## Tiago R D Costa

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8481933/publications.pdf

Version: 2024-02-01

25 papers

1,837 citations

623734 14 h-index 25 g-index

28 all docs

28 docs citations

28 times ranked 2399 citing authors

#	Article	IF	CITATIONS
1	Secretion systems in Gram-negative bacteria: structural and mechanistic insights. Nature Reviews Microbiology, 2015, 13, 343-359.	28.6	893
2	A comprehensive guide to pilus biogenesis in Gram-negative bacteria. Nature Reviews Microbiology, 2017, 15, 365-379.	28.6	221
3	Structure of the Bacterial Sex F Pilus Reveals an Assembly of a Stoichiometric Protein-Phospholipid Complex. Cell, 2016, 166, 1436-1444.e10.	28.9	122
4	Type IV secretion systems: Advances in structure, function, and activation. Molecular Microbiology, 2021, 115, 436-452.	2.5	119
5	Cryo-EM structure of the bacteria-killing type IV secretion system core complex from Xanthomonas citri. Nature Microbiology, 2018, 3, 1429-1440.	13.3	62
6	Cryo-EM structure of a type IV secretion system. Nature, 2022, 607, 191-196.	27.8	56
7	Use of chimeric type <scp>IV</scp> secretion systems to define contributions of outer membrane subassemblies for contactâ€dependent translocation. Molecular Microbiology, 2017, 105, 273-293.	2.5	49
8	The Cryoelectron Microscopy Structure of the Type 1 Chaperone-Usher Pilus Rod. Structure, 2017, 25, 1829-1838.e4.	3.3	46
9	Mating pair stabilization mediates bacterial conjugation species specificity. Nature Microbiology, 2022, 7, 1016-1027.	13.3	43
10	YopD Self-assembly and Binding to LcrV Facilitate Type III Secretion Activity by Yersinia pseudotuberculosis. Journal of Biological Chemistry, 2010, 285, 25269-25284.	3.4	24
11	The cryo-electron microscopy supramolecular structure of the bacterial stressosome unveils its mechanism of activation. Nature Communications, 2019, 10, 3005.	12.8	22
12	The Breadth and Molecular Basis of Hcp-Driven Type VI Secretion System Effector Delivery. MBio, 2021, 12, e0026221.	4.1	22
13	Cryo-EM Grid Preparation of Membrane Protein Samples for Single Particle Analysis. Frontiers in Molecular Biosciences, 2018, 5, 74.	3.5	19
14	The Legionella pneumophila Dot/Icm type IV secretion system and its effectors. Microbiology (United) Tj ETQq0	0 O <sub>1</sub> rgBT /0	Overlock 10 Tf
15	Type III secretion translocon assemblies that attenuate <i>Yersinia</i> virulence. Cellular Microbiology, 2013, 15, 1088-1110.	2.1	17
16	Structural Analysis of Protein Complexes by Cryo Electron Microscopy. Methods in Molecular Biology, 2017, 1615, 377-413.	0.9	15
17	Architecture of the outer-membrane core complex from a conjugative type IV secretion system. Nature Communications, 2021, 12, 6834.	12.8	15
18	Heterologous Complementation Studies With the YscX and YscY Protein Families Reveals a Specificity for Yersinia pseudotuberculosis Type III Secretion. Frontiers in Cellular and Infection Microbiology, 2018, 8, 80.	3.9	12

#	Article	IF	CITATIONS
19	Cryoelectron-microscopy structure of the enteropathogenic <i>Escherichia coli&lt;<math>l</math>i&gt; type III secretion system EspA filament. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .</i>	7.1	12
20	YopN and TyeA Hydrophobic Contacts Required for Regulating Ysc-Yop Type III Secretion Activity by Yersinia pseudotuberculosis. Frontiers in Cellular and Infection Microbiology, 2016, 6, 66.	3.9	11
21	Site-Directed Mutagenesis and Its Application in Studying the Interactions of T3S Components. Methods in Molecular Biology, 2017, 1531, 11-31.	0.9	11
22	Flagellin outer domain dimerization modulates motility in pathogenic and soil bacteria from viscous environments. Nature Communications, 2022, 13, 1422.	12.8	10
23	Coiled-coils in the YopD translocator family: A predicted structure unique to the YopD N-terminus contributes to full virulence of Yersinia pseudotuberculosis. Infection, Genetics and Evolution, 2012, 12, 1729-1742.	2.3	8
24	Genetically Engineered Frameshifted YopN-TyeA Chimeras Influence Type III Secretion System Function in Yersinia pseudotuberculosis. PLoS ONE, 2013, 8, e77767.	2.5	8
25	Measurement of Yersinia Translocon Pore Formation in Erythrocytes. Methods in Molecular Biology, 2019, 2010, 211-229.	0.9	2