

# Jon T Skare

## List of Publications by Year in descending order

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Version: 2024-02-01

29  
papers

1,431  
citations

394421

19  
h-index

501196

28  
g-index

31  
all docs

31  
docs citations

31  
times ranked

740  
citing authors

#	ARTICLE	IF	CITATIONS
1	Profiling of Temperature-Induced Changes in <i>Borrelia burgdorferi</i> Gene Expression by Using Whole Genome Arrays. <i>Infection and Immunity</i> , 2003, 71, 1689-1705.	2.2	263
2	The BosR regulatory protein of <i>Borrelia burgdorferi</i> interfaces with the RpoS regulatory pathway and modulates both the oxidative stress response and pathogenic properties of the Lyme disease spirochete. <i>Molecular Microbiology</i> , 2009, 74, 1344-1355.	2.5	115
3	<i>Borrelia burgdorferi</i> BBK32 Inhibits the Classical Pathway by Blocking Activation of the C1 Complement Complex. <i>PLoS Pathogens</i> , 2016, 12, e1005404.	4.7	111
4	The Oms66 (p66) protein is a <i>Borrelia burgdorferi</i> porin. <i>Infection and Immunity</i> , 1997, 65, 3654-3661.	2.2	106
5	Bioluminescent imaging of <i>Borrelia burgdorferi</i> in vivo demonstrates that the fibronectin-binding protein BBK32 is required for optimal infectivity. <i>Molecular Microbiology</i> , 2011, 82, 99-113.	2.5	97
6	Vascular binding of a pathogen under shear force through mechanistically distinct sequential interactions with host macromolecules. <i>Molecular Microbiology</i> , 2012, 86, 1116-1131.	2.5	75
7	Invasion of Eukaryotic Cells by <i>Borrelia burgdorferi</i> Requires $\beta$ 1 Integrins and Src Kinase Activity. <i>Infection and Immunity</i> , 2011, 79, 1338-1348.	2.2	61
8	A conservative amino acid change alters the function of BosR, the redox regulator of <i>Borrelia burgdorferi</i> . <i>Molecular Microbiology</i> , 2004, 54, 1352-1363.	2.5	57
9	Characterization of a Conditional <i>bosR</i> Mutant in <i>Borrelia burgdorferi</i> . <i>Infection and Immunity</i> , 2010, 78, 265-274.	2.2	56
10	Analysis of Mechanisms Associated with Loss of Infectivity of Clonal Populations of <i>Borrelia burgdorferi</i> B31MI. <i>Infection and Immunity</i> , 2001, 69, 3670-3677.	2.2	53
11	Lyme Disease Pathogenesis. <i>Current Issues in Molecular Biology</i> , 2022, 42, 473-518.	2.4	49
12	Biomechanics of <i>Borrelia burgdorferi</i> Vascular Interactions. <i>Cell Reports</i> , 2016, 16, 2593-2604.	6.4	48
13	Complement Evasion by Lyme Disease Spirochetes. <i>Trends in Microbiology</i> , 2020, 28, 889-899.	7.7	48
14	A high-throughput genetic screen identifies previously uncharacterized <i>Borrelia burgdorferi</i> genes important for resistance against reactive oxygen and nitrogen species. <i>PLoS Pathogens</i> , 2017, 13, e1006225.	4.7	36
15	Structural determination of the complement inhibitory domain of <i>Borrelia burgdorferi</i> BBK32 provides insight into classical pathway complement evasion by Lyme disease spirochetes. <i>PLoS Pathogens</i> , 2019, 15, e1007659.	4.7	33
16	Cloning and Molecular Characterization of Plasmid-Encoded Antigens of <i>Borrelia burgdorferi</i> . <i>Infection and Immunity</i> , 1999, 67, 4407-4417.	2.2	32
17	<i>Borrelia burgdorferi</i> gene expression profiling with membrane-based arrays. <i>Methods in Enzymology</i> , 2002, 358, 165-177.	1.0	31
18	Genome-wide screen identifies novel genes required for <i>Borrelia burgdorferi</i> survival in its Ixodes tick vector. <i>PLoS Pathogens</i> , 2019, 15, e1007644.	4.7	25

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19	The Classical Complement Pathway Is Required to Control <i>Borrelia burgdorferi</i> Levels During Experimental Infection. <i>Frontiers in Immunology</i> , 2018, 9, 959.	4.8	22
20	The BBA33 lipoprotein binds collagen and impacts <i>Borrelia burgdorferi</i> pathogenesis. <i>Molecular Microbiology</i> , 2015, 96, 68-83.	2.5	21
21	Genetic Transformation of <i>Borrelia burgdorferi</i> . <i>Current Protocols in Microbiology</i> , 2011, 20, Unit 12C.4.	6.5	20
22	The BB0646 protein demonstrates lipase and haemolytic activity associated with <i>Borrelia burgdorferi</i> , the aetiological agent of Lyme disease. <i>Molecular Microbiology</i> , 2012, 83, 319-334.	2.5	16
23	BB0744 Affects Tissue Tropism and Spatial Distribution of <i>Borrelia burgdorferi</i> . <i>Infection and Immunity</i> , 2015, 83, 3693-3703.	2.2	13
24	The intergenic small non-coding RNA <i>ittA</i> is required for optimal infectivity and tissue tropism in <i>Borrelia burgdorferi</i> . <i>PLoS Pathogens</i> , 2020, 16, e1008423.	4.7	13
25	A Structural Basis for Inhibition of the Complement Initiator Protease C1r by Lyme Disease Spirochetes. <i>Journal of Immunology</i> , 2021, 207, 2856-2867.	0.8	11
26	Detection of Bioluminescent <i>Borrelia burgdorferi</i> from In Vitro Cultivation and During Murine Infection. <i>Methods in Molecular Biology</i> , 2018, 1690, 241-257.	0.9	7
27	<i>Borrelia miyamotoi</i> FbpA and FbpB Are Immunomodulatory Outer Surface Lipoproteins With Distinct Structures and Functions. <i>Frontiers in Immunology</i> , 2022, 13, .	4.8	7
28	Live Imaging. <i>Current Issues in Molecular Biology</i> , 2022, 42, 385-408.	2.4	3
29	Minimal Role for the Alternative Pathway in Complement Activation By HIT Immune Complexes. <i>Blood</i> , 2021, 138, 2076-2076.	1.4	0