

# Stefan Jentsch

## List of Publications by Year in descending order

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27  
papers

3,538  
citations

394421

19  
h-index

552781

26  
g-index

32  
all docs

32  
docs citations

32  
times ranked

6677  
citing authors

#	ARTICLE	IF	CITATIONS
1	Nucleolar release of rDNA repeats for repair involves SUMO-mediated untethering by the Cdc48/p97 segregase. <i>Nature Communications</i> , 2021, 12, 4918.	12.8	12
2	A Selective Autophagy Pathway for Phase-Separated Endocytic Protein Deposits. <i>Molecular Cell</i> , 2020, 80, 764-778.e7.	9.7	82
3	ESCRT recruitment by the inner nuclear membrane protein Heh1 is regulated by Hub1-mediated alternative splicing. <i>Journal of Cell Science</i> , 2020, 133, .	2.0	14
4	Chaperone-Mediated Protein Disaggregation Triggers Proteolytic Clearance of Intra-nuclear Protein Inclusions. <i>Cell Reports</i> , 2020, 31, 107680.	6.4	43
5	Selective autophagy degrades nuclear pore complexes. <i>Nature Cell Biology</i> , 2020, 22, 159-166.	10.3	86
6	Slx5/Slx8-dependent ubiquitin hotspots on chromatin contribute to stress tolerance. <i>EMBO Journal</i> , 2019, 38, .	7.8	8
7	A SUMO-dependent pathway controls elongating RNA Polymerase II upon UV-induced damage. <i>Scientific Reports</i> , 2019, 9, 17914.	3.3	5
8	Receptor oligomerization guides pathway choice between proteasomal and autophagic degradation. <i>Nature Cell Biology</i> , 2017, 19, 732-739.	10.3	75
9	The INO80 Complex Removes H2A.Z to Promote Presynaptic Filament Formation during Homologous Recombination. <i>Cell Reports</i> , 2017, 19, 1294-1303.	6.4	51
10	Error-Prone Splicing Controlled by the Ubiquitin Relative Hub1. <i>Molecular Cell</i> , 2017, 67, 423-432.e4.	9.7	22
11	Pathway choice between proteasomal and autophagic degradation. <i>Autophagy</i> , 2017, 13, 1799-1800.	9.1	32
12	Identification of Substrates of Protein-Group SUMOylation. <i>Methods in Molecular Biology</i> , 2016, 1475, 219-231.	0.9	17
13	DNA-protein crosslink repair: proteases as DNA repair enzymes. <i>Trends in Biochemical Sciences</i> , 2015, 40, 67-71.	7.5	81
14	DNA-protein crosslink repair. <i>Nature Reviews Molecular Cell Biology</i> , 2015, 16, 455-460.	37.0	75
15	Mechanisms and principles of homology search during recombination. <i>Nature Reviews Molecular Cell Biology</i> , 2014, 15, 369-383.	37.0	153
16	The conserved ubiquitin-like protein Hub1 plays a critical role in splicing in human cells. <i>Journal of Molecular Cell Biology</i> , 2014, 6, 312-323.	3.3	30
17	A DNA-Dependent Protease Involved in DNA-Protein Crosslink Repair. <i>Cell</i> , 2014, 158, 327-338.	28.9	218
18	Autophagic Clearance of PolyQ Proteins Mediated by Ubiquitin-Atg8 Adaptors of the Conserved CUET Protein Family. <i>Cell</i> , 2014, 158, 549-563.	28.9	285

#	ARTICLE	IF	CITATIONS
19	Control of Nuclear Activities by Substrate-Selective and Protein-Group SUMOylation. Annual Review of Genetics, 2013, 47, 167-186.	7.6	214
20	Protein Group Modification and Synergy in the SUMO Pathway as Exemplified in DNA Repair. Cell, 2012, 151, 807-820.	28.9	404
21	Role of the ubiquitin-like protein Hub1 in splice-site usage and alternative splicing. Nature, 2011, 474, 173-178.	27.8	79
22	Travels with ubiquitin: from protein degradation to DNA repair. EMBO Molecular Medicine, 2011, 3, 72-74.	6.9	0
23	Regulatory Functions of Ubiquitin and SUMO in DNA Repair Pathways. Sub-Cellular Biochemistry, 2010, 54, 184-194.	2.4	7
24	Cdc48 (p97): a "molecular gearbox"™ in the ubiquitin pathway?. Trends in Biochemical Sciences, 2007, 32, 6-11.	7.5	264
25	The ubiquitin-like protein HUB1 forms SDS-resistant complexes with cellular proteins in the absence of ATP. EMBO Reports, 2003, 4, 1169-1174.	4.5	54
26	Sumo, ubiquitin's mysterious cousin. Nature Reviews Molecular Cell Biology, 2001, 2, 202-210.	37.0	685
27	Activation of a Membrane-Bound Transcription Factor by Regulated Ubiquitin/Proteasome-Dependent Processing. Cell, 2000, 102, 577-586.	28.9	540