

Frank Uhlmann

List of Publications by Year in descending order

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121
papers

12,285
citations

26610

56
h-index

29127

104
g-index

129
all docs

129
docs citations

129
times ranked

6659
citing authors

#	ARTICLE	IF	CITATIONS
1	Sister-chromatid separation at anaphase onset is promoted by cleavage of the cohesin subunit Scc1. <i>Nature</i> , 1999, 400, 37-42.	13.7	882
2	Cleavage of Cohesin by the CD Clan Protease Separin Triggers Anaphase in Yeast. <i>Cell</i> , 2000, 103, 375-386.	13.5	765
3	Cohesin relocation from sites of chromosomal loading to places of convergent transcription. <i>Nature</i> , 2004, 430, 573-578.	13.7	544
4	SMC complexes: from DNA to chromosomes. <i>Nature Reviews Molecular Cell Biology</i> , 2016, 17, 399-412.	16.1	455
5	Eco1-Dependent Cohesin Acetylation During Establishment of Sister Chromatid Cohesion. <i>Science</i> , 2008, 321, 563-566.	6.0	453
6	Cohesion between sister chromatids must be established during DNA replication. <i>Current Biology</i> , 1998, 8, 1095-1102.	1.8	431
7	Disjunction of Homologous Chromosomes in Meiosis I Depends on Proteolytic Cleavage of the Meiotic Cohesin Rec8 by Separin. <i>Cell</i> , 2000, 103, 387-398.	13.5	418
8	Splitting the Chromosome: Cutting the Ties That Bind Sister Chromatids. <i>Science</i> , 2000, 288, 1379-1384.	6.0	407
9	Phosphorylation of the Cohesin Subunit Scc1 by Polo/Cdc5 Kinase Regulates Sister Chromatid Separation in Yeast. <i>Cell</i> , 2001, 105, 459-472.	13.5	358
10	Identification of <i>cis</i> -acting sites for condensin loading onto budding yeast chromosomes. <i>Genes and Development</i> , 2008, 22, 2215-2227.	2.7	302
11	Biochemical reconstitution of topological DNA binding by the cohesin ring. <i>Nature</i> , 2014, 505, 367-371.	13.7	274
12	Establishment of Sister Chromatid Cohesion at the <i>S. cerevisiae</i> Replication Fork. <i>Molecular Cell</i> , 2006, 23, 787-799.	4.5	268
13	Degradation of a cohesin subunit by the N-end rule pathway is essential for chromosome stability. <i>Nature</i> , 2001, 410, 955-959.	13.7	264
14	Characterization of fission yeast cohesin: essential anaphase proteolysis of Rad21 phosphorylated in the S phase. <i>Genes and Development</i> , 2000, 14, 2757-2770.	2.7	256
15	Cdc14 Phosphatase Induces rDNA Condensation and Resolves Cohesin-Independent Cohesion during Budding Yeast Anaphase. <i>Cell</i> , 2004, 117, 471-482.	13.5	232
16	Downregulation of PP2A ^{Cdc55} Phosphatase by Separase Initiates Mitotic Exit in Budding Yeast. <i>Cell</i> , 2006, 125, 719-732.	13.5	230
17	DNA Entry into and Exit out of the Cohesin Ring by an Interlocking Gate Mechanism. <i>Cell</i> , 2015, 163, 1628-1640.	13.5	217
18	Rapid movement and transcriptional relocalization of human cohesin on DNA. <i>EMBO Journal</i> , 2016, 35, 2671-2685.	3.5	216

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19	A Model for ATP Hydrolysis-Dependent Binding of Cohesin to DNA. <i>Current Biology</i> , 2003, 13, 1930-1940.	1.8	191
20	Stabilization of microtubule dynamics at anaphase onset promotes chromosome segregation. <i>Nature</i> , 2005, 433, 171-176.	13.7	160
21	A non-proteolytic function of separase links the onset of anaphase to mitotic exit. <i>Nature Cell Biology</i> , 2003, 5, 249-254.	4.6	151
22	Orchestrating anaphase and mitotic exit: separase cleavage and localization of Slk19. <i>Nature Cell Biology</i> , 2001, 3, 771-777.	4.6	134
23	Deletion Analysis of the Large Subunit p140 in Human Replication Factor C Reveals Regions Required for Complex Formation and Replication Activities. <i>Journal of Biological Chemistry</i> , 1997, 272, 10058-10064.	1.6	131
24	The Dual Mechanism of Separase Regulation by Securin. <i>Current Biology</i> , 2002, 12, 973-982.	1.8	131
25	Cdk-counteracting phosphatases unlock mitotic exit. <i>Current Opinion in Cell Biology</i> , 2008, 20, 661-668.	2.6	119
26	A mechanism for chromosome segregation sensing by the NoCut checkpoint. <i>Nature Cell Biology</i> , 2009, 11, 477-483.	4.6	118
27	Establishment of DNA-DNA Interactions by the Cohesin Ring. <i>Cell</i> , 2018, 172, 465-477.e15.	13.5	116
28	Division of the Nucleolus and Its Release of CDC14 during Anaphase of Meiosis I Depends on Separase, SPO12, and SLK19. <i>Developmental Cell</i> , 2003, 4, 727-739.	3.1	115
29	Budding Yeast Wapl Controls Sister Chromatid Cohesion Maintenance and Chromosome Condensation. <i>Current Biology</i> , 2013, 23, 64-69.	1.8	114
30	The Scc2-Scc4 complex acts in sister chromatid cohesion and transcriptional regulation by maintaining nucleosome-free regions. <i>Nature Genetics</i> , 2014, 46, 1147-1151.	9.4	114
31	Ctf4 Links DNA Replication with Sister Chromatid Cohesion Establishment by Recruiting the Chl1 Helicase to the Replisome. <i>Molecular Cell</i> , 2016, 63, 371-384.	4.5	113
32	A Quantitative Model for Ordered Cdk Substrate Dephosphorylation during Mitotic Exit. <i>Cell</i> , 2011, 147, 803-814.	13.5	112
33	A Structure-Based Mechanism for DNA Entry into the Cohesin Ring. <i>Molecular Cell</i> , 2020, 79, 917-933.e9.	4.5	112
34	Hos1 Deacetylates Smc3 to Close the Cohesin Acetylation Cycle. <i>Molecular Cell</i> , 2010, 39, 677-688.	4.5	109
35	Chromosome Cohesion and Separation: From Men and Molecules. <i>Current Biology</i> , 2003, 13, R104-R114.	1.8	100
36	Bridging-induced phase separation induced by cohesin SMC protein complexes. <i>Science Advances</i> , 2021, 7, .	4.7	95

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37	In vivo analysis of cohesin architecture using FRET in the budding yeast <i>Saccharomyces cerevisiae</i> . <i>EMBO Journal</i> , 2007, 26, 3783-3793.	3.5	92
38	Irreversibility of mitotic exit is the consequence of systems-level feedback. <i>Nature</i> , 2009, 459, 592-595.	13.7	91
39	PP2A Cdc55 Phosphatase Imposes Ordered Cell-Cycle Phosphorylation by Opposing Threonine Phosphorylation. <i>Molecular Cell</i> , 2017, 65, 393-402.e3.	4.5	91
40	A Complex Consisting of Human Replication Factor C p40, p37, and p36 Subunits Is a DNA-dependent ATPase and an Intermediate in the Assembly of the Holoenzyme. <i>Journal of Biological Chemistry</i> , 1997, 272, 18974-18981.	1.6	89
41	A simple biophysical model emulates budding yeast chromosome condensation. <i>ELife</i> , 2015, 4, e05565.	2.8	87
42	Condensin, Chromatin Crossbarring and Chromosome Condensation. <i>Current Biology</i> , 2012, 22, R1012-R1021.	1.8	83
43	Preferential cleavage of chromatin-bound cohesin after targeted phosphorylation by Polo-like kinase. <i>EMBO Journal</i> , 2004, 23, 3144-3153.	3.5	82
44	The mechanism of sister chromatid cohesion. <i>Experimental Cell Research</i> , 2004, 296, 80-85.	1.2	79
45	Sli15INCENP Dephosphorylation Prevents Mitotic Checkpoint Reengagement Due to Loss of Tension at Anaphase Onset. <i>Current Biology</i> , 2010, 20, 1396-1401.	1.8	77
46	A quantitative model for cyclin-dependent kinase control of the cell cycle: revisited. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2011, 366, 3572-3583.	1.8	77
47	An Eco1-independent sister chromatid cohesion establishment pathway in <i>S. cerevisiae</i> . <i>Chromosoma</i> , 2013, 122, 121-134.	1.0	76
48	Condensin-mediated remodeling of the mitotic chromatin landscape in fission yeast. <i>Nature Genetics</i> , 2017, 49, 1553-1557.	9.4	75
49	Identification of Regions within the Four Small Subunits of Human Replication Factor C Required for Complex Formation and DNA Replication. <i>Journal of Biological Chemistry</i> , 1997, 272, 10065-10071.	1.6	74
50	Secured cutting: controlling separate at the metaphase to anaphase transition. <i>EMBO Reports</i> , 2001, 2, 487-492.	2.0	72
51	Chromosome cohesion and segregation in mitosis and meiosis. <i>Current Opinion in Cell Biology</i> , 2001, 13, 754-761.	2.6	71
52	Conserved features of cohesin binding along fission yeast chromosomes. <i>Genome Biology</i> , 2009, 10, R52.	13.9	71
53	Condensin aids sister chromatid decatenation by topoisomerase II. <i>Nucleic Acids Research</i> , 2014, 42, 340-348.	6.5	68
54	Cloning and characterization of promoter and 5' UTR of the NMDA receptor subunit $\mu 2$: evidence for alternative splicing of 5' non-coding exon. <i>Gene</i> , 1998, 208, 259-269.	1.0	66

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55	Condensin-Dependent rDNA Decatenation Introduces a Temporal Pattern to Chromosome Segregation. <i>Current Biology</i> , 2008, 18, 1084-1089.	1.8	65
56	Cell-cycle regulation of cohesin stability along fission yeast chromosomes. <i>EMBO Journal</i> , 2008, 27, 111-121.	3.5	64
57	A Role for Chromatin Remodeling in Cohesin Loading onto Chromosomes. <i>Molecular Cell</i> , 2019, 74, 664-673.e5.	4.5	62
58	Separase cooperates with Zds1 and Zds2 to activate Cdc14 phosphatase in early anaphase. <i>Journal of Cell Biology</i> , 2008, 182, 873-883.	2.3	61
59	Cohesin loading and sliding. <i>Journal of Cell Science</i> , 2011, 124, 685-691.	1.2	60
60	Structural Studies Reveal the Functional Modularity of the Scc2-Scc4 Cohesin Loader. <i>Cell Reports</i> , 2015, 12, 719-725.	2.9	60
61	Evidence for cohesin sliding along budding yeast chromosomes. <i>Open Biology</i> , 2016, 6, 150178.	1.5	60
62	Identification of <i>chk</i> targets that control cytokinesis. <i>EMBO Journal</i> , 2015, 34, 81-96.	3.5	56
63	A Brownian ratchet model for DNA loop extrusion by the cohesin complex. <i>ELife</i> , 2021, 10, .	2.8	56
64	System-level feedbacks make the anaphase switch irreversible. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 10016-10021.	3.3	55
65	Studies on Substrate Recognition by the Budding Yeast Separase. <i>Journal of Biological Chemistry</i> , 2004, 279, 1191-1196.	1.6	51
66	Structure of the cohesin loader Scc2. <i>Nature Communications</i> , 2017, 8, 13952.	5.8	49
67	Topological in vitro loading of the budding yeast cohesin ring onto DNA. <i>Life Science Alliance</i> , 2018, 1, e201800143.	1.3	49
68	Phosphoproteome dynamics during mitotic exit in budding yeast. <i>EMBO Journal</i> , 2018, 37, .	3.5	47
69	Division of Labor between PCNA Loaders in DNA Replication and Sister Chromatid Cohesion Establishment. <i>Molecular Cell</i> , 2020, 78, 725-738.e4.	4.5	45
70	Cell cycle regulation by feed-forward loops coupling transcription and phosphorylation. <i>Molecular Systems Biology</i> , 2009, 5, 236.	3.2	44
71	A matter of choice: the establishment of sister chromatid cohesion. <i>EMBO Reports</i> , 2009, 10, 1095-1102.	2.0	44
72	Displacement and re-accumulation of centromeric cohesin during transient pre-anaphase centromere splitting. <i>Chromosoma</i> , 2007, 116, 531-544.	1.0	42

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73	SMC complexes orchestrate the mitotic chromatin interaction landscape. <i>Current Genetics</i> , 2018, 64, 335-339.	0.8	41
74	Cdc14 and PP2A Phosphatases Cooperate to Shape Phosphoproteome Dynamics during Mitotic Exit. <i>Cell Reports</i> , 2019, 29, 2105-2119.e4.	2.9	40
75	System-level feedbacks control cell cycle progression. <i>FEBS Letters</i> , 2009, 583, 3992-3998.	1.3	38
76	A PxL motif promotes timely cell cycle substrate dephosphorylation by the Cdc14 phosphatase. <i>Nature Structural and Molecular Biology</i> , 2018, 25, 1093-1102.	3.6	31
77	SMC complexes: Lifting the lid on loop extrusion. <i>Current Opinion in Cell Biology</i> , 2022, 74, 13-22.	2.6	30
78	Facile synthesis of budding yeast α -factor and its use to synchronize cells of α mating type. <i>Yeast</i> , 2012, 29, 233-240.	0.8	27
79	Comparison of loop extrusion and diffusion capture as mitotic chromosome formation pathways in fission yeast. <i>Nucleic Acids Research</i> , 2021, 49, 1294-1312.	6.5	27
80	Separase regulation during mitosis. <i>Biochemical Society Symposia</i> , 2003, 70, 243-251.	2.7	26
81	Passenger acrobatics. <i>Nature</i> , 2003, 426, 780-781.	13.7	25
82	Computational modelling of mitotic exit in budding yeast: the role of separase and Cdc14 endocycles. <i>Journal of the Royal Society Interface</i> , 2011, 8, 1128-1141.	1.5	24
83	Cell-Cycle Regulation of Dynamic Chromosome Association of the Condensin Complex. <i>Cell Reports</i> , 2018, 23, 2308-2317.	2.9	24
84	Mitotic exit in two dimensions. <i>Journal of Theoretical Biology</i> , 2007, 248, 560-573.	0.8	21
85	Structural studies of RFC^{C} tf18 reveal a novel chromatin recruitment role for Dcc1. <i>EMBO Reports</i> , 2017, 18, 558-568.	2.0	19
86	Fission yeast condensin contributes to interphase chromatin organization and prevents transcription-coupled DNA damage. <i>Genome Biology</i> , 2020, 21, 272.	3.8	19
87	Budding yeast relies on G ₁ cyclin specificity to couple cell cycle progression with morphogenetic development. <i>Science Advances</i> , 2021, 7, .	4.7	16
88	Chromosome condensation: Packaging the genome. <i>Current Biology</i> , 2001, 11, R384-R387.	1.8	15
89	Chromosome Segregation. <i>Developmental Cell</i> , 2002, 2, 381-382.	3.1	15
90	Cohesin branches out. <i>Nature</i> , 2008, 451, 777-778.	13.7	15

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91	Mitotic exit in mammalian cells. <i>Molecular Systems Biology</i> , 2009, 5, 324.	3.2	15
92	Cohesion, but not too close. <i>Current Biology</i> , 2001, 11, R378.	1.8	14
93	The "anaphase problem": how to disable the mitotic checkpoint when sisters split. <i>Biochemical Society Transactions</i> , 2010, 38, 1660-1666.	1.6	14
94	Observation of DNA intertwining along authentic budding yeast chromosomes. <i>Genes and Development</i> , 2017, 31, 2151-2161.	2.7	12
95	The Cks1/Cks2 axis fine-tunes Mll1 expression and is crucial for MLL-rearranged leukaemia cell viability. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2018, 1865, 105-116.	1.9	12
96	Conserved roles of chromatin remodellers in cohesin loading onto chromatin. <i>Current Genetics</i> , 2020, 66, 951-956.	0.8	12
97	A global view of substrate phosphorylation and dephosphorylation during budding yeast mitotic exit. <i>Microbial Cell</i> , 2018, 5, 389-392.	1.4	12
98	More than a separase. <i>Nature Cell Biology</i> , 2005, 7, 930-932.	4.6	11
99	Mediator recruits the cohesin loader Scc2 to RNA Pol II-transcribed genes and promotes sister chromatid cohesion. <i>Current Biology</i> , 2022, 32, 2884-2896.e6.	1.8	11
100	Structural Basis of Eco1-Mediated Cohesin Acetylation. <i>Scientific Reports</i> , 2017, 7, 44313.	1.6	10
101	Chromosome cohesion: A polymerase for chromosome bridges. <i>Current Biology</i> , 2000, 10, R698-R700.	1.8	9
102	Chromosome Biology: The Crux of the Ring. <i>Current Biology</i> , 2006, 16, R102-R105.	1.8	8
103	Cohesin subunit Rad21L, the new kid on the block has new ideas. <i>EMBO Reports</i> , 2011, 12, 183-184.	2.0	8
104	Open questions: Chromosome condensation - Why does a chromosome look like a chromosome?. <i>BMC Biology</i> , 2013, 11, 9.	1.7	7
105	Assessing Budding Yeast Phosphoproteome Dynamics in a Time-Resolved Manner using TMT10plex Mass Tag Labeling. <i>STAR Protocols</i> , 2020, 1, 100022.	0.5	7
106	Cell Cycle: The Art of Multi-Tasking. <i>Current Biology</i> , 2010, 20, R101-R103.	1.8	6
107	Nur1 Dephosphorylation Confers Positive Feedback to Mitotic Exit Phosphatase Activation in Budding Yeast. <i>PLoS Genetics</i> , 2015, 11, e1004907.	1.5	6
108	What is your assay for sister-chromatid cohesion?. <i>EMBO Journal</i> , 2007, 26, 4609-4618.	3.5	5

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109	Chromosome segregation: how to open cohesin without cutting the ring?. EMBO Journal, 2013, 32, 614-616.	3.5	4
110	A silent revolution in chromosome biology. Nature Reviews Molecular Cell Biology, 2014, 15, 431-431.	16.1	4
111	Separaseâ€“securin complex: a cunning way to control chromosome segregation. Nature Structural and Molecular Biology, 2017, 24, 337-339.	3.6	4
112	An In Vitro Assay for Monitoring Topological DNA Entrapment by the Chromosomal Cohesin Complex. Methods in Molecular Biology, 2017, 1515, 23-35.	0.4	4
113	Efficient Depletion of Fission Yeast Condensin by Combined Transcriptional Repression and Auxin-Induced Degradation. Methods in Molecular Biology, 2019, 2004, 25-33.	0.4	4
114	Keeping the genome in shape. Nature, 2002, 417, 135-136.	13.7	3
115	A role for condensin in mediating transcriptional adaptation to environmental stimuli. Life Science Alliance, 2021, 4, e202000961.	1.3	3
116	ESCRting DNA at the Cleavage Site During Cytokinesis. Science, 2012, 336, 166-167.	6.0	2
117	Chromosome Condensation: Weaving an Untangled Web. Current Biology, 2015, 25, R663-R666.	1.8	2
118	Building chromosomes without bricks. Science, 2017, 356, 1233-1234.	6.0	2
119	Divide and die another day. Current Opinion in Cell Biology, 2010, 22, 764-765.	2.6	0
120	Analysis of Cell Cycle Progression in the Budding Yeast <i>S. cerevisiae</i> . Methods in Molecular Biology, 2021, 2329, 265-276.	0.4	0
121	Structural characterisation of the <i>Chaetomium thermophilum</i> Ch1 helicase. PLoS ONE, 2021, 16, e0251261.	1.1	0