

Irene Silveira Schrank

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5835263/publications.pdf>

Version: 2024-02-01

13

papers

263

citations

1163117

8

h-index

1199594

12

g-index

13

all docs

13

docs citations

13

times ranked

307

citing authors

#	ARTICLE	IF	CITATIONS
1	The PII Superfamily Revised: A Novel Group and Evolutionary Insights. <i>Journal of Molecular Evolution</i> , 2009, 68, 322-336.	1.8	68
2	New insights on the biology of swine respiratory tract mycoplasmas from a comparative genome analysis. <i>BMC Genomics</i> , 2013, 14, 175.	2.8	63
3	Unravelling the Transcriptome Profile of the Swine Respiratory Tract Mycoplasmas. <i>PLoS ONE</i> , 2014, 9, e110327.	2.5	34
4	Mycoplasma non-coding RNA: identification of small RNAs and targets. <i>BMC Genomics</i> , 2016, 17, 743.	2.8	27
5	Unveiling <i>Mycoplasma hyopneumoniae</i> Promoters: Sequence Definition and Genomic Distribution. <i>DNA Research</i> , 2012, 19, 103-115.	3.4	15
6	<i>Mycoplasma hyopneumoniae</i> Transcription Unit Organization: Genome Survey and Prediction. <i>DNA Research</i> , 2011, 18, 413-422.	3.4	12
7	Intrinsic terminators in <i>Mycoplasma hyopneumoniae</i> transcription. <i>BMC Genomics</i> , 2015, 16, 273.	2.8	12
8	Evaluation of growth and gene expression of <i>Mycoplasma hyopneumoniae</i> and <i>Mycoplasma hyorhinis</i> in defined medium. <i>Molecular Biology Reports</i> , 2018, 45, 2469-2479.	2.3	10
9	Genome organization in <i>Mycoplasma hyopneumoniae</i> : identification of promoter-like sequences. <i>Molecular Biology Reports</i> , 2014, 41, 5395-5402.	2.3	8
10	Repetitive Elements in <i>Mycoplasma hyopneumoniae</i> Transcriptional Regulation. <i>PLoS ONE</i> , 2016, 11, e0168626.	2.5	7
11	Global analysis of sRNA target genes in <i>Mycoplasma hyopneumoniae</i> . <i>BMC Genomics</i> , 2018, 19, 767.	2.8	5
12	The influence of regulatory elements on <i>Mycoplasma hyopneumoniae</i> 7448 transcriptional response during oxidative stress and heat shock. <i>Molecular Biology Reports</i> , 2022, 49, 139-147.	2.3	2
13	Functional characterization of the putative FAD synthase from <i>Mycoplasma hyopneumoniae</i> . <i>FEMS Microbiology Letters</i> , 2021, 368, .	1.8	0