

Kerry S Bloom

List of Publications by Year in descending order

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186
papers

11,335
citations

22099

59
h-index

34900

98
g-index

218
all docs

218
docs citations

218
times ranked

6655
citing authors

#	ARTICLE	IF	CITATIONS
1	Disruption of mitotic spindle orientation in a yeast dynein mutant.. Proceedings of the National Academy of Sciences of the United States of America, 1993, 90, 10096-10100.	3.3	384
2	Spindle dynamics and cell cycle regulation of dynein in the budding yeast, <i>Saccharomyces cerevisiae</i> .. Journal of Cell Biology, 1995, 130, 687-700.	2.3	369
3	Systematic exploration of essential yeast gene function with temperature-sensitive mutants. Nature Biotechnology, 2011, 29, 361-367.	9.4	352
4	Fractionation of hen oviduct chromatin into transcriptionally active and inactive regions after selective micrococcal nuclease digestion. Cell, 1978, 15, 141-150.	13.5	326
5	Yeast centromere DNA is in a unique and highly ordered structure in chromosomes and small circular minichromosomes. Cell, 1982, 29, 305-317.	13.5	326
6	Astral Microtubule Dynamics in Yeast: A Microtubule-based Searching Mechanism for Spindle Orientation and Nuclear Migration into the Bud. Journal of Cell Biology, 1997, 139, 985-994.	2.3	280
7	Molecular architecture of a kinetochore- microtubule attachment site. Nature Cell Biology, 2006, 8, 581-585.	4.6	263
8	Two different types of double-strand breaks in <i>Saccharomyces cerevisiae</i> are repaired by similar RAD52-independent, nonhomologous recombination events.. Molecular and Cellular Biology, 1994, 14, 1293-1301.	1.1	220
9	The polarity and dynamics of microtubule assembly in the budding yeast <i>Saccharomyces cerevisiae</i> . Nature Cell Biology, 2000, 2, 36-41.	4.6	216
10	Localization and anchoring of mRNA in budding yeast. Current Biology, 1999, 9, 569-S1.	1.8	208
11	Genetic manipulation of centromere function.. Molecular and Cellular Biology, 1987, 7, 2397-2405.	1.1	206
12	Budding Yeast Chromosome Structure and Dynamics during Mitosis. Journal of Cell Biology, 2001, 152, 1255-1266.	2.3	198
13	Centromeres: unique chromatin structures that drive chromosome segregation. Nature Reviews Molecular Cell Biology, 2011, 12, 320-332.	16.1	186
14	Point centromeres contain more than a single centromere-specific Cse4 (CENP-A) nucleosome. Journal of Cell Biology, 2011, 195, 573-582.	2.3	185
15	The role of the proteins Kar9 and Myo2 in orienting the mitotic spindle of budding yeast. Current Biology, 2000, 10, 1497-1506.	1.8	178
16	In Vivo Protein Architecture of the Eukaryotic Kinetochore with Nanometer Scale Accuracy. Current Biology, 2009, 19, 694-699.	1.8	170
17	Chromosome Congression by Kinesin-5 Motor-Mediated Disassembly of Longer Kinetochore Microtubules. Cell, 2008, 135, 894-906.	13.5	168
18	Dynamic Positioning of Mitotic Spindles in Yeast:. Molecular Biology of the Cell, 2000, 11, 3949-3961.	0.9	150

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19	Pericentric Chromatin Is Organized into an Intramolecular Loop in Mitosis. <i>Current Biology</i> , 2008, 18, 81-90.	1.8	148
20	Control of Microtubule Dynamics by Stu2p Is Essential for Spindle Orientation and Metaphase Chromosome Alignment in Yeast. <i>Molecular Biology of the Cell</i> , 2001, 12, 2870-2880.	0.9	146
21	Stable Kinetochore-Microtubule Attachment Constrains Centromere Positioning in Metaphase. <i>Current Biology</i> , 2004, 14, 1962-1967.	1.8	144
22	Molecular architecture of the kinetochore-microtubule attachment site is conserved between point and regional centromeres. <i>Journal of Cell Biology</i> , 2008, 181, 587-594.	2.3	144
23	Chromosome Fragmentation after Induction of a Double-Strand Break Is an Active Process Prevented by the RMX Repair Complex. <i>Current Biology</i> , 2004, 14, 2107-2112.	1.8	140
24	Dynamic Microtubules Lead the Way for Spindle Positioning. <i>Nature Reviews Molecular Cell Biology</i> , 2004, 5, 481-492.	16.1	130
25	Cohesin, condensin, and the intramolecular centromere loop together generate the mitotic chromatin spring. <i>Journal of Cell Biology</i> , 2011, 193, 1167-1180.	2.3	126
26	Tension-dependent Regulation of Microtubule Dynamics at Kinetochores Can Explain Metaphase Congression in Yeast. <i>Molecular Biology of the Cell</i> , 2005, 16, 3764-3775.	0.9	124
27	Phosphorylation of β -Tubulin Regulates Microtubule Organization in Budding Yeast. <i>Developmental Cell</i> , 2001, 1, 621-631.	3.1	115
28	Mitotic Spindle Form and Function. <i>Genetics</i> , 2012, 190, 1197-1224.	1.2	115
29	Control of spindle polarity and orientation in <i>Saccharomyces cerevisiae</i> . <i>Trends in Cell Biology</i> , 2001, 11, 160-166.	3.6	105
30	Mechanisms of force generation by end-on kinetochore-microtubule attachments. <i>Current Opinion in Cell Biology</i> , 2010, 22, 57-67.	2.6	104
31	Chemical Genetics Reveals a Role for Mps1 Kinase in Kinetochore Attachment during Mitosis. <i>Current Biology</i> , 2005, 15, 160-165.	1.8	100
32	Integrating high-throughput genetic interaction mapping and high-content screening to explore yeast spindle morphogenesis. <i>Journal of Cell Biology</i> , 2010, 188, 69-81.	2.3	100
33	Chromatin conformation of yeast centromeres.. <i>Journal of Cell Biology</i> , 1984, 99, 1559-1568.	2.3	97
34	Mechanisms of Microtubule-Based Kinetochore Positioning in the Yeast Metaphase Spindle. <i>Biophysical Journal</i> , 2003, 84, 3529-3546.	0.2	93
35	Mps1 Phosphorylation of Dam1 Couples Kinetochores to Microtubule Plus Ends at Metaphase. <i>Current Biology</i> , 2006, 16, 1489-1501.	1.8	93
36	Chromatin structure of altered yeast centromeres.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1988, 85, 175-179.	3.3	89

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37	The Differential Roles of Budding Yeast Tem1p, Cdc15p, and Bub2p Protein Dynamics in Mitotic Exit. <i>Molecular Biology of the Cell</i> , 2004, 15, 1519-1532.	0.9	89
38	Centromere Tethering Confines Chromosome Domains. <i>Molecular Cell</i> , 2013, 52, 819-831.	4.5	88
39	The Role of Actin in Spindle Orientation Changes during the <i>Saccharomyces cerevisiae</i> Cell Cycle. <i>Journal of Cell Biology</i> , 1999, 146, 1019-1032.	2.3	86
40	Rho GTPase regulation of exocytosis in yeast is independent of GTP hydrolysis and polarization of the exocyst complex. <i>Journal of Cell Biology</i> , 2005, 170, 583-594.	2.3	86
41	Microtubule Dynamics from Mating through the First Zygotic Division in the Budding Yeast <i>Saccharomyces cerevisiae</i> . <i>Journal of Cell Biology</i> , 1999, 144, 977-987.	2.3	85
42	A dynamin-like protein encoded by the yeast sporulation gene SP015. <i>Nature</i> , 1991, 349, 713-715.	13.7	84
43	Thin-foil magnetic force system for high-numerical-aperture microscopy. <i>Review of Scientific Instruments</i> , 2006, 77, 023702.	0.6	84
44	Identification of a Mid-anaphase Checkpoint in Budding Yeast. <i>Journal of Cell Biology</i> , 1997, 136, 345-354.	2.3	79
45	Imaging green fluorescent protein fusion proteins in <i>Saccharomyces cerevisiae</i> . <i>Current Biology</i> , 1997, 7, 701-704.	1.8	78
46	Pericentric Chromatin Is an Elastic Component of the Mitotic Spindle. <i>Current Biology</i> , 2007, 17, 741-748.	1.8	78
47	Analysis of the complexity and frequency of zein genes in the maize genome. <i>Biochemistry</i> , 1980, 19, 1644-1650.	1.2	76
48	The centromere frontier: Kinetochore components, microtubule-based motility, and the CEN-value paradox. <i>Cell</i> , 1993, 73, 621-624.	13.5	76
49	Yeast Kinetochores Do Not Stabilize Stu2p-dependent Spindle Microtubule Dynamics. <i>Molecular Biology of the Cell</i> , 2003, 14, 4181-4195.	0.9	75
50	Acquisition and processing of a conditional dicentric chromosome in <i>Saccharomyces cerevisiae</i> . <i>Molecular and Cellular Biology</i> , 1989, 9, 1368-1370.	1.1	73
51	SUMO-targeted ubiquitin ligase (STUbL) Slx5 regulates proteolysis of centromeric histone H3 variant Cse4 and prevents its mislocalization to euchromatin. <i>Molecular Biology of the Cell</i> , 2016, 27, 1500-1510.	0.9	73
52	A phosphatidylinositol transfer protein integrates phosphoinositide signaling with lipid droplet metabolism to regulate a developmental program of nutrient stress-induced membrane biogenesis. <i>Molecular Biology of the Cell</i> , 2014, 25, 712-727.	0.9	71
53	How the kinetochore couples microtubule force and centromere stretch to move chromosomes. <i>Nature Cell Biology</i> , 2016, 18, 382-392.	4.6	70
54	The Minus End-Directed Motor Kar3 Is Required for Coupling Dynamic Microtubule Plus Ends to the Cortical Shmoo Tip in Budding Yeast. <i>Current Biology</i> , 2003, 13, 1423-1428.	1.8	69

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55	Microtubule dynamics drive enhanced chromatin motion and mobilize telomeres in response to DNA damage. <i>Molecular Biology of the Cell</i> , 2017, 28, 1701-1711.	0.9	69
56	Structural Analysis and Sequence Organization of Yeast Centromeres. <i>Cold Spring Harbor Symposia on Quantitative Biology</i> , 1983, 47, 1175-1185.	2.0	69
57	FBW7 Loss Promotes Chromosomal Instability and Tumorigenesis via Cyclin E1/CDK2-Mediated Phosphorylation of CENP-A. <i>Cancer Research</i> , 2017, 77, 4881-4893.	0.4	68
58	Nucleosome depletion alters the chromatin structure of <i>Saccharomyces cerevisiae</i> centromeres.. <i>Molecular and Cellular Biology</i> , 1990, 10, 5721-5727.	1.1	67
59	β -Tubulin C354 Mutations that Severely Decrease Microtubule Dynamics Do Not Prevent Nuclear Migration in Yeast. <i>Molecular Biology of the Cell</i> , 2002, 13, 2919-2932.	0.9	65
60	The microtubule-based motor Kar3 and plus end-binding protein Bim1 provide structural support for the anaphase spindle. <i>Journal of Cell Biology</i> , 2008, 180, 91-100.	2.3	64
61	Centromeric Heterochromatin: The Primordial Segregation Machine. <i>Annual Review of Genetics</i> , 2014, 48, 457-484.	3.2	64
62	Enrichment of dynamic chromosomal crosslinks drive phase separation of the nucleolus. <i>Nucleic Acids Research</i> , 2017, 45, 11159-11173.	6.5	64
63	Coordinated Spindle Assembly and Orientation Requires Clb5p-Dependent Kinase in Budding Yeast. <i>Journal of Cell Biology</i> , 2000, 148, 441-452.	2.3	61
64	Towards building a chromosome segregation machine. <i>Nature</i> , 2010, 463, 446-456.	13.7	61
65	Differential kinetochore protein requirements for establishment versus propagation of centromere activity in <i>Saccharomyces cerevisiae</i> . <i>Journal of Cell Biology</i> , 2003, 160, 833-843.	2.3	58
66	Beyond the code: the mechanical properties of DNA as they relate to mitosis. <i>Chromosoma</i> , 2008, 117, 103-110.	1.0	58
67	Bud6 Directs Sequential Microtubule Interactions with the Bud Tip and Bud Neck during Spindle Morphogenesis in <i>Saccharomyces cerevisiae</i> . <i>Molecular Biology of the Cell</i> , 2000, 11, 3689-3702.	0.9	57
68	Nuclear migration: Cortical anchors for cytoplasmic dynein. <i>Current Biology</i> , 2001, 11, R326-R329.	1.8	57
69	Dicentric Chromosome Stretching during Anaphase Reveals Roles of Sir2/Ku in Chromatin Compaction in Budding Yeast. <i>Molecular Biology of the Cell</i> , 2001, 12, 2800-2812.	0.9	57
70	Function and Assembly of DNA Looping, Clustering, and Microtubule Attachment Complexes within a Eukaryotic Kinetochore. <i>Molecular Biology of the Cell</i> , 2009, 20, 4131-4139.	0.9	56
71	Pericentric chromatin loops function as a nonlinear spring in mitotic force balance. <i>Journal of Cell Biology</i> , 2013, 200, 757-772.	2.3	56
72	Kinesin-8 molecular motors: putting the brakes on chromosome oscillations. <i>Trends in Cell Biology</i> , 2008, 18, 307-310.	3.6	55

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73	A 3D Map of the Yeast Kinetochores Reveals the Presence of Core and Accessory Centromere-Specific Histone. <i>Current Biology</i> , 2013, 23, 1939-1944.	1.8	55
74	Centromeres: An Integrated Protein/DNA Complex Required for Chromosome Movement. <i>Annual Review of Cell Biology</i> , 1991, 7, 311-336.	26.0	54
75	Counting Kinetochores Protein Numbers in Budding Yeast Using Genetically Encoded Fluorescent Proteins. <i>Methods in Cell Biology</i> , 2008, 85, 127-151.	0.5	54
76	Itâ€™s a kar9ochore to capture microtubules. <i>Nature Cell Biology</i> , 2000, 2, E96-E98.	4.6	53
77	The kinetochores protein Ndc10p is required for spindle stability and cytokinesis in yeast. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 5408-5413.	3.3	50
78	Design Features of a Mitotic Spindle: Balancing Tension and Compression at a Single Microtubule Kinetochores Interface in Budding Yeast. <i>Annual Review of Genetics</i> , 2008, 42, 335-359.	3.2	49
79	Determining absolute protein numbers by quantitative fluorescence microscopy. <i>Methods in Cell Biology</i> , 2014, 123, 347-365.	0.5	49
80	Microtubule motors in eukaryotic spindle assembly and maintenance. <i>Seminars in Cell and Developmental Biology</i> , 2010, 21, 248-254.	2.3	46
81	Bub1 Kinase and Sgo1 Modulate Pericentric Chromatin in Response to Altered Microtubule Dynamics. <i>Current Biology</i> , 2012, 22, 471-481.	1.8	46
82	DNA loops generate intracentromere tension in mitosis. <i>Journal of Cell Biology</i> , 2015, 210, 553-564.	2.3	46
83	Nuclear congression is driven by cytoplasmic microtubule plus end interactions in <i>S. cerevisiae</i> . <i>Journal of Cell Biology</i> , 2006, 172, 27-39.	2.3	45
84	The <i>Saccharomyces cerevisiae</i> Spindle Pole Body Is a Dynamic Structure. <i>Molecular Biology of the Cell</i> , 2003, 14, 3494-3505.	0.9	43
85	ASH1 mRNA Localization in Three Acts. <i>Molecular Biology of the Cell</i> , 2001, 12, 2567-2577.	0.9	42
86	DNA relaxation dynamics as a probe for the intracellular environment. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 9250-9255.	3.3	42
87	Chapter 10 A High-Resolution Multimode Digital Microscope System. <i>Methods in Cell Biology</i> , 1998, 56, 185-215.	0.5	41
88	Measuring Nanometer Scale Gradients in Spindle Microtubule Dynamics Using Model Convolution Microscopy. <i>Molecular Biology of the Cell</i> , 2006, 17, 4069-4079.	0.9	40
89	Entropy gives rise to topologically associating domains. <i>Nucleic Acids Research</i> , 2016, 44, 5540-5549.	6.5	40
90	Centromere Structure and Function. <i>Progress in Molecular and Subcellular Biology</i> , 2017, 56, 515-539.	0.9	40

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91	Tightly centromere-linked gene (SPO15) essential for meiosis in the yeast <i>Saccharomyces cerevisiae</i> . <i>Molecular and Cellular Biology</i> , 1986, 6, 158-167.	1.1	39
92	ChromoShake: a chromosome dynamics simulator reveals that chromatin loops stiffen centromeric chromatin. <i>Molecular Biology of the Cell</i> , 2016, 27, 153-166.	0.9	39
93	Nuclear and Spindle Dynamics in Budding Yeast. <i>Molecular Biology of the Cell</i> , 1998, 9, 1627-1631.	0.9	38
94	The yeast DNA damage checkpoint proteins control a cytoplasmic response to DNA damage. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 11358-11363.	3.3	38
95	Esperanto for histones: CENP-A, not CenH3, is the centromeric histone H3 variant. <i>Chromosome Research</i> , 2013, 21, 101-106.	1.0	37
96	Ending the Rules: Widefield Microscopy and the Abbe Limit of Resolution. <i>Journal of Cellular Physiology</i> , 2014, 229, 132-138.	2.0	37
97	Systematic Triple-Mutant Analysis Uncovers Functional Connectivity between Pathways Involved in Chromosome Regulation. <i>Cell Reports</i> , 2013, 3, 2168-2178.	2.9	36
98	Kar9p-independent Microtubule Capture at Bud6p Cortical Sites Primes Spindle Polarity before Bud Emergence in <i>Saccharomyces cerevisiae</i> . <i>Molecular Biology of the Cell</i> , 2002, 13, 4141-4155.	0.9	35
99	The path of DNA in the kinetochore. <i>Current Biology</i> , 2006, 16, R276-R278.	1.8	35
100	Individual pericentromeres display coordinated motion and stretching in the yeast spindle. <i>Journal of Cell Biology</i> , 2013, 203, 407-416.	2.3	35
101	Tension-dependent nucleosome remodeling at the pericentromere in yeast. <i>Molecular Biology of the Cell</i> , 2012, 23, 2560-2570.	0.9	33
102	Cell structure and dynamics. <i>Current Opinion in Cell Biology</i> , 2007, 19, 1-4.	2.6	32
103	Model Convolution: A Computational Approach to Digital Image Interpretation. <i>Cellular and Molecular Bioengineering</i> , 2010, 3, 163-170.	1.0	32
104	The spatial segregation of pericentric cohesin and condensin in the mitotic spindle. <i>Molecular Biology of the Cell</i> , 2013, 24, 3909-3919.	0.9	32
105	The structure of a primitive kinetochore. <i>Trends in Biochemical Sciences</i> , 1989, 14, 223-227.	3.7	31
106	Dyskerin, tRNA genes, and condensin tether pericentric chromatin to the spindle axis in mitosis. <i>Journal of Cell Biology</i> , 2014, 207, 189-199.	2.3	31
107	A Kinesin-5, Cin8, Recruits Protein Phosphatase 1 to Kinetochores and Regulates Chromosome Segregation. <i>Current Biology</i> , 2018, 28, 2697-2704.e3.	1.8	30
108	Nuclear oscillations and nuclear filament formation accompany single-strand annealing repair of a dicentric chromosome in <i>Saccharomyces cerevisiae</i> . <i>Journal of Cell Science</i> , 2003, 116, 561-569.	1.2	29

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109	Polymer models of interphase chromosomes. <i>Nucleus</i> , 2014, 5, 376-390.	0.6	29
110	Microtubule dynamics in the budding yeast mating pathway. <i>Journal of Cell Science</i> , 2006, 119, 3485-3490.	1.2	28
111	A High-Resolution Multimode Digital Microscope System. <i>Methods in Cell Biology</i> , 2007, 81, 187-218.	0.5	28
112	Spatial signals link exit from mitosis to spindle position. <i>ELife</i> , 2016, 5, .	2.8	26
113	Chromosome integrity at a double-strand break requires exonuclease 1 and MRX. <i>DNA Repair</i> , 2011, 10, 102-110.	1.3	25
114	R-loops at centromeric chromatin contribute to defects in kinetochore integrity and chromosomal instability in budding yeast. <i>Molecular Biology of the Cell</i> , 2021, 32, 74-89.	0.9	25
115	Centromere dynamics. <i>Current Opinion in Genetics and Development</i> , 2007, 17, 151-156.	1.5	24
116	Inferring Latent States and Refining Force Estimates via Hierarchical Dirichlet Process Modeling in Single Particle Tracking Experiments. <i>PLoS ONE</i> , 2015, 10, e0137633.	1.1	24
117	Polo kinase Cdc5 associates with centromeres to facilitate the removal of centromeric cohesin during mitosis. <i>Molecular Biology of the Cell</i> , 2016, 27, 2286-2300.	0.9	24
118	Tension Management in the Kinetochore. <i>Current Biology</i> , 2010, 20, R1040-R1048.	1.8	23
119	Pat1 protects centromere-specific histone H3 variant Cse4 from Psh1-mediated ubiquitination. <i>Molecular Biology of the Cell</i> , 2015, 26, 2067-2079.	0.9	23
120	tRNA Genes Affect Chromosome Structure and Function via Local Effects. <i>Molecular and Cellular Biology</i> , 2019, 39, .	1.1	23
121	Tension sensors reveal how the kinetochore shares its load. <i>BioEssays</i> , 2017, 39, 1600216.	1.2	22
122	RotoStep: A Chromosome Dynamics Simulator Reveals Mechanisms of Loop Extrusion. <i>Cold Spring Harbor Symposia on Quantitative Biology</i> , 2017, 82, 101-109.	2.0	22
123	Common Features of the Pericentromere and Nucleolus. <i>Genes</i> , 2019, 10, 1029.	1.0	20
124	Hypothesis testing via integrated computer modeling and digital fluorescence microscopy. <i>Methods</i> , 2007, 41, 232-237.	1.9	19
125	The regulation of chromosome segregation via centromere loops. <i>Critical Reviews in Biochemistry and Molecular Biology</i> , 2019, 54, 352-370.	2.3	19
126	The rDNA is biomolecular condensate formed by polymer-polymer phase separation and is sequestered in the nucleolus by transcription and R-loops. <i>Nucleic Acids Research</i> , 2021, 49, 4586-4598.	6.5	19

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127	FluoroSim: A Visual Problem-Solving Environment for Fluorescence Microscopy. Eurographics Workshop on Visual Computing for Biomedicine, 2008, 2008, 151-158.	4.0	19
128	Chapter 19 Using Green Fluorescent Protein Fusion Proteins to Quantitate Microtubule and Spindle Dynamics in Budding Yeast. Methods in Cell Biology, 1998, 61, 369-383.	0.5	18
129	Persistent mechanical linkage between sister chromatids throughout anaphase. Chromosoma, 2009, 118, 633-645.	1.0	18
130	Genetic dissection of centromere function.. Molecular and Cellular Biology, 1993, 13, 3156-3166.	1.1	16
131	The role of centromere-binding factor 3 (CBF3) in spindle stability, cytokinesis, and kinetochore attachment. Biochemistry and Cell Biology, 2005, 83, 696-702.	0.9	16
132	Nonrandom Distribution of Interhomolog Recombination Events Induced by Breakage of a Dicentric Chromosome in <i>Saccharomyces cerevisiae</i> . Genetics, 2013, 194, 69-80.	1.2	16
133	Chromatin structures of <i>Kluyveromyces lactis</i> centromeres in <i>K. lactis</i> and <i>Saccharomyces cerevisiae</i> . Chromosoma, 1993, 102, 660-667.	1.0	15
134	Geometric partitioning of cohesin and condensin is a consequence of chromatin loops. Molecular Biology of the Cell, 2018, 29, 2737-2750.	0.9	15
135	Cdk1 phosphorylation of Esp1/Separase functions with PP2A and Slk19 to regulate pericentric Cohesin and anaphase onset. PLoS Genetics, 2018, 14, e1007029.	1.5	15
136	Selective excision of the centromere chromatin complex from <i>Saccharomyces cerevisiae</i> .. Journal of Cell Biology, 1988, 107, 9-15.	2.3	14
137	Statistical mechanics of chromosomes: <i>in vivo</i> and <i>in silico</i> approaches reveal high-level organization and structure arise exclusively through mechanical feedback between loop extruders and chromatin substrate properties. Nucleic Acids Research, 2020, 48, 11284-11303.	6.5	14
138	The Locomotor Activity of Fish: An Analogy to the Kinetics of an Opposed First-Order Chemical Reaction. Transactions of the American Fisheries Society, 1975, 104, 752-754.	0.6	13
139	Heterogeneity and maintenance of centromere plasmid copy number in <i>Saccharomyces cerevisiae</i> . Chromosoma, 1990, 99, 281-288.	1.0	13
140	mRNA localization: motile RNA, asymmetric anchors. Current Opinion in Microbiology, 1999, 2, 604-609.	2.3	13
141	Yeast weighs in on the elusive spindle matrix: New filaments in the nucleus. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 4757-4759.	3.3	13
142	Lessons learned from counting molecules: how to lure CENP-A into the kinetochore. Open Biology, 2014, 4, 140191.	1.5	13
143	Shaping centromeres to resist mitotic spindle forces. Journal of Cell Science, 2022, 135, .	1.2	12
144	Microtubule composition: Cryptography of dynamic polymers. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 6839-6840.	3.3	11

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145	Transient crosslinking kinetics optimize gene cluster interactions. <i>PLoS Computational Biology</i> , 2019, 15, e1007124.	1.5	10
146	Fork pausing allows centromere DNA loop formation and kinetochore assembly. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 11784-11789.	3.3	9
147	Behavior of dicentric chromosomes in budding yeast. <i>PLoS Genetics</i> , 2021, 17, e1009442.	1.5	9
148	[26] High-resolution video and digital-enhanced differential interference contrast light microscopy of cell division in budding yeast. <i>Methods in Enzymology</i> , 1998, 298, 317-331.	0.4	8
149	The SUMO deconjugating peptidase Smt4 contributes to the mechanism required for transition from sister chromatid arm cohesion to sister chromatid pericentromere separation. <i>Cell Cycle</i> , 2015, 14, 2206-2218.	1.3	8
150	Cdc7-mediated phosphorylation of Cse4 regulates high-fidelity chromosome segregation in budding yeast. <i>Molecular Biology of the Cell</i> , 2021, 32, ar15.	0.9	8
151	Centromeres and telomeres: structural element's of eukaryotic chromosomes. <i>Current Opinion in Cell Biology</i> , 1989, 1, 526-532.	2.6	7
152	Chromosome Segregation: Seeing Is Believing. <i>Current Biology</i> , 2005, 15, R500-R503.	1.8	7
153	Performance of deep learning restoration methods for the extraction of particle dynamics in noisy microscopy image sequences. <i>Molecular Biology of the Cell</i> , 2021, 32, 903-914.	0.9	7
154	A Cohesin-Based Partitioning Mechanism Revealed upon Transcriptional Inactivation of Centromere. <i>PLoS Genetics</i> , 2016, 12, e1006021.	1.5	7
155	The nucleosome repeat length of <i>Kluyveromyces lactis</i> is 16 bp longer than that of <i>Saccharomyces cerevisiae</i> . <i>Nucleic Acids Research</i> , 1993, 21, 2247-2248.	6.5	6
156	A High-Resolution Multimode Digital Microscope System. <i>Methods in Cell Biology</i> , 2013, 114, 179-210.	0.5	6
157	AI-Assisted Forward Modeling of Biological Structures. <i>Frontiers in Cell and Developmental Biology</i> , 2019, 7, 279.	1.8	6
158	DNA damage reduces heterogeneity and coherence of chromatin motions. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	3.3	6
159	Hitching a ride. <i>EMBO Reports</i> , 2006, 7, 985-987.	2.0	5
160	Stu2 uses a 15-nm parallel coiled coil for kinetochore localization and concomitant regulation of the mitotic spindle. <i>Molecular Biology of the Cell</i> , 2018, 29, 285-294.	0.9	5
161	UV-induced damage and repair in centromere DNA of yeast. <i>Molecular Genetics and Genomics</i> , 1987, 210, 16-22.	2.4	4
162	NoCut: Cytokinesis in Check. <i>Cell</i> , 2006, 125, 17-18.	13.5	4

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