

# Robert G Quivey Jr

## List of Publications by Year in descending order

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45  
papers

2,325  
citations

236925

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docs citations

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times ranked

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citing authors

#	ARTICLE	IF	CITATIONS
1	Analysis of the <i>Streptococcus mutans</i> Proteome during Acid and Oxidative Stress Reveals Modules of Protein Coexpression and an Expanded Role for the TreR Transcriptional Regulator. <i>MSystems</i> , 2022, 7, e0127221.	3.8	8
2	Prediction of early childhood caries onset and oral microbiota. <i>Molecular Oral Microbiology</i> , 2021, 36, 255-257.	2.7	3
3	<i>Streptococcus mutans</i> SpxA2 relays the signal of cell envelope stress from LiaR to effectors that maintain cell wall and membrane homeostasis. <i>Molecular Oral Microbiology</i> , 2020, 35, 118-128.	2.7	10
4	Disruption of <i>scp</i> -Rhamnose Biosynthesis Results in Severe Growth Defects in <i>Streptococcus mutans</i> . <i>Journal of Bacteriology</i> , 2020, 202, .	2.2	14
5	<i>Streptococcus mutans</i> requires mature rhamnose-glucose polysaccharides for proper pathophysiology, morphogenesis and cellular division. <i>Molecular Microbiology</i> , 2019, 112, 944-959.	2.5	27
6	Inactivation of <i>Streptococcus mutans</i> genes <i>lytST</i> and <i>dltAD</i> impairs its pathogenicity in vivo. <i>Journal of Oral Microbiology</i> , 2019, 11, 1607505.	2.7	18
7	Characterization of the Trehalose Utilization Operon in <i>Streptococcus mutans</i> Reveals that the TreR Transcriptional Regulator Is Involved in Stress Response Pathways and Toxin Production. <i>Journal of Bacteriology</i> , 2018, 200, .	2.2	24
8	A Drug Repositioning Approach Reveals that <i>Streptococcus mutans</i> Is Susceptible to a Diverse Range of Established Antimicrobials and Nonantibiotics. <i>Antimicrobial Agents and Chemotherapy</i> , 2018, 62, .	3.2	23
9	Vitamin D Compounds Are Bactericidal against <i>Streptococcus mutans</i> and Target the Bacitracin-Associated Efflux System. <i>Antimicrobial Agents and Chemotherapy</i> , 2018, 62, .	3.2	31
10	Deficiency of BrpA in <i>Streptococcus mutans</i> reduces virulence in rat caries model. <i>Molecular Oral Microbiology</i> , 2018, 33, 353-363.	2.7	17
11	Diverted Total Synthesis of Carolacton-Inspired Analogs Yields Three Distinct Phenotypes in <i>Streptococcus mutans</i> Biofilms. <i>Journal of the American Chemical Society</i> , 2017, 139, 7188-7191.	13.7	27
12	Extracellular DNA and lipoteichoic acids interact with exopolysaccharides in the extracellular matrix of <i>Streptococcus mutans</i> biofilms. <i>Biofouling</i> , 2017, 33, 722-740.	2.2	63
13	<i>Candida albicans</i> Carriage in Children with Severe Early Childhood Caries (S-ECC) and Maternal Relatedness. <i>PLoS ONE</i> , 2016, 11, e0164242.	2.5	84
14	What Are We Learning and What Can We Learn from the Human Oral Microbiome Project?. <i>Current Oral Health Reports</i> , 2016, 3, 56-63.	1.6	12
15	PlsX deletion impacts fatty acid synthesis and acid adaptation in <i>Streptococcus mutans</i> . <i>Microbiology (United Kingdom)</i> , 2016, 162, 662-671.	1.8	5
16	A Modified Chromogenic Assay for Determination of the Ratio of Free Intracellular NAD <sup>+</sup> /NADH in <i>Streptococcus mutans</i> . <i>Bio-protocol</i> , 2016, 6, .	0.4	9
17	Î <sup>2</sup> -Phosphoglucomutase contributes to aciduricity in <i>Streptococcus mutans</i> . <i>Microbiology (United Kingdom)</i> 10.1093/mic/kgy118	1.8	18
18	<i>Streptococcus mutans</i> : a new Gram-positive paradigm?. <i>Microbiology (United Kingdom)</i> , 2013, 159, 436-445.	1.8	174

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19	The Streptococcus mutans Aminotransferase Encoded by <i>ilvE</i> Is Regulated by CodY and CcpA. Journal of Bacteriology, 2013, 195, 3552-3562.	2.2	24
20	The Branched-Chain Amino Acid Aminotransferase Encoded by <i>ilvE</i> Is Involved in Acid Tolerance in Streptococcus mutans. Journal of Bacteriology, 2012, 194, 2010-2019.	2.2	78
21	Mutation of the NADH Oxidase Gene ( <i>nox</i> ) Reveals an Overlap of the Oxygen- and Acid-Mediated Stress Responses in Streptococcus mutans. Applied and Environmental Microbiology, 2012, 78, 1215-1227.	3.1	46
22	Cardiolipin biosynthesis in Streptococcus mutans is regulated in response to external pH. Microbiology (United Kingdom), 2012, 158, 2133-2143.	1.8	30
23	Role of DNA base excision repair in the mutability and virulence of <i>Streptococcus mutans</i> . Molecular Microbiology, 2012, 85, 361-377.	2.5	17
24	Responses of Lactic Acid Bacteria to Acid Stress. , 2011, , 23-53.		7
25	Two Spx Proteins Modulate Stress Tolerance, Survival, and Virulence in <i>Streptococcus mutans</i> . Journal of Bacteriology, 2010, 192, 2546-2556.	2.2	109
26	Alkali production associated with malolactic fermentation by oral streptococci and protection against acid, oxidative, or starvation damage. Canadian Journal of Microbiology, 2010, 56, 539-547.	1.7	45
27	Role of Clp Proteins in Expression of Virulence Properties of <i>Streptococcus mutans</i> . Journal of Bacteriology, 2009, 191, 2060-2068.	2.2	84
28	Role of Unsaturated Fatty Acid Biosynthesis in Virulence of Streptococcus mutans. Infection and Immunity, 2007, 75, 1537-1539.	2.2	58
29	Smx Nuclease Is the Major, Low-pH-Inducible Apurinic/Apyrimidinic Endonuclease in Streptococcus mutans. Journal of Bacteriology, 2005, 187, 2705-2714.	2.2	15
30	The putative autolysin regulator LytR in Streptococcus mutans plays a role in cell division and is growth-phase regulated. Microbiology (United Kingdom), 2005, 151, 625-631.	1.8	91
31	The F-ATPase Operon Promoter of Streptococcus mutans Is Transcriptionally Regulated in Response to External pH. Journal of Bacteriology, 2004, 186, 8524-8528.	2.2	82
32	The <i>fabM</i> Gene Product of Streptococcus mutans Is Responsible for the Synthesis of Monounsaturated Fatty Acids and Is Necessary for Survival at Low pH. Journal of Bacteriology, 2004, 186, 4152-4158.	2.2	111
33	Shifts in the Membrane Fatty Acid Profile of <i>Streptococcus mutans</i> Enhance Survival in Acidic Environments. Applied and Environmental Microbiology, 2004, 70, 929-936.	3.1	189
34	Low pH-induced membrane fatty acid alterations in oral bacteria. FEMS Microbiology Letters, 2004, 238, 291-295.	1.8	107
35	Low pH-induced membrane fatty acid alterations in oral bacteria. FEMS Microbiology Letters, 2004, 238, 291-295.	1.8	60
36	Genetic and Biochemical Characterization of the F-ATPase Operon from <i>S. streptococcus sanguis</i> 10904. Journal of Bacteriology, 2003, 185, 1525-1533.	2.2	45

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37	Genetics of Acid Adaptation in Oral Streptococci. Critical Reviews in Oral Biology and Medicine, 2001, 12, 301-314.	4.4	109
38	Shifts in membrane fatty acid profiles associated with acid adaptation of <i>Streptococcus mutans</i> . FEMS Microbiology Letters, 2000, 189, 89-92.	1.8	105
39	Adaptation of oral streptococci to low pH. Advances in Microbial Physiology, 2000, 42, 239-274.	2.4	124
40	[33] Physiologic homeostasis and stress responses in oral biofilms. Methods in Enzymology, 1999, 310, 441-460.	1.0	38
41	Use of proteomics and PCR to elucidate changes in protein expression in oral streptococci. Cytotechnology, 1998, 20, 165-179.	0.7	2
42	Cloning and nucleotide sequence analysis of the <i>Streptococcus mutans</i> membrane-bound, proton-translocating ATPase operon. Gene, 1996, 183, 87-96.	2.2	51
43	Acid adaptation in <i>Streptococcus mutans</i> UA159 alleviates sensitization to environmental stress due to RecA deficiency. FEMS Microbiology Letters, 1995, 126, 257-262.	1.8	77
44	In vivo inactivation of the <i>Streptococcus mutans</i> recA gene mediated by PCR amplification and cloning of a recA DNA fragment. Gene, 1992, 116, 35-42.	2.2	31
45	Polymerase chain reaction amplification, cloning, sequence determination and homologies of streptococcal ATPase-encoding DNAs. Gene, 1991, 97, 63-68.	2.2	22