

Claudia Sala

List of Publications by Year in descending order

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Version: 2024-02-01

48
papers

3,198
citations

201674

27
h-index

223800

46
g-index

53
all docs

53
docs citations

53
times ranked

4652
citing authors

#	ARTICLE	IF	CITATIONS
1	Benzothiazinones Kill <i>Mycobacterium tuberculosis</i> by Blocking Arabinan Synthesis. <i>Science</i> , 2009, 324, 801-804.	12.6	660
2	<i>Mycobacterium tuberculosis</i> Differentially Activates cGAS- and Inflammasome-Dependent Intracellular Immune Responses through ESX-1. <i>Cell Host and Microbe</i> , 2015, 17, 799-810.	11.0	341
3	Extremely potent human monoclonal antibodies from COVID-19 convalescent patients. <i>Cell</i> , 2021, 184, 1821-1835.e16.	28.9	180
4	Towards a new tuberculosis drug: pyridomycin – nature's isoniazid. <i>EMBO Molecular Medicine</i> , 2012, 4, 1032-1042.	6.9	175
5	Lansoprazole is an antituberculous prodrug targeting cytochrome bc1. <i>Nature Communications</i> , 2015, 6, 7659.	12.8	141
6	The PhoP-Dependent ncRNA Mcr7 Modulates the TAT Secretion System in <i>Mycobacterium tuberculosis</i> . <i>PLoS Pathogens</i> , 2014, 10, e1004183.	4.7	127
7	Simple Model for Testing Drugs against Nonreplicating <i>Mycobacterium tuberculosis</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2010, 54, 4150-4158.	3.2	117
8	Virulence Regulator EspR of <i>Mycobacterium tuberculosis</i> Is a Nucleoid-Associated Protein. <i>PLoS Pathogens</i> , 2012, 8, e1002621.	4.7	115
9	Streptomycin-Starved <i>Mycobacterium tuberculosis</i> 18b, a Drug Discovery Tool for Latent Tuberculosis. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 5782-5789.	3.2	88
10	Anticytolytic Screen Identifies Inhibitors of Mycobacterial Virulence Protein Secretion. <i>Cell Host and Microbe</i> , 2014, 16, 538-548.	11.0	83
11	The Inosine Monophosphate Dehydrogenase, GuaB2, Is a Vulnerable New Bactericidal Drug Target for Tuberculosis. <i>ACS Infectious Diseases</i> , 2017, 3, 5-17.	3.8	83
12	<i>E</i> C forms a filamentous structure in the cell envelope of <i>Mycobacterium tuberculosis</i> and impacts ESX secretion. <i>Molecular Microbiology</i> , 2017, 103, 26-38.	2.5	77
13	Assessing the essentiality of the decaprenylphospho-d-arabinofuranose pathway in <i>Mycobacterium tuberculosis</i> using conditional mutants. <i>Molecular Microbiology</i> , 2014, 92, 194-211.	2.5	76
14	Development of a repressible mycobacterial promoter system based on two transcriptional repressors. <i>Nucleic Acids Research</i> , 2010, 38, e134-e134.	14.5	74
15	Transcriptional Regulation of furA and katG upon Oxidative Stress in <i>Mycobacterium smegmatis</i> . <i>Journal of Bacteriology</i> , 2001, 183, 6801-6806.	2.2	67
16	<i>Mycobacterium tuberculosis</i> FurA Autoregulates Its Own Expression. <i>Journal of Bacteriology</i> , 2003, 185, 5357-5362.	2.2	61
17	Genome-wide regulon and crystal structure of Blal (Rv1846c) from <i>Mycobacterium tuberculosis</i> . <i>Molecular Microbiology</i> , 2009, 71, 1102-1116.	2.5	61
18	Transcription facilitated genome-wide recruitment of topoisomerase I and DNA gyrase. <i>PLoS Genetics</i> , 2017, 13, e1006754.	3.5	56

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19	In Vitro and In Vivo Activities of Three Oxazolidinones against Nonreplicating Mycobacterium tuberculosis. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 3217-3223.	3.2	53
20	Genome-Wide Definition of the SigF Regulon in Mycobacterium tuberculosis. <i>Journal of Bacteriology</i> , 2012, 194, 2001-2009.	2.2	46
21	High-resolution detection of DNA binding sites of the global transcriptional regulator GlxR in <i>Corynebacterium glutamicum</i> . <i>Microbiology (United Kingdom)</i> , 2013, 159, 12-22.	1.8	44
22	High-resolution transcriptome and genome-wide dynamics of RNA polymerase and NusA in Mycobacterium tuberculosis. <i>Nucleic Acids Research</i> , 2013, 41, 961-977.	14.5	41
23	The Phosphatidyl- <i>myo</i> -Inositol Mannosyltransferase PimA Is Essential for Mycobacterium tuberculosis Growth <i>In Vitro</i> and <i>In Vivo</i> . <i>Journal of Bacteriology</i> , 2014, 196, 3441-3451.	2.2	37
24	Tuberculosis drugs: new candidates and how to find more. <i>Future Microbiology</i> , 2011, 6, 617-633.	2.0	36
25	Characterization of DprE1-Mediated Benzothiazinone Resistance in Mycobacterium tuberculosis. <i>Antimicrobial Agents and Chemotherapy</i> , 2016, 60, 6451-6459.	3.2	36
26	EspL is essential for virulence and stabilizes EspE, EspF and EspH levels in Mycobacterium tuberculosis. <i>PLoS Pathogens</i> , 2018, 14, e1007491.	4.7	33
27	Bioluminescence for Assessing Drug Potency against Nonreplicating Mycobacterium tuberculosis. <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 4012-4019.	3.2	30
28	<i>EspL</i> regulates the <i>ESX-1</i> secretion system in response to <i>ATP</i> levels in <i>Mycobacterium tuberculosis</i> . <i>Molecular Microbiology</i> , 2014, 93, 1057-1065.	2.5	27
29	CtrA Protein Rv3789 Is Required for Arabinosylation of Arabinogalactan in Mycobacterium tuberculosis. <i>Journal of Bacteriology</i> , 2015, 197, 3686-3697.	2.2	26
30	The <i>katG</i> mRNA of Mycobacterium tuberculosis and Mycobacterium smegmatis is processed at its 5' end and is stabilized by both a polypurine sequence and translation initiation. <i>BMC Molecular Biology</i> , 2008, 9, 33.	3.0	22
31	Essential Nucleoid Associated Protein mlHF (Rv1388) Controls Virulence and Housekeeping Genes in Mycobacterium tuberculosis. <i>Scientific Reports</i> , 2018, 8, 14214.	3.3	19
32	Genomic and transcriptomic analysis of the streptomycin-dependent Mycobacterium tuberculosis strain 18b. <i>BMC Genomics</i> , 2016, 17, 190.	2.8	18
33	Whole-Genome Sequencing for Comparative Genomics and De Novo Genome Assembly. <i>Methods in Molecular Biology</i> , 2015, 1285, 1-16.	0.9	15
34	Dissecting Regulatory Networks in Host-Pathogen Interaction Using ChIP-on-chip Technology. <i>Cell Host and Microbe</i> , 2009, 5, 430-437.	11.0	14
35	Sigma Factor F Does Not Prevent Rifampin Inhibition of RNA Polymerase or Cause Rifampin Tolerance in <i>Mycobacterium tuberculosis</i> . <i>Journal of Bacteriology</i> , 2010, 192, 5472-5479.	2.2	14
36	Promoter mutagenesis for fine-tuning expression of essential genes in <i>Mycobacterium tuberculosis</i> . <i>Microbial Biotechnology</i> , 2018, 11, 238-247.	4.2	13

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37	Mycobacterium ulcerans Mouse Model Refinement for Pre-Clinical Profiling of Vaccine Candidates. PLoS ONE, 2016, 11, e0167059.	2.5	12
38	Vaccines as remedy for antimicrobial resistance and emerging infections. Current Opinion in Immunology, 2020, 65, 102-106.	5.5	11
39	Multicenter analysis of sputum microbiota in tuberculosis patients. PLoS ONE, 2020, 15, e0240250.	2.5	10
40	Rv3852 (H-NS) of Mycobacterium tuberculosis Is Not Involved in Nucleoid Compaction and Virulence Regulation. Journal of Bacteriology, 2017, 199, .	2.2	9
41	Antibodies, epicenter of SARS-CoV-2 immunology. Cell Death and Differentiation, 2021, 28, 821-824.	11.2	9
42	Assessing essentiality of transketolase in Mycobacterium tuberculosis using an inducible protein degradation system. FEMS Microbiology Letters, 2014, 358, 30-35.	1.8	8
43	Polarly Localized EccE Is Required for ESX-1 Function and Stabilization of ESX-1 Membrane Proteins in Mycobacterium tuberculosis. Journal of Bacteriology, 2020, 202, .	2.2	7
44	Host-Directed Therapies and Anti-Virulence Compounds to Address Anti-Microbial Resistant Tuberculosis Infection. Applied Sciences (Switzerland), 2020, 10, 2688.	2.5	6
45	DNA replication in phage P4: Characterization of replicon II. Plasmid, 2006, 56, 216-222.	1.4	2
46	FasR Regulates Fatty Acid Biosynthesis and Is Essential for Virulence of Mycobacterium tuberculosis. Frontiers in Microbiology, 2020, 11, 586285.	3.5	1
47	Bacteriophage P4 sut1: a mutation suppressing transcription termination. Journal of General Virology, 2007, 88, 1041-1047.	2.9	0
48	Editorial on Special Issue "Tuberculosis Drug Discovery and Development 2019". Applied Sciences (Switzerland), 2020, 10, 6069.	2.5	0