

Winfried Hausner

List of Publications by Year in descending order

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Version: 2024-02-01

13
papers

458
citations

1040056

9
h-index

1125743

13
g-index

14
all docs

14
docs citations

14
times ranked

388
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Insights into synthesis and function of KsgA/Dim1-dependent rRNA modifications in archaea. <i>Nucleic Acids Research</i> , 2021, 49, 1662-1687. | 14.5 | 20 |
| 2 | CopR, a Global Regulator of Transcription to Maintain Copper Homeostasis in <i>Pyrococcus furiosus</i> . <i>Frontiers in Microbiology</i> , 2020, 11, 613532. | 3.5 | 10 |
| 3 | Next Generation DNA-Seq and Differential RNA-Seq Allow Re-annotation of the <i>Pyrococcus furiosus</i> DSM 3638 Genome and Provide Insights Into Archaeal Antisense Transcription. <i>Frontiers in Microbiology</i> , 2019, 10, 1603. | 3.5 | 15 |
| 4 | Inward-facing conformation of a multidrug resistance MATE family transporter. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 12275-12284. | 7.1 | 36 |
| 5 | The Transcriptional Regulator TFB-RF1 Activates Transcription of a Putative ABC Transporter in <i>Pyrococcus furiosus</i> . <i>Frontiers in Microbiology</i> , 2018, 9, 838. | 3.5 | 7 |
| 6 | Genome-wide binding analysis of the transcriptional regulator TrmBL1 in <i>Pyrococcus furiosus</i> . <i>BMC Genomics</i> , 2016, 17, 40. | 2.8 | 37 |
| 7 | The TrmB family: a versatile group of transcriptional regulators in Archaea. <i>Extremophiles</i> , 2014, 18, 925-936. | 2.3 | 27 |
| 8 | Activation of a Chimeric Rpb5/RpoH Subunit Using Library Selection. <i>PLoS ONE</i> , 2014, 9, e87485. | 2.5 | 6 |
| 9 | Genetic engineering of <i>Pyrococcus furiosus</i> to use chitin as a carbon source. <i>BMC Biotechnology</i> , 2013, 13, 9. | 3.3 | 23 |
| 10 | Activation of Archaeal Transcription Mediated by Recruitment of Transcription Factor B. <i>Journal of Biological Chemistry</i> , 2012, 287, 18863-18871. | 3.4 | 38 |
| 11 | Shuttle Vector-Based Transformation System for <i>Pyrococcus furiosus</i> . <i>Applied and Environmental Microbiology</i> , 2010, 76, 3308-3313. | 3.1 | 80 |
| 12 | Promoter architecture and response to a positive regulator of archaeal transcription. <i>Molecular Microbiology</i> , 2005, 56, 625-637. | 2.5 | 49 |
| 13 | Activation of archaeal transcription by recruitment of the TATA-binding protein. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 5097-5102. | 7.1 | 108 |