

# Xiang Chen

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2411664/publications.pdf>

Version: 2024-02-01

25  
papers

2,101  
citations

567281

15  
h-index

642732

23  
g-index

26  
all docs

26  
docs citations

26  
times ranked

2048  
citing authors

#	ARTICLE	IF	CITATIONS
1	Proteasome subunit Rpn13 is a novel ubiquitin receptor. <i>Nature</i> , 2008, 453, 481-488.	27.8	553
2	Ubiquitin docking at the proteasome through a novel pleckstrin-homology domain interaction. <i>Nature</i> , 2008, 453, 548-552.	27.8	290
3	Rpn1 provides adjacent receptor sites for substrate binding and deubiquitination by the proteasome. <i>Science</i> , 2016, 351, .	12.6	234
4	Gates, Channels, and Switches: Elements of the Proteasome Machine. <i>Trends in Biochemical Sciences</i> , 2016, 41, 77-93.	7.5	223
5	A bis-Benzylidene Piperidone Targeting Proteasome Ubiquitin Receptor RPN13/ADRM1 as a Therapy for Cancer. <i>Cancer Cell</i> , 2013, 24, 791-805.	16.8	137
6	Chemical and structural studies provide a mechanistic basis for recognition of the MYC G-quadruplex. <i>Nature Communications</i> , 2018, 9, 4229.	12.8	131
7	Structure of Proteasome Ubiquitin Receptor hRpn13 and Its Activation by the Scaffolding Protein hRpn2. <i>Molecular Cell</i> , 2010, 38, 404-415.	9.7	102
8	Prokaryotic Ubiquitin-Like Protein Pup Is Intrinsically Disordered. <i>Journal of Molecular Biology</i> , 2009, 392, 208-217.	4.2	97
9	Structures of Rpn1 T1:Rad23 and hRpn13:hPLIC2 Reveal Distinct Binding Mechanisms between Substrate Receptors and Shuttle Factors of the Proteasome. <i>Structure</i> , 2016, 24, 1257-1270.	3.3	72
10	Structure of hRpn10 Bound to UBQLN2 UBL Illustrates Basis for Complementarity between Shuttle Factors and Substrates at the Proteasome. <i>Journal of Molecular Biology</i> , 2019, 431, 939-955.	4.2	41
11	Proteasome interaction with ubiquitinated substrates: from mechanisms to therapies. <i>FEBS Journal</i> , 2021, 288, 5231-5251.	4.7	40
12	Structure of E3 ligase E6AP with a proteasome-binding site provided by substrate receptor hRpn10. <i>Nature Communications</i> , 2020, 11, 1291.	12.8	29
13	Covalent Rpn13-Binding Inhibitors for the Treatment of Ovarian Cancer. <i>ACS Omega</i> , 2018, 3, 11917-11929.	3.5	25
14	Structural Plasticity Allows UCH37 to Be Primed by RPN13 or Locked Down by INO80G. <i>Molecular Cell</i> , 2015, 57, 767-768.	9.7	21
15	Structure-guided bifunctional molecules hit a DEUBAD-lacking hRpn13 species upregulated in multiple myeloma. <i>Nature Communications</i> , 2021, 12, 7318.	12.8	18
16	Identifying and Studying Ubiquitin Receptors by NMR. <i>Methods in Molecular Biology</i> , 2012, 832, 279-303.	0.9	17
17	Novel TDP2-ubiquitin interactions and their importance for the repair of topoisomerase II-mediated DNA damage. <i>Nucleic Acids Research</i> , 2016, 44, gkw719.	14.5	17
18	Cryo-EM Reveals Unanchored M1-Ubiquitin Chain Binding at hRpn11 of the 26S Proteasome. <i>Structure</i> , 2020, 28, 1206-1217.e4.	3.3	17

#	ARTICLE	IF	CITATIONS
19	Metabolic plasticity of IDH1-mutant glioma cell lines is responsible for low sensitivity to glutaminase inhibition. <i>Cancer &amp; Metabolism</i> , 2020, 8, 23.	5.0	14
20	Measuring ubiquitin chain linkage: Rap80 uses a molecular ruler mechanism for ubiquitin linkage specificity. <i>EMBO Journal</i> , 2009, 28, 2307-2308.	7.8	12
21	Nuclear destruction: A suicide mission by AKIRIN2 brings intact proteasomes into the nucleus. <i>Molecular Cell</i> , 2022, 82, 13-14.	9.7	5
22	The CD8 $\zeta$ hinge is intrinsically disordered with a dynamic exchange that includes proline cis-trans isomerization. <i>Journal of Magnetic Resonance</i> , 2022, 340, 107234.	2.1	5
23	<sup>1</sup> H, <sup>15</sup> N, <sup>13</sup> C resonance assignments for <i>Saccharomyces cerevisiae</i> Rad23 UBL domain. <i>Biomolecular NMR Assignments</i> , 2016, 10, 291-295.	0.8	1
24	An optimized protocol for acquiring and processing cryo-EM data of human 26S proteasome with M1-Ub6. <i>STAR Protocols</i> , 2021, 2, 100278.	1.2	0
25	Ubiquitination. , 2020, , 1-11.		0