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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	The paradox of autophagy and its implication in cancer etiology and therapy. Apoptosis: an International Journal on Programmed Cell Death, 2009, 14, 376-391.	4.9	192
3	The Dependence Receptor UNC5H2/B Triggers Apoptosis via PP2A-Mediated Dephosphorylation of DAP Kinase. Molecular Cell, 2010, 40, 863-876.	9.7	111
4	Golgi organization is regulated by proteasomal degradation. Nature Communications, 2020, 11, 409.	12.8	73
5	Revealing the cellular degradome by mass spectrometry analysis of proteasome-cleaved peptides. Nature Biotechnology, 2018, 36, 1110-1116.	17.5	33
6	Metabolic alterations in the tumor microenvironment and their role in oncogenesis. Cancer Letters, 2020, 484, 65-71.	7.2	32
7	Post-translational modification profiling – A novel tool for mapping the protein modification landscape in cancer. Experimental Biology and Medicine, 2016, 241, 1475-1482.	2.4	21
8	SUMOylation of linker histone H1 drives chromatin condensation and restriction of embryonic cell fate identity. Molecular Cell, 2022, 82, 106-122.e9.	9.7	19
9	PKD at the crossroads of necrosis and autophagy. Autophagy, 2012, 8, 433-434.	9.1	15
10	DAPk silencing by DNA methylation conveys resistance to anti EGFR drugs in lung cancer cells. Cell Cycle, 2012, 11, 2051-2051.	2.6	11
11	Histone degradation by the proteasome regulates chromatin and cellular plasticity. FEBS Journal, 2022, 289, 3304-3316.	4.7	11
12	Maintaining Golgi Homeostasis: A Balancing Act of Two Proteolytic Pathways. Cells, 2022, 11, 780.	4.1	6
13	Gatekeepers of the Gut: The Roles of Proteasomes at the Gastrointestinal Barrier. Biomolecules, 2021, 11, 989.	4.0	5
14	Altered Protein Abundance and Localization Inferred from Sites of Alternative Modification by Ubiquitin and SUMO. Journal of Molecular Biology, 2021, 433, 167219.	4.2	4
15	Phenotypic Screen Identifies JAK2 as a Major Regulator of FAT10 Expression. ACS Chemical Biology, 2019, 14, 2538-2545.	3.4	3
16	Post-Translational Modification Profiling-Functional Proteomics for the Analysis of Immune Regulation. Methods in Molecular Biology, 2017, 1647, 139-152.	0.9	2