## Jonathon Baker

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/180852/publications.pdf Version: 2024-02-01



IONATHON RAKER

| #  | Article  | IF  | CITATIONS |
|----|--|-----|-----------|
| 1  | Ecology of the Oral Microbiome: Beyond Bacteria. Trends in Microbiology, 2017, 25, 362-374.  | 7.7 | 222       |
| 2  | Host Factor SAMHD1 Restricts DNA Viruses in Non-Dividing Myeloid Cells. PLoS Pathogens, 2013, 9, e1003481.   | 4.7 | 151       |
| 3  | Development of a Bacteriophage Cocktail to Constrain the Emergence of Phage-Resistant Pseudomonas aeruginosa. Frontiers in Microbiology, 2020, 11, 327.  | 3.5 | 92        |
| 4  | Acidâ€adaptive mechanisms of <i>Streptococcus mutans</i> –the more we know, the more we don't.<br>Molecular Oral Microbiology, 2017, 32, 107-117.  | 2.7 | 75        |
| 5  | Identification of the Bacterial Biosynthetic Gene Clusters of the Oral Microbiome Illuminates the<br>Unexplored Social Language of Bacteria during Health and Disease. MBio, 2019, 10, .   | 4.1 | 73        |
| 6  | Exploiting the Oral Microbiome to Prevent Tooth Decay: Has Evolution Already Provided the Best<br>Tools?. Frontiers in Microbiology, 2018, 9, 3323.  | 3.5 | 70        |
| 7  | Deep metagenomics examines the oral microbiome during dental caries, revealing novel taxa and co-occurrences with host molecules. Genome Research, 2021, 31, 64-74.  | 5.5 | 59        |
| 8  | Streptococcus mutans NADH Oxidase Lies at the Intersection of Overlapping Regulons Controlled by<br>Oxygen and NAD <sup>+</sup> Levels. Journal of Bacteriology, 2014, 196, 2166-2177.   | 2.2 | 54        |
| 9  | Cariogenic <i>Streptococcus mutans</i> Produces Tetramic Acid Strain-Specific Antibiotics That<br>Impair Commensal Colonization. ACS Infectious Diseases, 2020, 6, 563-571.  | 3.8 | 40        |
| 10 | <i>Klebsiella</i> and <i>Providencia</i> emerge as lone survivors following long-term starvation of<br>oral microbiota. Proceedings of the National Academy of Sciences of the United States of America,<br>2019, 116, 8499-8504.      | 7.1 | 30        |
| 11 | Transcriptional profile of glucoseâ€shocked and acidâ€adapted strains of <i>Streptococcus mutans</i> .<br>Molecular Oral Microbiology, 2015, 30, 496-517.  | 2.7 | 27        |
| 12 | Characterization of the Trehalose Utilization Operon in Streptococcus mutans Reveals that the TreR<br>Transcriptional Regulator Is Involved in Stress Response Pathways and Toxin Production. Journal of<br>Bacteriology, 2018, 200, . | 2.2 | 24        |
| 13 | Development and comparison of a quantitative TaqMan-MGB real-time PCR assay to three other methods of quantifying vaccinia virions. Journal of Virological Methods, 2014, 196, 126-132.  | 2.1 | 23        |
| 14 | Loss of NADH Oxidase Activity in Streptococcus mutans Leads to Rex-Mediated Overcompensation in<br>NAD <sup>+</sup> Regeneration by Lactate Dehydrogenase. Journal of Bacteriology, 2015, 197, 3645-3657.                              | 2.2 | 23        |
| 15 | Precision Reengineering of the Oral Microbiome for Caries Management. Advances in Dental Research, 2019, 30, 34-39.  | 3.6 | 20        |
| 16 | Multi-Omics Study of Keystone Species in a Cystic Fibrosis Microbiome. International Journal of<br>Molecular Sciences, 2021, 22, 12050.  | 4.1 | 14        |
| 17 | Caries-Associated Biosynthetic Gene Clusters in <i>Streptococcus mutans</i> . Journal of Dental Research, 2020, 99, 969-976.   | 5.2 | 13        |
| 18 | <i>Streptococcus mutans</i> SpxA2 relays the signal of cell envelope stress from LiaR to effectors that maintain cell wall and membrane homeostasis. Molecular Oral Microbiology, 2020, 35, 118-128.                                   | 2.7 | 10        |

JONATHON BAKER

| #  | Article  | IF             | CITATIONS  |
|----|--|----------------|------------|
| 19 | Composite Long- and Short-Read Sequencing Delivers a Complete Genome Sequence of B04Sm5, a<br>Reutericyclin- and Mutanocyclin-Producing Strain of Streptococcus mutans. Microbiology Resource<br>Announcements, 2020, 9, . | 0.6            | 9          |
| 20 | Complete Genomes of Clade G6 <i>Saccharibacteria</i> Suggest a Divergent Ecological Niche and Lifestyle. MSphere, 2021, 6, e0053021.   | 2.9            | 9          |
| 21 | A Modified Chromogenic Assay for Determination of the Ratio of Free Intracellular NAD+/NADH in Streptococcus mutans. Bio-protocol, 2016, 6, .  | 0.4            | 9          |
| 22 | Analysis of the Streptococcus mutans Proteome during Acid and Oxidative Stress Reveals Modules of<br>Protein Coexpression and an Expanded Role for the TreR Transcriptional Regulator. MSystems, 2022, 7,<br>e0127221.     | 3.8            | 8          |
| 23 | Tetramic Acids Mutanocyclin and Reutericyclin A, Produced by Streptococcus mutans Strain B04Sm5<br>Modulate the Ecology of an in vitro Oral Biofilm. Frontiers in Oral Health, 2021, 2, 796140.                            | 3.0            | 5          |
| 24 | <i>mucG, mucH,</i> and <i>mucl</i> Modulate Production of Mutanocyclin and Reutericyclins in<br>Streptococcus mutans B04Sm5. Journal of Bacteriology, 2022, 204, e0004222.   | 2.2            | 4          |
| 25 | Complete Genome Sequence of Strain JB001, a Member of Saccharibacteria Clade G6 (" Candidatus) Tj ETQq1  | 10.7843<br>0.6 | 14 rgBT /O |
| 26 | Identification of Bacterial Biosynthetic Gene Associated with Caries. Methods in Molecular Biology, 2021, 2327, 161-189.   | 0.9            | 2          |
| 27 | Complete Genome Sequence of " <i>Candidatus</i> Nanosynbacter―Strain HMT-348_TM7c-JB, a Member<br>of <i>Saccharibacteria</i> Clade G1. Microbiology Resource Announcements, 2022, , e0002322.                              | 0.6            | 2          |