

Anthony A Hyman

List of Publications by Year in descending order

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Version: 2024-02-01

225
papers

50,751
citations

1094

112
h-index

1928

207
g-index

240
all docs

240
docs citations

240
times ranked

36977
citing authors

#	ARTICLE	IF	CITATIONS
1	Sequence-dependent surface condensation of a pioneer transcription factor on DNA. <i>Nature Physics</i> , 2022, 18, 271-276.	6.5	73
2	Characterization of RNA content in individual phase-separated coacervate microdroplets. <i>Nature Communications</i> , 2022, 13, 2626.	5.8	14
3	Phosphofructokinase relocalizes into subcellular compartments with liquid-like properties in vivo. <i>Biophysical Journal</i> , 2021, 120, 1170-1186.	0.2	39
4	Reentrant liquid condensate phase of proteins is stabilized by hydrophobic and non-ionic interactions. <i>Nature Communications</i> , 2021, 12, 1085.	5.8	245
5	Feedback control of PLK1 by Apo1 ensures accurate chromosome segregation. <i>Cell Reports</i> , 2021, 36, 109343.	2.9	15
6	HspB8 prevents aberrant phase transitions of FUS by chaperoning its folded RNA-binding domain. <i>ELife</i> , 2021, 10, .	2.8	42
7	Local thermodynamics govern formation and dissolution of <i>Caenorhabditis elegans</i> P granule condensates. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	64
8	Biomolecular condensates at the nexus of cellular stress, protein aggregation disease and ageing. <i>Nature Reviews Molecular Cell Biology</i> , 2021, 22, 196-213.	16.1	535
9	ESI mutagenesis: a one-step method for introducing mutations into bacterial artificial chromosomes. <i>Life Science Alliance</i> , 2021, 4, e202000836.	1.3	2
10	Quantitative theory for the diffusive dynamics of liquid condensates. <i>ELife</i> , 2021, 10, .	2.8	22
11	ASCB Keith Porter Lecture. <i>Molecular Biology of the Cell</i> , 2020, 31, 2864-2867.	0.9	1
12	Protein condensates as aging Maxwell fluids. <i>Science</i> , 2020, 370, 1317-1323.	6.0	247
13	Partitioning of cancer therapeutics in nuclear condensates. <i>Science</i> , 2020, 368, 1386-1392.	6.0	281
14	Condensation of Ded1p Promotes a Translational Switch from Housekeeping to Stress Protein Production. <i>Cell</i> , 2020, 181, 818-831.e19.	13.5	130
15	RNA-Induced Conformational Switching and Clustering of G3BP Drive Stress Granule Assembly by Condensation. <i>Cell</i> , 2020, 181, 346-361.e17.	13.5	557
16	Drops and fibers – how biomolecular condensates and cytoskeletal filaments influence each other. <i>Emerging Topics in Life Sciences</i> , 2020, 4, 247-261.	1.1	54
17	Soluble tubulin is significantly enriched at mitotic centrosomes. <i>Journal of Cell Biology</i> , 2019, 218, 3977-3985.	2.3	26
18	Kinetically distinct phases of tau on microtubules regulate kinesin motors and severing enzymes. <i>Nature Cell Biology</i> , 2019, 21, 1086-1092.	4.6	113

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19	FUS pathology in ALS is linked to alterations in multiple ALS-associated proteins and rescued by drugs stimulating autophagy. <i>Acta Neuropathologica</i> , 2019, 138, 67-84.	3.9	94
20	Directed Growth of Biomimetic Microcompartments. <i>Advanced Biology</i> , 2019, 3, e1800314.	3.0	25
21	Inhibition of <sc>CPAP</sc> â€“tubulin interaction prevents proliferation of centrosomeâ€“amplified cancer cells. <i>EMBO Journal</i> , 2019, 38, .	3.5	24
22	Phosphatase PP2A and microtubule-mediated pulling forces disassemble centrosomes during mitotic exit. <i>Biology Open</i> , 2018, 7, .	0.6	32
23	Controlling compartmentalization by non-membrane-bound organelles. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2018, 373, 20170193.	1.8	132
24	RNA buffers the phase separation behavior of prion-like RNA binding proteins. <i>Science</i> , 2018, 360, 918-921.	6.0	837
25	Isogenic FUS-eGFP iPSC Reporter Lines Enable Quantification of FUS Stress Granule Pathology that Is Rescued by Drugs Inducing Autophagy. <i>Stem Cell Reports</i> , 2018, 10, 375-389.	2.3	95
26	Impaired DNA damage response signaling by FUS-NLS mutations leads to neurodegeneration and FUS aggregate formation. <i>Nature Communications</i> , 2018, 9, 335.	5.8	217
27	Phase separation of a yeast prion protein promotes cellular fitness. <i>Science</i> , 2018, 359, .	6.0	534
28	Protein Dynamics in Complex DNA Lesions. <i>Molecular Cell</i> , 2018, 69, 1046-1061.e5.	4.5	128
29	Organization and Function of Non-dynamic Biomolecular Condensates. <i>Trends in Biochemical Sciences</i> , 2018, 43, 81-94.	3.7	160
30	Salt-Dependent Rheology and Surface Tension of Protein Condensates Using Optical Traps. <i>Physical Review Letters</i> , 2018, 121, 258101.	2.9	125
31	Positioning of Particles in Active Droplets. <i>Physical Review Letters</i> , 2018, 121, 158102.	2.9	24
32	Phase Transitions Drive the Formation of Vesicular Stomatitis Virus Replication Compartments. <i>MBio</i> , 2018, 9, .	1.8	183
33	A Userâ€™s Guide for Phase Separation Assays with Purified Proteins. <i>Journal of Molecular Biology</i> , 2018, 430, 4806-4820.	2.0	195
34	A Molecular Grammar Governing the Driving Forces for Phase Separation of Prion-like RNA Binding Proteins. <i>Cell</i> , 2018, 174, 688-699.e16.	13.5	1,372
35	Rab5 and Alsin regulate stress-activated cytoprotective signaling on mitochondria. <i>ELife</i> , 2018, 7, .	2.8	65
36	Different Material States of Pub1 Condensates Define Distinct Modes of Stress Adaptation and Recovery. <i>Cell Reports</i> , 2018, 23, 3327-3339.	2.9	183

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37	The replicative helicase MCM recruits cohesin acetyltransferase ESCO2 to mediate centromeric sister chromatid cohesion. <i>EMBO Journal</i> , 2018, 37, .	3.5	50
38	Biomolecular condensates: organizers of cellular biochemistry. <i>Nature Reviews Molecular Cell Biology</i> , 2017, 18, 285-298.	16.1	3,771
39	ATP as a biological hydrotrope. <i>Science</i> , 2017, 356, 753-756.	6.0	677
40	A liquid reservoir for silent chromatin. <i>Nature</i> , 2017, 547, 168-169.	13.7	20
41	Stem cells: the new "model organism". <i>Molecular Biology of the Cell</i> , 2017, 28, 1409-1411.	0.9	22
42	The Centrosome Is a Selective Condensate that Nucleates Microtubules by Concentrating Tubulin. <i>Cell</i> , 2017, 169, 1066-1077.e10.	13.5	533
43	An aberrant phase transition of stress granules triggered by misfolded protein and prevented by chaperone function. <i>EMBO Journal</i> , 2017, 36, 1669-1687.	3.5	370
44	Growth and division of active droplets provides a model for protocells. <i>Nature Physics</i> , 2017, 13, 408-413.	6.5	304
45	Genome-scale single-cell mechanical phenotyping reveals disease-related genes involved in mitotic rounding. <i>Nature Communications</i> , 2017, 8, 1266.	5.8	52
46	Local Nucleation of Microtubule Bundles through Tubulin Concentration into a Condensed Tau Phase. <i>Cell Reports</i> , 2017, 20, 2304-2312.	2.9	278
47	RNA gets in phase. <i>Journal of Cell Biology</i> , 2017, 216, 2235-2237.	2.3	23
48	<scp>CPAP</scp> promotes timely cilium disassembly to maintain neural progenitor pool. <i>EMBO Journal</i> , 2016, 35, 803-819.	3.5	208
49	TransgeneOmics " A transgenic platform for protein localization based function exploration. <i>Methods</i> , 2016, 96, 69-74.	1.9	13
50	Polo-like kinase phosphorylation determines <i>Caenorhabditis elegans</i> centrosome size and density by biasing SPD-5 toward an assembly-competent conformation. <i>Biology Open</i> , 2016, 5, 1431-1440.	0.6	53
51	In vitro Reconstitution of a Membrane Switch Mechanism for the Polarity Protein LGL. <i>Journal of Molecular Biology</i> , 2016, 428, 4828-4842.	2.0	15
52	Rheology of the Active Cell Cortex in Mitosis. <i>Biophysical Journal</i> , 2016, 111, 589-600.	0.2	119
53	Amyloid-like Self-Assembly of a Cellular Compartment. <i>Cell</i> , 2016, 166, 637-650.	13.5	294
54	Polar Positioning of Phase-Separated Liquid Compartments in Cells Regulated by an mRNA Competition Mechanism. <i>Cell</i> , 2016, 166, 1572-1584.e16.	13.5	283

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55	Are aberrant phase transitions a driver of cellular aging?. <i>BioEssays</i> , 2016, 38, 959-968.	1.2	234
56	A locus in <i>Pristionchus pacificus</i> that is responsible for the ability to give rise to fertile offspring at higher temperatures. <i>Biology Open</i> , 2016, 5, 1111-1117.	0.6	6
57	The Mitotic Spindle in the One-Cell <i>C. elegans</i> Embryo Is Positioned with High Precision and Stability. <i>Biophysical Journal</i> , 2016, 111, 1773-1784.	0.2	27
58	Molecular basis for CPAP-tubulin interaction in controlling centriolar and ciliary length. <i>Nature Communications</i> , 2016, 7, 11874.	5.8	66
59	PLEKHA7 Recruits PDZD11 to Adherens Junctions to Stabilize Nectins. <i>Journal of Biological Chemistry</i> , 2016, 291, 11016-11029.	1.6	28
60	Ki-67 acts as a biological surfactant to disperse mitotic chromosomes. <i>Nature</i> , 2016, 535, 308-312.	13.7	392
61	Site-Specific Cryo-focused Ion Beam Sample Preparation Guided by 3D Correlative Microscopy. <i>Biophysical Journal</i> , 2016, 110, 860-869.	0.2	172
62	Visualizing the molecular sociology at the HeLa cell nuclear periphery. <i>Science</i> , 2016, 351, 969-972.	6.0	493
63	Priority of discovery in the life sciences. <i>ELife</i> , 2016, 5, .	2.8	28
64	Suppression of Ostwald ripening in active emulsions. <i>Physical Review E</i> , 2015, 92, 012317.	0.8	146
65	Mitotic cells contract actomyosin cortex and generate pressure to round against or escape epithelial confinement. <i>Nature Communications</i> , 2015, 6, 8872.	5.8	79
66	Temperature Dependence of Cell Division Timing Accounts for a Shift in the Thermal Limits of <i>C. elegans</i> and <i>C. briggsae</i> . <i>Cell Reports</i> , 2015, 10, 647-653.	2.9	85
67	Cdk1-dependent mitotic enrichment of cortical myosin II promotes cell rounding against confinement. <i>Nature Cell Biology</i> , 2015, 17, 148-159.	4.6	131
68	Emergent Properties of the Metaphase Spindle. <i>Cold Spring Harbor Perspectives in Biology</i> , 2015, 7, a015784.	2.3	40
69	Method. <i>Methods in Cell Biology</i> , 2015, 129, 369-382.	0.5	18
70	Sestrin 2 Protein Regulates Platelet-derived Growth Factor Receptor β^2 ($Pdgfr\beta^2$) Expression by Modulating Proteasomal and Nrf2 Transcription Factor Functions. <i>Journal of Biological Chemistry</i> , 2015, 290, 9738-9752.	1.6	17
71	Regulated assembly of a supramolecular centrosome scaffold in vitro. <i>Science</i> , 2015, 348, 808-812.	6.0	170
72	A Human Interactome in Three Quantitative Dimensions Organized by Stoichiometries and Abundances. <i>Cell</i> , 2015, 163, 712-723.	13.5	1,132

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73	A focused ion beam milling and lift-out approach for site-specific preparation of frozen-hydrated lamellas from multicellular organisms. <i>Journal of Structural Biology</i> , 2015, 192, 262-269.	1.3	125
74	A Liquid-to-Solid Phase Transition of the ALS Protein FUS Accelerated by Disease Mutation. <i>Cell</i> , 2015, 162, 1066-1077.	13.5	2,182
75	Quantitative comparison of a human cancer cell surface proteome between interphase and mitosis. <i>EMBO Journal</i> , 2015, 34, 251-265.	3.5	41
76	Coiled-Coil Proteins Facilitated the Functional Expansion of the Centrosome. <i>PLoS Computational Biology</i> , 2014, 10, e1003657.	1.5	32
77	Products of the Parkinson's disease-related glyoxalase DJ-1, D-lactate and glycolate, support mitochondrial membrane potential and neuronal survival. <i>Biology Open</i> , 2014, 3, 777-784.	0.6	49
78	Pericentriolar material structure and dynamics. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2014, 369, 20130459.	1.8	250
79	Conserved TCP domain of Sas-4/CPAP is essential for pericentriolar material tethering during centrosome biogenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E354-63.	3.3	70
80	Encouraging innovation. <i>Molecular Biology of the Cell</i> , 2014, 25, 427-428.	0.9	1
81	Timing and mechanism of the initial cue establishing handed left-right asymmetry in <i>Caenorhabditis elegans</i> embryos. <i>Genesis</i> , 2014, 52, 572-580.	0.8	38
82	Centrosomes are autocatalytic droplets of pericentriolar material organized by centrioles. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E2636-45.	3.3	187
83	Liquid-Liquid Phase Separation in Biology. <i>Annual Review of Cell and Developmental Biology</i> , 2014, 30, 39-58.	4.0	2,234
84	The <i>Caenorhabditis elegans</i> pericentriolar material components SPD-2 and SPD-5 are monomeric in the cytoplasm before incorporation into the PCM matrix. <i>Molecular Biology of the Cell</i> , 2014, 25, 2984-2992.	0.9	31
85	Quantification of surface tension and internal pressure generated by single mitotic cells. <i>Scientific Reports</i> , 2014, 4, 6213.	1.6	151
86	The Segmentation of Microtubules in Electron Tomograms Using Amira. <i>Methods in Molecular Biology</i> , 2014, 1136, 261-278.	0.4	29
87	XMAP215 activity sets spindle length by controlling the total mass of spindle microtubules. <i>Nature Cell Biology</i> , 2013, 15, 1116-1122.	4.6	115
88	A genomic toolkit to investigate kinesin and myosin motor function in cells. <i>Nature Cell Biology</i> , 2013, 15, 325-334.	4.6	104
89	A Systematic Mammalian Genetic Interaction Map Reveals Pathways Underlying Ricin Susceptibility. <i>Cell</i> , 2013, 152, 909-922.	13.5	332
90	Principles of PAR polarity in <i>Caenorhabditis elegans</i> embryos. <i>Nature Reviews Molecular Cell Biology</i> , 2013, 14, 315-322.	16.1	85

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91	Synergy between XMAP215 and EB1 increases microtubule growth rates to physiological levels. <i>Nature Cell Biology</i> , 2013, 15, 688-693.	4.6	160
92	Spatial Organization of the Cell Cytoplasm by Position-Dependent Phase Separation. <i>Physical Review Letters</i> , 2013, 111, 088101.	2.9	131
93	Funding Innovative Science. <i>Science</i> , 2013, 339, 119-119.	6.0	2
94	Stoichiometry of chromatin-associated protein complexes revealed by label-free quantitative mass spectrometry-based proteomics. <i>Nucleic Acids Research</i> , 2013, 41, e28-e28.	6.5	222
95	C11ORF24 Is a Novel Type I Membrane Protein That Cycles between the Golgi Apparatus and the Plasma Membrane in Rab6-Positive Vesicles. <i>PLoS ONE</i> , 2013, 8, e82223.	1.1	3
96	BICD2, dynactin, and LIS1 cooperate in regulating dynein recruitment to cellular structures. <i>Molecular Biology of the Cell</i> , 2012, 23, 4226-4241.	0.9	231
97	Sds22 and Repo-Man stabilize chromosome segregation by counteracting Aurora B on anaphase kinetochores. <i>Journal of Cell Biology</i> , 2012, 198, 173-183.	2.3	69
98	One-step purification of assembly-competent tubulin from diverse eukaryotic sources. <i>Molecular Biology of the Cell</i> , 2012, 23, 4393-4401.	0.9	125
99	APC15 mediates CDC20 autoubiquitylation by APC/CMCC and disassembly of the mitotic checkpoint complex. <i>Nature Structural and Molecular Biology</i> , 2012, 19, 1116-1123.	3.6	118
100	A Genome-Scale Resource for In Vivo Tag-Based Protein Function Exploration in <i>C. elegans</i> . <i>Cell</i> , 2012, 150, 855-866.	13.5	253
101	Beyond Oil and Water—Phase Transitions in Cells. <i>Science</i> , 2012, 337, 1047-1049.	6.0	217
102	Automated tracing of microtubules in electron tomograms of plastic embedded samples of <i>Caenorhabditis elegans</i> embryos. <i>Journal of Structural Biology</i> , 2012, 178, 129-138.	1.3	101
103	Functional Repurposing Revealed by Comparing <i>S. pombe</i> and <i>S. cerevisiae</i> Genetic Interactions. <i>Cell</i> , 2012, 149, 1339-1352.	13.5	154
104	High-efficiency counterselection recombineering for site-directed mutagenesis in bacterial artificial chromosomes. <i>Nature Methods</i> , 2012, 9, 103-109.	9.0	52
105	GTSE1 Is a Microtubule Plus-End Tracking Protein That Regulates EB1-Dependent Cell Migration. <i>PLoS ONE</i> , 2012, 7, e51259.	1.1	52
106	Tracking mechanics and volume of globular cells with atomic force microscopy using a constant-height clamp. <i>Nature Protocols</i> , 2012, 7, 143-154.	5.5	45
107	Organelle Growth Control through Limiting Pools of Cytoplasmic Components. <i>Current Biology</i> , 2012, 22, R330-R339.	1.8	190
108	Diverse transcription factor binding features revealed by genome-wide ChIP-seq in <i>C. elegans</i> . <i>Genome Research</i> , 2011, 21, 245-254.	2.4	224

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109	Purification of Tubulin from Porcine Brain. <i>Methods in Molecular Biology</i> , 2011, 777, 15-28.	0.4	68
110	Polarization of PAR Proteins by Advective Triggering of a Pattern-Forming System. <i>Science</i> , 2011, 334, 1137-1141.	6.0	290
111	A High-Resolution <i>C.Âlegans</i> Essential Gene Network Based on Phenotypic Profiling of a Complex Tissue. <i>Cell</i> , 2011, 145, 470-482.	13.5	193
112	Samurai Sword Sets Spindle Size. <i>Cell</i> , 2011, 147, 1224-1225.	13.5	1
113	Beyond Stereospecificity: Liquids and Mesoscale Organization of Cytoplasm. <i>Developmental Cell</i> , 2011, 21, 14-16.	3.1	147
114	3K1334 A single-cell RNAi screen for regulators of mitotic cell mechanics identifies diseases-associated genes(<i>Cell biology 4</i> ,The 49th Annual Meeting of the Biophysical Society of Japan). <i>Seibutsu Butsuri</i> , 2011, 51, S146.	0.0	0
115	Novel asymmetrically localizing components of human centrosomes identified by complementary proteomics methods. <i>EMBO Journal</i> , 2011, 30, 1520-1535.	3.5	278
116	Hydrostatic pressure and the actomyosin cortex drive mitotic cell rounding. <i>Nature</i> , 2011, 469, 226-230.	13.7	576
117	Force probing cell shape changes to molecular resolution. <i>Trends in Biochemical Sciences</i> , 2011, 36, 444-450.	3.7	27
118	Limiting Amounts of Centrosome Material Set Centrosome Size in <i>C.Âlegans</i> Embryos. <i>Current Biology</i> , 2011, 21, 1259-1267.	1.8	198
119	Atomic Force Microscopy to Study Mechanics of Living Mitotic Mammalian Cells. <i>Japanese Journal of Applied Physics</i> , 2011, 50, 08LA01.	0.8	1
120	Proliferating versus differentiating stem and cancer cells exhibit distinct midbody-release behaviour. <i>Nature Communications</i> , 2011, 2, 503.	5.8	139
121	Systematic Phosphorylation Analysis of Human Mitotic Protein Complexes. <i>Science Signaling</i> , 2011, 4, rs12.	1.6	87
122	Whither systems biology. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2011, 366, 3635-3637.	1.8	16
123	PAR proteins diffuse freely across the anteriorâ€“posterior boundary in polarized <i>C. elegans</i> embryos. <i>Journal of Cell Biology</i> , 2011, 193, 583-594.	2.3	106
124	The First Cell Cycle of the <i>Caenorhabditis elegans</i> Embryo: Spatial and Temporal Control of an Asymmetric Cell Division. <i>Results and Problems in Cell Differentiation</i> , 2011, 53, 109-133.	0.2	19
125	Active liquid-like behavior of nucleoli determines their size and shape in <i>Xenopus laevis</i> oocytes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 4334-4339.	3.3	1,004
126	XMAP215 polymerase activity is built by combining multiple tubulin-binding TOG domains and a basic lattice-binding region. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 2741-2746.	3.3	143

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127	Centrosome Size Sets Mitotic Spindle Length in <i>Caenorhabditis elegans</i> Embryos. <i>Current Biology</i> , 2010, 20, 353-358.	1.8	181
128	LGL Can Partition the Cortex of One-Cell <i>Caenorhabditis elegans</i> Embryos into Two Domains. <i>Current Biology</i> , 2010, 20, 1296-1303.	1.8	92
129	Phenotypic profiling of the human genome by time-lapse microscopy reveals cell division genes. <i>Nature</i> , 2010, 464, 721-727.	13.7	768
130	Live-cell imaging RNAi screen identifies PP2A ^{B55} and importin ^{β1} as key mitotic exit regulators in human cells. <i>Nature Cell Biology</i> , 2010, 12, 886-893.	4.6	315
131	Cortical domain correction repositions the polarity boundary to match the cytokinesis furrow in <i>C. elegans</i> embryos. <i>Development (Cambridge)</i> , 2010, 137, 1743-1753.	1.2	46
132	Genome-Wide Identification of Binding Sites Defines Distinct Functions for <i>Caenorhabditis elegans</i> PHA-4/FOXA in Development and Environmental Response. <i>PLoS Genetics</i> , 2010, 6, e1000848.	1.5	165
133	Automated tracking and analysis of centrosomes in early <i>Caenorhabditis elegans</i> embryos. <i>Bioinformatics</i> , 2010, 26, i13-i20.	1.8	25
134	FRAP Analysis of Membrane-Associated Proteins: Lateral Diffusion and Membrane-Cytoplasmic Exchange. <i>Biophysical Journal</i> , 2010, 99, 2443-2452.	0.2	63
135	Quantitative Interaction Proteomics and Genome-wide Profiling of Epigenetic Histone Marks and Their Readers. <i>Cell</i> , 2010, 142, 967-980.	13.5	710
136	Sororin Mediates Sister Chromatid Cohesion by Antagonizing Wapl. <i>Cell</i> , 2010, 143, 737-749.	13.5	325
137	Systematic Analysis of Human Protein Complexes Identifies Chromosome Segregation Proteins. <i>Science</i> , 2010, 328, 593-599.	6.0	465
138	Membrane Invaginations Reveal Cortical Sites that Pull on Mitotic Spindles in One-Cell <i>C. elegans</i> Embryos. <i>PLoS ONE</i> , 2010, 5, e12301.	1.1	96
139	HAUS, the 8-Subunit Human Augmin Complex, Regulates Centrosome and Spindle Integrity. <i>Current Biology</i> , 2009, 19, 816-826.	1.8	231
140	Comparative profiling identifies C13orf3 as a component of the Ska complex required for mammalian cell division. <i>EMBO Journal</i> , 2009, 28, 1453-1465.	3.5	89
141	Growth, fluctuation and switching at microtubule plus ends. <i>Nature Reviews Molecular Cell Biology</i> , 2009, 10, 569-574.	16.1	152
142	Germline P Granules Are Liquid Droplets That Localize by Controlled Dissolution/Condensation. <i>Science</i> , 2009, 324, 1729-1732.	6.0	2,267
143	EB1 Recognizes the Nucleotide State of Tubulin in the Microtubule Lattice. <i>PLoS ONE</i> , 2009, 4, e7585.	1.1	137
144	BAC TransgeneOmics: a high-throughput method for exploration of protein function in mammals. <i>Nature Methods</i> , 2008, 5, 409-415.	9.0	568

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145	The Mammalian SPD-2 Ortholog Cep192 Regulates Centrosome Biogenesis. <i>Current Biology</i> , 2008, 18, 136-141.	1.8	169
146	Characterization of Protein Dynamics in Asymmetric Cell Division by Scanning Fluorescence Correlation Spectroscopy. <i>Biophysical Journal</i> , 2008, 95, 5476-5486.	0.2	52
147	Efficient chaperone-mediated tubulin biogenesis is essential for cell division and cell migration in <i>C. elegans</i> . <i>Developmental Biology</i> , 2008, 313, 320-334.	0.9	66
148	XMAP215 Is a Processive Microtubule Polymerase. <i>Cell</i> , 2008, 132, 79-88.	13.5	479
149	A Protein Domain-Based Interactome Network for <i>C. elegans</i> Early Embryogenesis. <i>Cell</i> , 2008, 134, 534-545.	13.5	196
150	Cell cycle progression requires the CDC-48 ^{UFD-1/NPL-4} complex for efficient DNA replication. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 12879-12884.	3.3	69
151	Building a spindle of the correct length in human cells requires the interaction between TPX2 and Aurora A. <i>Journal of Cell Biology</i> , 2008, 182, 289-300.	2.3	178
152	Two-photon fluorescence imaging and correlation analysis applied to protein dynamics in <i>C. elegans</i> embryo. , 2008, , .		9
153	Acto-myosin reorganization and PAR polarity in <i>C. elegans</i> . <i>Development (Cambridge)</i> , 2007, 134, 1035-1043.	1.2	102
154	Functional Interaction between Phosducin-like Protein 2 and Cytosolic Chaperonin Is Essential for Cytoskeletal Protein Function and Cell Cycle Progression. <i>Molecular Biology of the Cell</i> , 2007, 18, 2336-2345.	0.9	50
155	The Rho GTPase-activating proteins RGA-3 and RGA-4 are required to set the initial size of PAR domains in <i>Caenorhabditis elegans</i> one-cell embryos. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 14976-14981.	3.3	112
156	The <i>C. elegans</i> RSA Complex Localizes Protein Phosphatase 2A to Centrosomes and Regulates Mitotic Spindle Assembly. <i>Cell</i> , 2007, 128, 115-127.	13.5	87
157	Stress Generation and Filament Turnover during Actin Ring Constriction. <i>PLoS ONE</i> , 2007, 2, e696.	1.1	99
158	Genome-scale RNAi profiling of cell division in human tissue culture cells. <i>Nature Cell Biology</i> , 2007, 9, 1401-1412.	4.6	270
159	Microtubule polymerases and depolymerases. <i>Current Opinion in Cell Biology</i> , 2007, 19, 31-35.	2.6	267
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