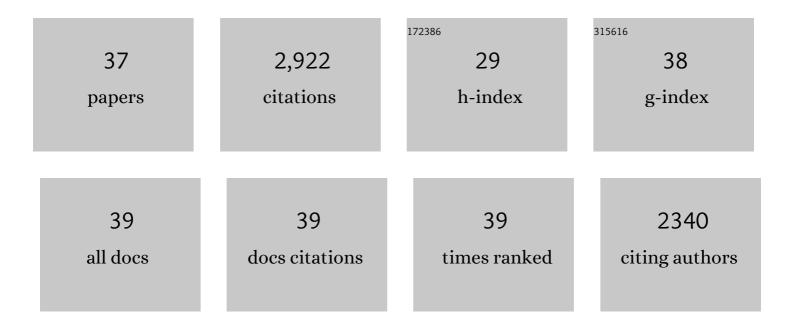


## List of Publications by Year in descending order

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YINTI

| #  | Article   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | Trace Element Analysis of Borrelia burgdorferi by Inductively Coupled Plasma-Sector Field Mass<br>Spectrometry. Methods in Molecular Biology, 2018, 1690, 83-94.  | 0.4  | 1         |
| 2  | BosR Is A Novel Fur Family Member Responsive to Copper and Regulating Copper Homeostasis in<br>Borrelia burgdorferi. Journal of Bacteriology, 2017, 199, .  | 1.0  | 12        |
| 3  | A high-throughput genetic screen identifies previously uncharacterized Borrelia burgdorferi genes<br>important for resistance against reactive oxygen and nitrogen species. PLoS Pathogens, 2017, 13,<br>e1006225.                      | 2.1  | 36        |
| 4  | Lyme borreliosis. Nature Reviews Disease Primers, 2016, 2, 16090.   | 18.1 | 530       |
| 5  | BosR Functions as a Repressor of the ospAB Operon in Borrelia burgdorferi. PLoS ONE, 2014, 9, e109307.  | 1.1  | 26        |
| 6  | Emergence of Ixodes scapularis and Borrelia burgdorferi, the Lyme disease vector and agent, in Ohio.<br>Frontiers in Cellular and Infection Microbiology, 2014, 4, 70.  | 1.8  | 23        |
| 7  | TRIF Mediates Toll-Like Receptor 2-Dependent Inflammatory Responses to Borrelia burgdorferi.<br>Infection and Immunity, 2013, 81, 402-410.  | 1.0  | 54        |
| 8  | Tick-Specific Borrelial Antigens Appear to Be Upregulated in American but Not European Patients With<br>Lyme Arthritis, a Late Manifestation of Lyme Borreliosis. Journal of Infectious Diseases, 2013, 208,<br>934-941.                | 1.9  | 16        |
| 9  | <i><scp>B</scp>orrelia burgdorferi</i> oxidative stress regulator <scp>BosR</scp> directly represses lipoproteins primarily expressed in the tick during mammalian infection. Molecular Microbiology, 2013, 89, 1140-1153.              | 1.2  | 40        |
| 10 | A novel iron―and copperâ€binding protein in the <scp>L</scp> yme disease spirochaete. Molecular<br>Microbiology, 2012, 86, 1441-1451.   | 1.2  | 50        |
| 11 | Borrelia burgdorferi RST1 (OspC Type A) Genotype Is Associated with Greater Inflammation and More<br>Severe Lyme Disease. American Journal of Pathology, 2011, 178, 2726-2739.  | 1.9  | 105       |
| 12 | Burden and viability of <i>Borrelia burgdorferi</i> in skin and joints of patients with erythema migrans or lyme arthritis. Arthritis and Rheumatism, 2011, 63, 2238-2247.  | 6.7  | 124       |
| 13 | Ehrlichia chaffeensis Induces Monocyte Inflammatory Responses through MyD88, ERK, and NF-κB but<br>Not through TRIF, Interleukin-1 Receptor 1 (IL-1R1)/IL-18R1, or Toll-Like Receptors. Infection and Immunity,<br>2011, 79, 4947-4956. | 1.0  | 32        |
| 14 | Treg cell numbers and function in patients with antibioticâ€refractory or antibioticâ€responsive lyme<br>arthritis. Arthritis and Rheumatism, 2010, 62, 2127-2137.  | 6.7  | 49        |
| 15 | Oxygen-Limiting Conditions Enrich for Fimbriate Cells of Uropathogenic <i>Proteus<br/>mirabilis</i> and <i>Escherichia coli</i> . Journal of Bacteriology, 2009, 191, 1382-1392.  | 1.0  | 44        |
| 16 | A Differential Role for BB0365 in the Persistence ofBorrelia burgdorferiin Mice and Ticks. Journal of<br>Infectious Diseases, 2008, 197, 148-155.   | 1.9  | 52        |
| 17 | Outer Surface Protein B Is Critical for Borrelia burgdorferi Adherence and Survival within Ixodes<br>Ticks. PLoS Pathogens, 2007, 3, e33.   | 2.1  | 78        |
| 18 | Role of Outer Surface Protein D in the <i>Borrelia burgdorferi</i> Life Cycle. Infection and Immunity, 2007, 75, 4237-4244.   | 1.0  | 36        |

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|----|---|------|-----------|
| 19 | The Lyme disease agent Borrelia burgdorferi requires BB0690, a Dps homologue, to persist within ticks. Molecular Microbiology, 2007, 63, 694-710.   | 1.2  | 110       |
| 20 | Coinfection withBorrelia burgdorferisensu stricto andBorrelia gariniialters the course of murine<br>Lyme borreliosis. FEMS Immunology and Medical Microbiology, 2007, 49, 224-234.  | 2.7  | 15        |
| 21 | Borrelia burgdorferi Lacking BBK32, a Fibronectin-Binding Protein, Retains Full Pathogenicity.<br>Infection and Immunity, 2006, 74, 3305-3313.  | 1.0  | 87        |
| 22 | Association of Linear Plasmid 28-1 with an Arthritic Phenotype of Borrelia burgdorferi. Infection and<br>Immunity, 2005, 73, 7208-7215.   | 1.0  | 33        |
| 23 | Development of an Intranasal Vaccine To Prevent Urinary Tract Infection by Proteus mirabilis.<br>Infection and Immunity, 2004, 72, 66-75.   | 1.0  | 67        |
| 24 | Use of Translational Fusion of the MrpH Fimbrial Adhesin-Binding Domain with the Cholera Toxin A2<br>Domain, Coexpressed with the Cholera Toxin B Subunit, as an Intranasal Vaccine To Prevent<br>Experimental Urinary Tract Infection by Proteus mirabilis. Infection and Immunity, 2004, 72, 7306-7310. | 1.0  | 37        |
| 25 | Proteus mirabilis Genes That Contribute to Pathogenesis of Urinary Tract Infection: Identification of 25 Signature-Tagged Mutants Attenuated at Least 100-Fold. Infection and Immunity, 2004, 72, 2922-2938.  | 1.0  | 172       |
| 26 | TROSPA, an Ixodes scapularis Receptor for Borrelia burgdorferi. Cell, 2004, 119, 457-468.   | 13.5 | 348       |
| 27 | Visualization of Proteus mirabilis within the Matrix of Urease-Induced Bladder Stones during Experimental Urinary Tract Infection. Infection and Immunity, 2002, 70, 389-394.   | 1.0  | 88        |
| 28 | Vaccines for Proteus mirabilis in urinary tract infection. International Journal of Antimicrobial<br>Agents, 2002, 19, 461-465.   | 1.1  | 44        |
| 29 | Identification of MrpI as the sole recombinase that regulates the phase variation of MR/P fimbria, a<br>bladder colonization factor of uropathogenic Proteus mirabilis. Molecular Microbiology, 2002, 45,<br>865-874.   | 1.2  | 66        |
| 30 | Repression of bacterial motility by a novel fimbrial gene product. EMBO Journal, 2001, 20, 4854-4862.   | 3.5  | 81        |
| 31 | Identification of DNA Sequences from a Second Pathogenicity Island of UropathogenicEscherichia<br>coliCFT073: Probes Specific for Uropathogenic Populations. Journal of Infectious Diseases, 2001, 184,<br>1041-1049.   | 1.9  | 49        |
| 32 | Pathogenesis of Proteus mirabilisurinary tract infection. Microbes and Infection, 2000, 2, 1497-1505.   | 1.0  | 149       |
| 33 | Identification of protease and rpoN-associated genes of uropathogenic Proteus mirabilis by negative selection in a mouse model of ascending urinary tract infection. Microbiology (United Kingdom), 1999, 145, 185-195.   | 0.7  | 68        |
| 34 | Requirement of MrpH for Mannose-Resistant <i>Proteus</i> -Like Fimbria-Mediated Hemagglutination<br>by <i>Proteus mirabilis</i> . Infection and Immunity, 1999, 67, 2822-2833.  | 1.0  | 55        |
| 35 | MrpB Functions as the Terminator for Assembly of <i>Proteus mirabilis</i> Mannose-Resistant<br><i>Proteus</i> -Like Fimbriae. Infection and Immunity, 1998, 66, 1759-1763.  | 1.0  | 16        |
| 36 | In vivo phase variation of MR/P fimbrial gene expression in Proteus mirabilis infecting the urinary tract. Molecular Microbiology, 1997, 23, 1009-1019.   | 1.2  | 91        |

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| 37 | Proteus mirabilis mannose-resistant, Proteus-like fimbriae: MrpG is located at the fimbrial tip and is required for fimbrial assembly. Infection and Immunity, 1997, 65, 1327-1334. | 1.0 | 31        |