgdorferi, in rabbits, a host putatively resistant to long-term infection with this spirochete. <i>FEMS Immunology and Medical Microbiology</i> , 2007 , 50, 421-9		20
47	Isolation and characterization of a Treponema pallidum major 60-kilodalton protein resembling the groEL protein of Escherichia coli. <i>Journal of Bacteriology</i> , 1990 , 172, 2862-70	3.5	20
46	The Nucleotide Excision Repair Pathway Protects Borrelia burgdorferi from Nitrosative Stress in Ixodes scapularis Ticks. <i>Frontiers in Microbiology</i> , 2016 , 7, 1397	5.7	19
45	The Microaerophilic Nature of Treponema pallidurn. Sexually Transmitted Diseases, 1982, 9, 1-8	2.4	18
44	Peaceful coexistence amongst Borrelia plasmids: getting by with a little help from their friends?. <i>Plasmid</i> , 2013 , 70, 161-7	3.3	17
43	Genome-wide screen identifies novel genes required for Borrelia burgdorferi survival in its Ixodes tick vector. <i>PLoS Pathogens</i> , 2019 , 15, e1007644	7.6	16

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42	Infectivity of the highly transformable BBE02- lp56- mutant of Borrelia burgdorferi, the Lyme disease spirochete, via ticks. <i>Infection and Immunity</i> , 2006 , 74, 3678-81	3.7	16
41	Cryo-electron tomography of periplasmic flagella in Borrelia burgdorferi reveals a distinct cytoplasmic ATPase complex. <i>PLoS Biology</i> , 2018 , 16, e3000050	9.7	16
40	Interaction of spirochetes with the host. <i>Research in Microbiology</i> , 1992 , 143, 629-39	4	15
39	A Retrospective Study on Genetic Heterogeneity within Treponema Strains: Subpopulations Are Genetically Distinct in a Limited Number of Positions. <i>PLoS Neglected Tropical Diseases</i> , 2015 , 9, e00041	16 ⁸	15
38	Serum antibodies to Borrelia burgdorferi, Anaplasma phagocytophilum, and Babesia microti in recaptured white-footed mice. <i>Journal of Wildlife Diseases</i> , 2013 , 49, 294-302	1.3	14
37	Analysis of the intergenic sequences provided by Feria-Arroyo et al. does not support the claim of high Borrelia burgdorferi tick infection rates in Texas and northeastern Mexico. <i>Parasites and Vectors</i> , 2014 , 7, 467	4	12
36	Specific Th1 cell lines that confer protective immunity against experimental Borrelia burgdorferi infection in mice. <i>Journal of Leukocyte Biology</i> , 1998 , 63, 542-9	6.5	12
35	Serum requirement for the multiplication of Treponema pallidum in a tissue-culture system: association of growth-promoting activity with the protein fraction. <i>Sexually Transmitted Diseases</i> , 1986 , 13, 207-13	2.4	12
34	The Borrelia burgdorferi Glycosaminoglycan Binding Protein Bgp in the B31 Strain Is Not Essential for Infectivity despite Facilitating Adherence and Tissue Colonization. <i>Infection and Immunity</i> , 2018 , 86,	3.7	12
33	The genome sequence of Treponema pallidum, the syphilis spirochete: will clinicians benefit?. <i>Current Opinion in Infectious Diseases</i> , 2000 , 13, 29-36	5.4	11
32	Lyme Disease Pathogenesis. Current Issues in Molecular Biology, 2021, 42, 473-518	2.9	11
31	Enhanced Protective Immunogenicity of Homodimeric Borrelia burgdorferi Outer Surface Protein C. <i>Vaccine Journal</i> , 2017 , 24,		10
30	The dynamic proteome of Lyme disease Borrelia. <i>Genome Biology</i> , 2006 , 7, 209	18.3	10
29	Long-term incorporation of tritiated adenine into deoxyribonucleic acid and ribonucleic acid by Treponema pallidum (Nichols strain). <i>Infection and Immunity</i> , 1980 , 29, 1040-9	3.7	10
28	The thermophilic, homohexameric aminopeptidase of Borrelia burgdorferi is a member of the M29 family of metallopeptidases. <i>Infection and Immunity</i> , 2005 , 73, 2253-61	3.7	9
27	The intergenic small non-coding RNA ittA is required for optimal infectivity and tissue tropism in Borrelia burgdorferi. <i>PLoS Pathogens</i> , 2020 , 16, e1008423	7.6	8
26	The Genus Treponema 2006 , 211-234		8
25	How do lyme borrelia organisms cause disease? The quest for virulence determinants(). <i>The Open Neurology Journal</i> , 2012 , 6, 119-23	0.4	7

24	In Vitro Cultivation of the Syphilis Spirochete Treponema pallidum. <i>Current Protocols</i> , 2021 , 1, e44		7
23	Hiding in Plain Sight: Colonic Spirochetosis in Humans. <i>Journal of Bacteriology</i> , 2019 , 201,	3.5	6
22	Response to Esteve-Gassent et al.: flaB sequences obtained from Texas PCR products are identical to the positive control strain Borrelia burgdorferi B31. <i>Parasites and Vectors</i> , 2015 , 8, 310	4	6
21	Treponema 2015 , 1-42		5
20	Out of the woods: the remarkable genomes of the genus Borrelia. <i>Journal of Bacteriology</i> , 2011 , 193, 6812-4	3.5	5
19	Isolated pontine progressive multifocal leukoencephalopathy: unusual magnetic resonance imaging features. <i>Journal of Neuroimaging</i> , 2002 , 12, 63-6	2.8	5
18	Parameters Affecting Continuous Culture of Treponema pallidum Strains. MBio, 2021, 12,	7.8	5
17	Illuminating the agent of syphilis: the Treponema pallidum genome project. <i>Electrophoresis</i> , 1998 , 19, 551-3	3.6	4
16	A selective antibiotic for Lyme disease. <i>Cell</i> , 2021 , 184, 5405-5418.e16	56.2	4
15	YebC regulates variable surface antigen VlsE expression and is required for host immune evasion in Borrelia burgdorferi. <i>PLoS Pathogens</i> , 2020 , 16, e1008953	7.6	4
14	The Genus Treponema 1992 , 3537-3559		4
13	OptiSol Corneal Storage Medium and Transmission of Treponema Pallidum. <i>Cornea</i> , 1995 , 14, 595???60	03.1	3
12	Demonstration of Treponema pallidum in a cutaneous gumma by indirect immunofluorescence. <i>Archives of Dermatology</i> , 1983 , 119, 677-680		3
11	Susceptibility of Treponema pallidum subsp. to Doxycycline. <i>Antimicrobial Agents and Chemotherapy</i> , 2020 , 64,	5.9	3
10	vls Antigenic Variation Systems of Lyme Disease Borrelia: Eluding Host Immunity through both Random, Segmental Gene Conversion and Framework Heterogeneity 2015 , 471-489		2
9	Serum antibodies to whole-cell and recombinant antigens of Borrelia burgdorferi in cottontail rabbits. <i>Journal of Wildlife Diseases</i> , 2012 , 48, 12-20	1.3	2
8	Construction of small genome BAC library for functional and genomic applications. <i>Methods in Molecular Biology</i> , 2004 , 255, 47-56	1.4	1
7	Molecular Mechanism for Rotational Switching of the Bacterial Flagellar Motor		1

LIST OF PUBLICATIONS

6 Origin of modern syphilis and emergence of a contemporary pandemic cluster 1 BBB07 contributes to, but is not essential for, infection in mice. Microbiology (United Kingdom), 2.9 2020, 166, 988-994 Comparison of transcriptional profiles of Treponema pallidum during experimental infection of 7.6 О rabbits and in vitro culture: Highly similar, yet different. PLoS Pathogens, 2021, 17, e1009949 Catching up with Lyme Disease Antigenic Variation Computationally. Trends in Microbiology, 2018, 12.4 26, 644-645 Kinetics and In Vivo Induction of Genetic Variation of vlsE in Borrelia burgdorferi. Infection and 2 3.7 Immunity, 1999, 67, 468-468 Comparative Pathogenomics of Spirochetes 141-159