Highâ€resolution insights into binding of unfolded poly SlpA

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Citation Report

#	Article	IF	Citations
1	Generation of a Highly Active Folding Enzyme by Combining a Parvulin-Type Prolyl Isomerase from SurA with an Unrelated Chaperone Domain. Journal of Molecular Biology, 2013, 425, 4089-4098.	2.0	9
2	Molecular function of the prolyl cis/trans isomerase and metallochaperone SlyD. Biological Chemistry, 2013, 394, 965-975.	1.2	21
3	Structural basis for PTPA interaction with the invariant C-terminal tail of PP2A. Biological Chemistry, 2014, 395, 881-889.	1.2	15
4	Elucidation of Abiotic Stress Signaling in Plants. , 2015, , .		5
5	Plant Immunophilins: A Protein Family with Diverse Functions Beyond Protein Folding Activity. , 2015, , 367-395.		1
6	Molecular insights into substrate recognition and catalytic mechanism of the chaperone and FKBP peptidyl-prolyl isomerase SlyD. BMC Biology, 2016, 14, 82.	1.7	26
7	Lactobacillus slpA promotes ESC growth through the ERK1/2 pathway. Cytotechnology, 2017, 69, 117-122.	0.7	1
8	Targeting the molecular chaperone SlyD to inhibit bacterial growth with a small molecule. Scientific Reports, 2017, 7, 42141.	1.6	12
9	Structure-Based Insights into the Dynamics and Function of Two-Domain SlpA from Escherichia coli. Biochemistry, 2017, 56, 6533-6543.	1.2	2
10	Identification of Substrates of Cytoplasmic Peptidyl-Prolyl Cis/Trans Isomerases and Their Collective Essentiality in Escherichia Coli. International Journal of Molecular Sciences, 2020, 21, 4212.	1.8	7