

# Ugo Mayor

## List of Publications by Year in descending order

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Version: 2024-02-01

41  
papers

2,661  
citations

331259

21  
h-index

301761

39  
g-index

46  
all docs

46  
docs citations

46  
times ranked

3296  
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of substrates for human deubiquitinating enzymes (DUBs): An up-to-date review and a case study for neurodevelopmental disorders. <i>Seminars in Cell and Developmental Biology</i> , 2022, 132, 120-131.	2.3	4
2	A Proteomic Approach for Systematic Mapping of Substrates of Human Deubiquitinating Enzymes. <i>International Journal of Molecular Sciences</i> , 2021, 22, 4851.	1.8	6
3	Solvent-Based Elimination of Organic Matter from Marine-Collected Plastics. <i>Environments - MDPI</i> , 2021, 8, 68.	1.5	3
4	Neddylation inhibition ameliorates steatosis in NAFLD by boosting hepatic fatty acid oxidation via the DEPTOR-mTOR axis. <i>Molecular Metabolism</i> , 2021, 53, 101275.	3.0	22
5	The ubiquitin ligase Ariadne-1 regulates neurotransmitter release via ubiquitination of NSF. <i>Journal of Biological Chemistry</i> , 2021, 296, 100408.	1.6	6
6	Identification of proximal SUMO-dependent interactors using SUMO-ID. <i>Nature Communications</i> , 2021, 12, 6671.	5.8	27
7	Multi-Omics Integration Highlights the Role of Ubiquitination in CCl4-Induced Liver Fibrosis. <i>International Journal of Molecular Sciences</i> , 2020, 21, 9043.	1.8	12
8	SUMOylation in the control of cholesterol homeostasis. <i>Open Biology</i> , 2020, 10, 200054.	1.5	14
9	How to Inactivate Human Ubiquitin E3 Ligases by Mutation. <i>Frontiers in Cell and Developmental Biology</i> , 2020, 8, 39.	1.8	31
10	The role of SUMOylation during development. <i>Biochemical Society Transactions</i> , 2020, 48, 463-478.	1.6	27
11	Mass Spectrometry-Based Characterization of Ub- and UbL-Modified Proteins. <i>Methods in Molecular Biology</i> , 2020, 2051, 265-276.	0.4	3
12	Detailed Dissection of UBE3A-Mediated DDI1 Ubiquitination. <i>Frontiers in Physiology</i> , 2019, 10, 534.	1.3	17
13	Impaired proteostasis in rare neurological diseases. <i>Seminars in Cell and Developmental Biology</i> , 2019, 93, 164-177.	2.3	14
14	Basal mitophagy is widespread in <i>Drosophila</i> but minimally affected by loss of Pink1 or parkin. <i>Journal of Cell Biology</i> , 2018, 217, 1613-1622.	2.3	253
15	Quantitative proteomics reveals neuronal ubiquitination of Rngo/Ddi1 and several proteasomal subunits by Ube3a, accounting for the complexity of Angelman syndrome. <i>Human Molecular Genetics</i> , 2018, 27, 1955-1971.	1.4	30
16	Ubiquitylation Dynamics of the Clock Cell Proteome and TIMELESS during a Circadian Cycle. <i>Cell Reports</i> , 2018, 23, 2273-2282.	2.9	29
17	Neuronal Proteomic Analysis of the Ubiquitinated Substrates of the Disease-Linked E3 Ligases Parkin and Ube3a. <i>BioMed Research International</i> , 2018, 2018, 1-14.	0.9	12
18	A comprehensive platform for the analysis of ubiquitin-like protein modifications using in vivo biotinylation. <i>Scientific Reports</i> , 2017, 7, 40756.	1.6	58

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19	Quantitative proteomic analysis of Parkin substrates in Drosophila neurons. <i>Molecular Neurodegeneration</i> , 2017, 12, 29.	4.4	77
20	In Vivo Strategies to Isolate and Characterize the Neuronal Ubiquitinated Proteome. <i>Neuromethods</i> , 2017, , 179-189.	0.2	2
21	Multi-story Parkin. <i>Oncotarget</i> , 2017, 8, 50327-50328.	0.8	2
22	Analysis of SUMOylated Proteins in Cells and In Vivo Using the bioSUMO Strategy. <i>Methods in Molecular Biology</i> , 2016, 1475, 161-169.	0.4	4
23	Isolation of Ubiquitinated Proteins to High Purity from In Vivo Samples. <i>Methods in Molecular Biology</i> , 2016, 1449, 193-202.	0.4	8
24	Proteomic Analysis of the Ubiquitin Landscape in the Drosophila Embryonic Nervous System and the Adult Photoreceptor Cells. <i>PLoS ONE</i> , 2015, 10, e0139083.	1.1	39
25	<scp>USP</scp> 30 deubiquitylates mitochondrial <scp>P</scp> arkin substrates and restricts apoptotic cell death. <i>EMBO Reports</i> , 2015, 16, 618-627.	2.0	136
26	Ube3a, the E3 ubiquitin ligase causing Angelman syndrome and linked to autism, regulates protein homeostasis through the proteasomal shuttle Rpn10. <i>Cellular and Molecular Life Sciences</i> , 2014, 71, 2747-2758.	2.4	77
27	Using in Vivo Biotinylated Ubiquitin to Describe a Mitotic Exit Ubiquitome from Human Cells. <i>Molecular and Cellular Proteomics</i> , 2014, 13, 2411-2425.	2.5	37
28	Ubiquitin Profiling in Liver Using a Transgenic Mouse with Biotinylated Ubiquitin. <i>Journal of Proteome Research</i> , 2014, 13, 3016-3026.	1.8	31
29	Scavenger Receptors Mediate the Role of SUMO and Ftz-f1 in Drosophila Steroidogenesis. <i>PLoS Genetics</i> , 2013, 9, e1003473.	1.5	58
30	Ubiquitination site preferences in anaphase promoting complex/cyclosome (APC/C) substrates. <i>Open Biology</i> , 2013, 3, 130097.	1.5	39
31	Deciphering Tissue-Specific Ubiquitylation by Mass Spectrometry. <i>Methods in Molecular Biology</i> , 2012, 832, 65-80.	0.4	11
32	A Novel Strategy to Isolate Ubiquitin Conjugates Reveals Wide Role for Ubiquitination during Neural Development. <i>Molecular and Cellular Proteomics</i> , 2011, 10, M110.002188.	2.5	77
33	Solution structure of a protein denatured state and folding intermediate. <i>Nature</i> , 2005, 437, 1053-1056.	13.7	233
34	The complete folding pathway of a protein from nanoseconds to microseconds. <i>Nature</i> , 2003, 421, 863-867.	13.7	449
35	The Denatured State of Engrailed Homeodomain under Denaturing and Native Conditions. <i>Journal of Molecular Biology</i> , 2003, 333, 977-991.	2.0	88
36	Unifying features in protein-folding mechanisms. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 13286-13291.	3.3	225

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37	Crystal Structures of Engrailed Homeodomain Mutants. Journal of Biological Chemistry, 2003, 278, 43699-43708.	1.6	39
38	Structural insights in the folding of small single-domain proteins. Italian Journal of Biochemistry, 2003, 52, 154-61.	0.3	0
39	Protein folding and unfolding in microseconds to nanoseconds by experiment and simulation. Proceedings of the National Academy of Sciences of the United States of America, 2000, 97, 13518-13522.	3.3	303
40	From snapshot to movie: phi analysis of protein folding transition states taken one step further. Proceedings of the National Academy of Sciences of the United States of America, 1999, 96, 14854-14859.	3.3	145
41	Konpartimentu-espezifikoko gertuko biotinizazioa: XPO1en esportazio-kargoak identifikatzeko hurbilketa berria. Ekaia (journal), 0, , .	0.0	0