

# Felix Mueller-Planitz

## List of Publications by Year in descending order

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Version: 2024-02-01

26  
papers

902  
citations

471509

17  
h-index

580821

25  
g-index

29  
all docs

29  
docs citations

29  
times ranked

1486  
citing authors

#	ARTICLE	IF	CITATIONS
1	A case of convergent evolution: Several viral and bacterial pathogens hijack RSK kinases through a common linear motif. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	7.1	14
2	A critical role for linker DNA in higher-order folding of chromatin fibers. <i>Nucleic Acids Research</i> , 2021, 49, 2537-2551.	14.5	19
3	Nucleosome Positioning and Spacing: From Mechanism to Function. <i>Journal of Molecular Biology</i> , 2021, 433, 166847.	4.2	26
4	A CDK-regulated chromatin segregase promoting chromosome replication. <i>Nature Communications</i> , 2021, 12, 5224.	12.8	6
5	The biogenesis and function of nucleosome arrays. <i>Nature Communications</i> , 2021, 12, 7011.	12.8	12
6	Structural Architecture of the Nucleosome Remodeler ISWI Determined from Cross-Linking, Mass Spectrometry, SAXS, and Modeling. <i>Structure</i> , 2018, 26, 282-294.e6.	3.3	11
7	Remodeling and Repositioning of Nucleosomes in Nucleosomal Arrays. <i>Methods in Molecular Biology</i> , 2018, 1805, 349-370.	0.9	5
8	Myosin Va's adaptor protein melanophilin enforces track selection on the microtubule and actin networks in vitro. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E4714-E4723.	7.1	28
9	Kinesin-2 motors adapt their stepping behavior for processive transport on axonemes and microtubules. <i>EMBO Reports</i> , 2017, 18, 1947-1956.	4.5	26
10	A Poly-ADP-Ribose Trigger Releases the Auto-Inhibition of a Chromatin Remodeling Oncogene. <i>Molecular Cell</i> , 2017, 68, 860-871.e7.	9.7	70
11	Concerted regulation of ISWI by an autoinhibitory domain and the H4 N-terminal tail. <i>ELife</i> , 2017, 6, .	6.0	28
12	Integrative Modeling of the ISWI Chromatin Remodeling Enzyme from Cross-Linking/Mass Spectrometry and Saxs Data. <i>Biophysical Journal</i> , 2016, 110, 237a.	0.5	0
13	Crossfinder-assisted mapping of protein crosslinks formed by site-specifically incorporated crosslinkers. <i>Bioinformatics</i> , 2015, 31, 2043-2045.	4.1	18
14	Nucleosome Spacing Generated by ISWI and CHD1 Remodelers Is Constant Regardless of Nucleosome Density. <i>Molecular and Cellular Biology</i> , 2015, 35, 1588-1605.	2.3	52
15	ISWI Remodelling of Physiological Chromatin Fibres Acetylated at Lysine 16 of Histone H4. <i>PLoS ONE</i> , 2014, 9, e88411.	2.5	24
16	Rapid Purification of Recombinant Histones. <i>PLoS ONE</i> , 2014, 9, e104029.	2.5	45
17	Nucleosome sliding mechanisms: new twists in a looped history. <i>Nature Structural and Molecular Biology</i> , 2013, 20, 1026-1032.	8.2	92
18	No need for a power stroke in ISWI-mediated nucleosome sliding. <i>EMBO Reports</i> , 2013, 14, 1092-1097.	4.5	18

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19	The Myosin Chaperone UNC-45 Is Organized in Tandem Modules to Support Myofilament Formation in <i>C.Âlegans</i> . <i>Cell</i> , 2013, 152, 183-195.	28.9	94
20	The ATPase domain of ISWI is an autonomous nucleosome remodeling machine. <i>Nature Structural and Molecular Biology</i> , 2013, 20, 82-89.	8.2	77
21	Probing the Conformation of the ISWI ATPase Domain With Genetically Encoded Photoreactive Crosslinkers and Mass Spectrometry. <i>Molecular and Cellular Proteomics</i> , 2012, 11, M111.012088.	3.8	45
22	Regulation of a heterodimeric kinesin-2 through an unprocessive motor domain that is turned processive by its partner. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 10460-10465.	7.1	62
23	The DNA binding CXC domain of MSL2 is required for faithful targeting the Dosage Compensation Complex to the X chromosome. <i>Nucleic Acids Research</i> , 2010, 38, 3209-3221.	14.5	65
24	Coupling between ATP Binding and DNA Cleavage by DNA Topoisomerase II. <i>Journal of Biological Chemistry</i> , 2008, 283, 17463-17476.	3.4	23
25	DNA topoisomerase II selects DNA cleavage sites based on reactivity rather than binding affinity. <i>Nucleic Acids Research</i> , 2007, 35, 3764-3773.	14.5	23
26	Interdomain Communication in DNA Topoisomerase II. <i>Journal of Biological Chemistry</i> , 2006, 281, 23395-23404.	3.4	18