

# Rolf Hilker

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6499807/publications.pdf>

Version: 2024-02-01

16  
papers

3,268  
citations

516710

16  
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888059

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18  
docs citations

18  
times ranked

6834  
citing authors

#	ARTICLE	IF	CITATIONS
1	BNT162b2 vaccine induces neutralizing antibodies and poly-specific T cells in humans. <i>Nature</i> , 2021, 595, 572-577.	27.8	583
2	COVID-19 vaccine BNT162b1 elicits human antibody and TH1 T cell responses. <i>Nature</i> , 2020, 586, 594-599.	27.8	1,520
3	Key Parameters of Tumor Epitope Immunogenicity Revealed Through a Consortium Approach Improve Neoantigen Prediction. <i>Cell</i> , 2020, 183, 818-834.e13.	28.9	287
4	Genome Analysis of the Carbapenem- and Colistin-Resistant <i>Escherichia coli</i> Isolate NRZ14408 Reveals Horizontal Gene Transfer Pathways towards Panresistance and Enhanced Virulence. <i>Antimicrobial Agents and Chemotherapy</i> , 2017, 61, .	3.2	9
5	<i>Sinorhizobium meliloti</i> YbeY is an endoribonuclease with unprecedented catalytic features, acting as silencing enzyme in riboregulation. <i>Nucleic Acids Research</i> , 2017, 45, 1371-1391.	14.5	29
6	Genome-wide identification of transcriptional start sites in the haloarchaeon <i>Haloferax volcanii</i> based on differential RNA-Seq (dRNA-Seq). <i>BMC Genomics</i> , 2016, 17, 629.	2.8	125
7	Intraclonal genome diversity of the major <i>Pseudomonas aeruginosa</i> clones <i>C</i> and <i>PA</i> 14. <i>Environmental Microbiology Reports</i> , 2016, 8, 227-234.	2.4	41
8	ReadXplorer 2 detailed read mapping analysis and visualization from one single source. <i>Bioinformatics</i> , 2016, 32, 3702-3708.	4.1	96
9	Presence of a widely disseminated <i>Listeria monocytogenes</i> serotype 4b clone in India. <i>Emerging Microbes and Infections</i> , 2016, 5, 1-4.	6.5	17
10	Cyclic Di-GMP Regulates Multiple Cellular Functions in the Symbiotic Alphaproteobacterium <i>Sinorhizobium meliloti</i> . <i>Journal of Bacteriology</i> , 2016, 198, 521-535.	2.2	56
11	A detailed view of the intracellular transcriptome of <i>Listeria monocytogenes</i> in murine macrophages using RNA-seq. <i>Frontiers in Microbiology</i> , 2015, 6, 1199.	3.5	36
12	Interclonal gradient of virulence in the <i>Pseudomonas aeruginosa</i> pangenome from disease and environment. <i>Environmental Microbiology</i> , 2015, 17, 29-46.	3.8	113
13	Deletion of the Sm1 encoding motif in the lsm gene results in distinct changes in the transcriptome and enhanced swarming activity of <i>Haloferax</i> cells. <i>Biochimie</i> , 2015, 117, 129-137.	2.6	27
14	Serotype 1 and 8 Pneumococci Evade Sensing by Inflammasomes in Human Lung Tissue. <i>PLoS ONE</i> , 2015, 10, e0137108.	2.5	31
15	ReadXplorer visualization and analysis of mapped sequences. <i>Bioinformatics</i> , 2014, 30, 2247-2254.	4.1	127
16	UniMoG a unifying framework for genomic distance calculation and sorting based on DCJ. <i>Bioinformatics</i> , 2012, 28, 2509-2511.	4.1	38