Rieko Ishima

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4032358/publications.pdf

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		1684188	1474206
13	82	5	9
papers	citations	h-index	g-index
13 all docs	13 docs citations	13 times ranked	125

#	Article	IF	CITATIONS
1	Relative domain orientation of the <scp>L289K HIV</scp> â€1 reverse transcriptase monomer. Protein Science, 2022, 31, e4307.	7.6	1
2	A synergy of activity, stability, and inhibitorâ€interaction of HIV â€1 protease mutants evolved under drugâ€pressure. Protein Science, 2021, 30, 571-582.	7.6	1
3	The reduced form of the antibody CH2 domain. Protein Science, 2021, 30, 1895-1903.	7.6	1
4	Retroviral RNase H: Structure, mechanism, and inhibition. The Enzymes, 2021, 50, 227-247.	1.7	4
5	Effect of Lysyl–tRNA Synthetase on the Maturation of HIV-1 Reverse Transcriptase. ACS Omega, 2020, 5, 16619-16627.	3.5	2
6	Large Multidomain Protein NMR: HIV-1 Reverse Transcriptase Precursor in Solution. International Journal of Molecular Sciences, 2020, 21, 9545.	4.1	1
7	Binding thermodynamics of metal ions to HIV-1 ribonuclease H domain. Journal of Thermal Analysis and Calorimetry, 2019, 135, 2647-2653.	3.6	8
8	19F NMR relaxation studies of fluorosubstituted tryptophans. Journal of Biomolecular NMR, 2019, 73, 401-409.	2.8	25
9	Determinants of Active-Site Inhibitor Interaction with HIV-1 RNase H. ACS Infectious Diseases, 2019, 5, 1963-1974.	3.8	10
10	NMR and MD studies combined to elucidate inhibitor and water interactions of HIV-1 protease and their modulations with resistance mutations. Journal of Biomolecular NMR, 2019, 73, 365-374.	2.8	9
11	Probing Structural Changes among Analogous Inhibitor-Bound Forms of HIV-1 Protease and a Drug-Resistant Mutant in Solution by Nuclear Magnetic Resonance. Biochemistry, 2018, 57, 1652-1662.	2.5	12
12	An NMR strategy to detect conformational differences in a protein complexed with highly analogous inhibitors in solution. Methods, 2018, 148, 9-18.	3.8	3
13	Entire-Dataset Analysis of NMR Fast-Exchange Titration Spectra: A Mg2+Titration Analysis for HIV-1 Ribonuclease H Domain. Journal of Physical Chemistry B, 2016, 120, 12420-12431.	2.6	5