

Yumi Inoue

List of Publications by Year in descending order

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Version: 2024-02-01

11
papers

309
citations

933447

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1281871

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docs citations

13
times ranked

184
citing authors

#	ARTICLE	IF	CITATIONS
1	The FlhA linker mediates flagellar protein export switching during flagellar assembly. <i>Communications Biology</i> , 2021, 4, 646.	4.4	16
2	FliK-Driven Conformational Rearrangements of FlhA and FlhB Are Required for Export Switching of the Flagellar Protein Export Apparatus. <i>Journal of Bacteriology</i> , 2020, 202, .	2.2	16
3	The flexible linker of the secreted FliK ruler is required for export switching of the flagellar protein export apparatus. <i>Scientific Reports</i> , 2020, 10, 838.	3.3	16
4	Structural Insights into the Substrate Specificity Switch Mechanism of the Type III Protein Export Apparatus. <i>Structure</i> , 2019, 27, 965-976.e6.	3.3	39
5	Mutational analysis of the C-terminal cytoplasmic domain of FlhB, a transmembrane component of the flagellar type III protein export apparatus in <i>Salmonella</i> . <i>Genes To Cells</i> , 2019, 24, 408-421.	1.2	11
6	Insight into structural remodeling of the FlhA ring responsible for bacterial flagellar type III protein export. <i>Science Advances</i> , 2018, 4, eaao7054.	10.3	50
7	Novel insights into the mechanism of well-ordered assembly of bacterial flagellar proteins in <i>Salmonella</i> . <i>Scientific Reports</i> , 2018, 8, 1787.	3.3	36
8	Straight and rigid flagellar hook made by insertion of the FlgG specific sequence into FlgE. <i>Scientific Reports</i> , 2017, 7, 46723.	3.3	27
9	The role of intrinsically disordered C-terminal region of FliK in substrate specificity switching of the bacterial flagellar type III export apparatus. <i>Molecular Microbiology</i> , 2017, 105, 572-588.	2.5	30
10	Structural stability of flagellin subunit affects the rate of flagellin export in the absence of FlhS chaperone. <i>Molecular Microbiology</i> , 2016, 102, 405-416.	2.5	32
11	FliH and FlhI ensure efficient energy coupling of flagellar type III protein export in <i>Salmonella</i> . <i>MicrobiologyOpen</i> , 2016, 5, 424-435.	3.0	36