## **Robert Boissy**

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Comparative Genomic Analyses of 17 Clinical Isolates of Gardnerella vaginalis Provide Evidence of Multiple Genetically Isolated Clades Consistent with Subspeciation into Genovars. Journal of Bacteriology, 2012, 194, 3922-3937.	2.2	147
2	Comparative supragenomic analyses among the pathogens Staphylococcus aureus, Streptococcus pneumoniae, and Haemophilus influenzae Using a modification of the finite supragenome model. BMC Genomics, 2011, 12, 187.	2.8	50
3	<i>Cellulosilyticum ruminicola</i> , a Newly Described Rumen Bacterium That Possesses Redundant Fibrolytic-Protein-Encoding Genes and Degrades Lignocellulose with Multiple Carbohydrate- Borne Fibrolytic Enzymes. Applied and Environmental Microbiology, 2010, 76, 3818-3824.	3.1	71
4	Insights into the Genome of Large Sulfur Bacteria Revealed by Analysis of Single Filaments. PLoS Biology, 2007, 5, e230.	5.6	151
5	Comparative Genomic Analyses of Seventeen <i>Streptococcus pneumoniae</i> Strains: Insights into the Pneumococcal Supragenome. Journal of Bacteriology, 2007, 189, 8186-8195.	2.2	249
6	Characterization and modeling of the Haemophilus influenzae core and supragenomes based on the complete genomic sequences of Rd and 12 clinical nontypeable strains. Genome Biology, 2007, 8, R103.	9.6	228
7	Reply to the letter to the Editor: "N-Acetyltransferases and the susceptibility to benzidine-induced bladder carcinogenesis― International Journal of Cancer, 2007, 121, 1637-1639.	5.1	3
8	NAT2 slow acetylation and bladder cancer in workers exposed to benzidine. International Journal of Cancer, 2006, 118, 161-168.	5.1	62
9	Inherited polymorphism in the N-acetyltransferase 1 (NAT1) and 2 (NAT2) genes and susceptibility to	5.1	59