Melanie Schwarten

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Guidelines for the use and interpretation of assays for monitoring autophagy (3rd edition). Autophagy, 2016, 12, 1-222.	9.1	4,701
2	Nix directly binds to GABARAP: A possible crosstalk between apoptosis and autophagy. Autophagy, 2009, 5, 690-698.	9.1	212
3	BEST-TROSY experiments for time-efficient sequential resonance assignment of large disordered proteins. Journal of Biomolecular NMR, 2013, 55, 311-321.	2.8	193
4	Biochemical characterisation of TCTP questions its function as a guanine nucleotide exchange factor for Rheb. FEBS Letters, 2008, 582, 3005-3010.	2.8	76
5	Ras Homolog Enriched in Brain (Rheb) Enhances Apoptotic Signaling*. Journal of Biological Chemistry, 2010, 285, 33979-33991.	3.4	49
6	Transient Structure and SH3 Interaction Sites in an Intrinsically Disordered Fragment of the Hepatitis C Virus Protein NS5A. Journal of Molecular Biology, 2012, 420, 310-323.	4.2	49
7	Mechanism-based inhibition of an aldolase at high concentrations of its natural substrate acetaldehyde: structural insights and protective strategies. Chemical Science, 2016, 7, 4492-4502.	7.4	49
8	The Atg8 Family of Proteins—Modulating Shape and Functionality of Autophagic Membranes. Frontiers in Genetics, 2017, 8, 109.	2.3	36
9	Structure and potential function of γ â€aminobutyrate type A receptorâ€associated protein. FEBS Journal, 2009, 276, 4989-5005.	4.7	33
10	Pyroglutamate-modified Aβ(3-42) affects aggregation kinetics of Aβ(1-42) by accelerating primary and secondary pathways. Chemical Science, 2017, 8, 4996-5004.	7.4	33
11	Interaction of Bcl-2 with the Autophagy-related GABAA Receptor-associated Protein (GABARAP). Journal of Biological Chemistry, 2013, 288, 37204-37215.	3.4	27
12	Structural Analysis and Aggregation Propensity of Pyroglutamate Aβ(3-40) in Aqueous Trifluoroethanol. PLoS ONE, 2015, 10, e0143647.	2.5	27
13	Preparation of a Functional GABARAP–Lipid Conjugate in Nanodiscs and its Investigation by Solution NMR Spectroscopy. ChemBioChem, 2010, 11, 1967-1970.	2.6	26
14	The mammalian autophagy initiator complex contains 2 HORMA domain proteins. Autophagy, 2015, 11, 2300-2308.	9.1	26
15	Solution structure of Atg8 reveals conformational polymorphism of the N-terminal domain. Biochemical and Biophysical Research Communications, 2010, 395, 426-431.	2.1	23
16	Pyroglutamate-Modified Amyloid-β(3–42) Shows α-Helical Intermediates before Amyloid Formation. Biophysical Journal, 2017, 112, 1621-1633.	0.5	22
17	Interaction of Nonstructural Protein 5A of the Hepatitis C Virus with Src Homology 3 Domains Using Noncanonical Binding Sites. Biochemistry, 2013, 52, 6160-6168.	2.5	21
18	Amino Terminal Region of Dengue Virus NS4A Cytosolic Domain Binds to Highly Curved Liposomes. Viruses, 2015, 7, 4119-4130.	3.3	21

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19	Purification and Characterization of Recombinant N-Terminally Pyroglutamate-Modified Amyloid-β Variants and Structural Analysis by Solution NMR Spectroscopy. PLoS ONE, 2015, 10, e0139710.	2.5	21
20	Dengue virus NS4A cytoplasmic domain binding to liposomes is sensitive to membrane curvature. Biochimica Et Biophysica Acta - Biomembranes, 2015, 1848, 1119-1126.	2.6	20
21	The Disordered Region of the HCV Protein NS5A: Conformational Dynamics, SH3 Binding, and Phosphorylation. Biophysical Journal, 2015, 109, 1483-1496.	0.5	19
22	Comparative modeling of human NSF reveals a possible binding mode of GABARAP and GATEâ€16. Proteins: Structure, Function and Bioinformatics, 2009, 77, 637-646.	2.6	17
23	Conformational Polymorphism in Autophagy-Related Protein GATE-16. Biochemistry, 2015, 54, 5469-5479.	2.5	17
24	Direct binding to GABARAP family members is essential for HIV-1 Nef plasma membrane localization. Scientific Reports, 2017, 7, 5979.	3.3	11
25	Sequence-specific 1H, 13C, and 15N backbone assignment of the GTPase rRheb in its GDP-bound form. Biomolecular NMR Assignments, 2007, 1, 45-47.	0.8	10
26	Sequence-specific 1H, 13C, and 15N backbone assignment of the activated 21 kDa GTPase rRheb. Biomolecular NMR Assignments, 2007, 1, 105-108.	0.8	8
27	Hepatitis C virus NS5A is able to competitively displace c-Myc from the Bin1 SH3 domain <i>in vitro</i> . Journal of Peptide Science, 2014, 20, 334-340.	1.4	8
28	Sequence-specific 1H, 13C, and 15N resonance assignment of the autophagy-related protein Atg8. Biomolecular NMR Assignments, 2009, 3, 137-139.	0.8	5
29	Investigating Structure and Dynamics of Atg8 Family Proteins. Methods in Enzymology, 2017, 587, 115-142.	1.0	5
30	Deficiency of GABARAP but Not Its Paralogs Causes Enhanced EGF-Induced EGFR Degradation. Cells, 2020, 9, 1296.	4.1	3
31	Sequence-specific 1H, 15N, and 13C resonance assignments of the autophagy-related protein LC3C. Biomolecular NMR Assignments, 2016, 10, 41-43.	0.8	2
32	Structural Studies of Autophagy-Related Proteins. Methods in Molecular Biology, 2019, 1880, 17-56.	0.9	2
33	Analysis of the Bin1 SH3 interaction with peptides derived from the hepatitis C virus protein NS5A and c-Myc reveals that NS5A can competitively displace c-Myc in vitro. European Journal of Medical Research, 2014, 19, S10.	2.2	1
34	In Vitro Reconstitution of the Highly Active and Natively Folded Recombinant Human Superoxide Dismutase 1 Holoenzyme. ChemistrySelect, 2018, 3, 7627-7632.	1.5	1
35	The non-structural protein 5A (NS5A) of hepatitis C virus interacts with the SH3 domain of human Bin1 using non-canonical binding sites. European Journal of Medical Research, 2014, 19, .	2.2	0
36	Cytoplasmic Domain of Dengue Virus Protein NS4A Preferentially Binds Highly Curved Membranes. Biophysical Journal, 2015, 108, 246a.	0.5	0