Johannes van den Boom

List of Publications by Year in descending order

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#	Article	IF	Citations
1	Ubiquitin-directed AAA+ ATPase p97/VCP unfolds stable proteins crosslinked to DNA for proteolysis by SPRTN. Journal of Biological Chemistry, 2022, 298, 101976.	3.4	13
2	Targeted substrate loop insertion by VCP/p97 during PP1 complex disassembly. Nature Structural and Molecular Biology, 2021, 28, 964-971.	8.2	16
3	Protein Phosphatase-1 Complex Disassembly by p97 is Initiated through Multivalent Recognition of Catalytic and Regulatory Subunits by the p97 SEP-domain Adapters. Journal of Molecular Biology, 2020, 432, 6061-6074.	4.2	20
4	The chaperone Lhs1 contributes to the virulence of the fish-pathogenic oomycete Aphanomyces invadans. Fungal Biology, 2020, 124, 1024-1031.	2.5	5
5	The other side of the corona: nanoparticles inhibit the protease taspase1 in a size-dependent manner. Nanoscale, 2020, 12, 19093-19103.	5.6	7
6	Structure of the PUB Domain from Ubiquitin Regulatory X Domain Protein 1 (UBXD1) and Its Interaction with the p97 AAA+ ATPase. Biomolecules, 2019, 9, 876.	4.0	5
7	A Nonâ€Competitive Inhibitor of VCP/p97 and VPS4 Reveals Conserved Allosteric Circuits in Type I and II AAA ATPases. Angewandte Chemie - International Edition, 2018, 57, 1576-1580.	13.8	23
8	VCP/p97-Mediated Unfolding as a Principle in Protein Homeostasis and Signaling. Molecular Cell, 2018, 69, 182-194.	9.7	288
9	Ubiquitin-Independent Disassembly by a p97 AAA-ATPase Complex Drives PP1 Holoenzyme Formation. Molecular Cell, 2018, 72, 766-777.e6.	9.7	62
10	Human DHEA sulfation requires direct interaction between PAPS synthase 2 and DHEA sulfotransferase SULT2A1. Journal of Biological Chemistry, 2018, 293, 9724-9735.	3.4	29
11	Structural Characterization of the Loop at the Alpha-Subunit C-Terminus of the Mixed Lineage Leukemia Protein Activating Protease Taspase1. PLoS ONE, 2016, 11, e0151431.	2.5	8
12	VCP/p97 Extracts Sterically Trapped Ku70/80 Rings from DNA in Double-Strand Break Repair. Molecular Cell, 2016, 64, 189-198.	9.7	91
13	Structural Model of the Bilitranslocase Transmembrane Domain Supported by NMR and FRET Data. PLoS ONE, 2015, 10, e0135455.	2.5	8
14	3′-Phosphoadenosine 5′-Phosphosulfate (PAPS) Synthases, Naturally Fragile Enzymes Specifically Stabilized by Nucleotide Binding. Journal of Biological Chemistry, 2012, 287, 17645-17655.	3.4	37