## Robert Schmieder

## List of Publications by Year

 in descending orderSource: https:|/exaly.com/author-pdf/12108646/publications.pdf
Version: 2024-02-01

$1 \quad$ Allelic variation contributes to bacterial host specificity. Nature Communications, 2015, 6, 8754.
5.8

100

2 Genomic Comparison of the Closely-Related Salmonella enterica Serovars Enteritidis, Dublin and Gallinarum. PLoS ONE, 2015, 10, e0126883.

Species-Specific Viromes in the Ancestral Holobiont Hydra. PLoS ONE, 2014, 9, e109952.
1.1

Breath gas metabolites and bacterial metagenomes from cystic fibrosis airways indicate active pH
$4 \begin{aligned} & \text { Breath gas metabolites and bacterial metagenomes from cystic fibrosis air } \\ & \text { neutral 2,3-butanedione fermentation. ISME Journal, 2014, 8, 1247-1258. }\end{aligned}$
4.4

114

Clinical Insights from Metagenomic Analysis of Sputum Samples from Patients with Cystic Fibrosis.
Journal of Clinical Microbiology, 2014, 52, 425-437.
1.8

Comparative genomics of 274 Vibrio cholerae genomes reveals mobile functions structuring three
$6 \quad \begin{aligned} & \text { Comparative genomics of } 274 \text { Vibrio cholerae gen } \\ & \text { niche dimensions. BMC Genomics, 2014, 15, } 654 .\end{aligned}$
1.2

24

Metagenomics and metatranscriptomics: Windows on CF-associated viral and microbial communities.
Metagenomics and metatranscriptomics: Win
Journal of Cystic Fibrosis, 2013, 12, 154-164.
0.3

142

Combining de novo and reference-guided assembly with scaffold_builder. Source Code for Biology
$8 \quad \begin{aligned} & \text { Combining de novo and ref } \\ & \text { and Medicine, 2013, 8, } 23 .\end{aligned}$
1.7

59
$9 \quad$ Big data challenges and opportunities in high-throughput sequencing. Systems Biomedicine (Austin,) Tj ETQq1 10.684314 rgBTh $/ \mathrm{Ove}$

10 Mechanistic Model of Rothia mucilaginosa Adaptation toward Persistence in the CF Lung, Based on a Genome Reconstructed from Metagenomic Data. PLoS ONE, 2013, 8, e64285.

## 11 Spatial distribution of microbial communities in the cystic fibrosis lung. ISME Journal, 2012, 6, 471-474.

$4.4 \quad 156$

12 Identification and removal of ribosomal RNA sequences from metatranscriptomes. Bioinformatics, 2012, 28, 433-435.
1.8

212
SEQanswers: an open access community for collaboratively decoding genomes. Bioinform
28, 1272-1273. Microfluidic PCR Combined with Pyrosequencing for Identification of Allelic Variants with
Phenotypic Associations among Targeted Salmonella Genes. Applied and Environmental Microbiology,
1.4

14
2012, 78, 7480-7482.
Case Studies of the Spatial Heterogeneity of DNA Viruses in the Cystic Fibrosis Lung. American Journal
of Respiratory Cell and Molecular Biology, 2012, 46, 127-131.
1.4

102

Reference-independent comparative metagenomics using cross-assembly: crAss. Bioinformatics, 2012,
28, 3225-3231.

Insights into antibiotic resistance through metagenomic approaches. Future Microbiology, 2012, 7,
73-89.
1.0

251

21 Quality control and preprocessing of metagenomic datasets. Bioinformatics, 2011, 27, 863-864.

22 Metagenomic detection of phage-encoded platelet-binding factors in the human oral cavity.
Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 4547-4553.
3.3

123

TagCleaner: Identification and removal of tag sequences from genomic and metagenomic datasets.

