

Jason R Swedlow

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/3447931/jason-r-swedlow-publications-by-year.pdf>

Version: 2024-04-10

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

110 papers	7,506 citations	41 h-index	86 g-index
139 ext. papers	8,635 ext. citations	12.2 avg, IF	5.61 L-index

#	Paper	IF	Citations
110	The BioImage Archive - building a home for life-sciences microscopy data.. <i>Journal of Molecular Biology</i> , 2022 , 167505	6.5	3
109	MITI minimum information guidelines for highly multiplexed tissue images.. <i>Nature Methods</i> , 2022 , 19, 262-267	21.6	2
108	OME-NGFF: a next-generation file format for expanding bioimaging data-access strategies. <i>Nature Methods</i> , 2021 , 18, 1496-1498	21.6	13
107	A global view of standards for open image data formats and repositories. <i>Nature Methods</i> , 2021 , 18, 1440-1446	21.6	18
106	REMBI: Recommended Metadata for Biological Images-enabling reuse of microscopy data in biology. <i>Nature Methods</i> , 2021 , 18, 1418-1422	21.6	16
105	QUAREP-LiMi: a community endeavor to advance quality assessment and reproducibility in light microscopy. <i>Nature Methods</i> , 2021 , 18, 1423-1426	21.6	18
104	ISO 23494: Biotechnology [Provenance Information Model for Biological Specimen And Data. <i>Lecture Notes in Computer Science</i> , 2021 , 222-225	0.9	2
103	QUAREP-LiMi: A community-driven initiative to establish guidelines for quality assessment and reproducibility for instruments and images in light microscopy. <i>Journal of Microscopy</i> , 2021 , 284, 56-73	1.9	11
102	Numerically Enhanced Stimulated Emission Depletion Microscopy with Adaptive Optics for Deep-Tissue Super-Resolved Imaging. <i>ACS Nano</i> , 2020 , 14, 394-405	16.7	12
101	Community standards for open cell migration data. <i>GigaScience</i> , 2020 , 9,	7.6	9
100	Bringing Open Data to Whole Slide Imaging. <i>Lecture Notes in Computer Science</i> , 2019 , 2019, 3-10	0.9	11
99	Big Data in Correlative Imaging 2019 , 211-222		
98	Full volume super-resolution imaging of thick mitotic spindle using 3D AO STED microscope. <i>Biomedical Optics Express</i> , 2019 , 10, 1999-2009	3.5	17
97	Killing with proficiency: Integrated post-translational regulation of an offensive Type VI secretion system. <i>PLoS Pathogens</i> , 2018 , 14, e1007230	7.6	12
96	A call for public archives for biological image data. <i>Nature Methods</i> , 2018 , 15, 849-854	21.6	61
95	Visualization of Biomedical Data. <i>Annual Review of Biomedical Data Science</i> , 2018 , 1, 275-304	5.6	34
94	The Image Data Resource: A Bioimage Data Integration and Publication Platform. <i>Nature Methods</i> , 2017 , 14, 775-781	21.6	130

93	The Ndc80 complex targets Bod1 to human mitotic kinetochores. <i>Open Biology</i> , 2017 , 7,	7	7
92	The Open Microscopy Environment: open image informatics for the biological sciences 2016 ,		1
91	Metadata management for high content screening in OMERO. <i>Methods</i> , 2016 , 96, 27-32	4.6	23
90	CDK-dependent phosphorylation of PHD1 on serine 130 alters its substrate preference in cells. <i>Journal of Cell Science</i> , 2016 , 129, 191-205	5.3	13
89	BOD1 Is Required for Cognitive Function in Humans and Drosophila. <i>PLoS Genetics</i> , 2016 , 12, e1006022	6	15
88	Publishing and sharing multi-dimensional image data with OMERO. <i>Mammalian Genome</i> , 2015 , 26, 441-73.	2	20
87	OMERO and Bio-Formats 5: flexible access to large bioimaging datasets at scale 2015 ,		3
86	An open data ecosystem for cell migration research. <i>Trends in Cell Biology</i> , 2015 , 25, 55-8	18.3	21
85	Phosphorylated DegU manipulates cell fate differentiation in the Bacillus subtilis biofilm. <i>Journal of Bacteriology</i> , 2014 , 196, 16-27	3.5	40
84	Analysis of global RNA synthesis at the single cell level following hypoxia. <i>Journal of Visualized Experiments</i> , 2014 ,	1.6	2
83	A 3D cellular context for the macromolecular world. <i>Nature Structural and Molecular Biology</i> , 2014 , 21, 841-5	17.6	33
82	Quantitative fluorescence microscopy and image deconvolution. <i>Methods in Cell Biology</i> , 2013 , 114, 407-26		29
81	Quantitative analysis of digital microscope images. <i>Methods in Cell Biology</i> , 2013 , 114, 337-67	1.8	6
80	At the (kineto)chore, yeast really are like people. <i>Cell</i> , 2013 , 154, 959-961	56.2	
79	PHD1 links cell-cycle progression to oxygen sensing through hydroxylation of the centrosomal protein Cep192. <i>Developmental Cell</i> , 2013 , 26, 381-92	10.2	57
78	Web-based visualisation and analysis of 3D electron-microscopy data from EMDB and PDB. <i>Journal of Structural Biology</i> , 2013 , 184, 173-81	3.4	23
77	Bod1 regulates protein phosphatase 2A at mitotic kinetochores. <i>Nature Communications</i> , 2013 , 4, 2677	17.4	58
76	Ubiquitylation-dependent localization of PLK1 in mitosis. <i>Nature Cell Biology</i> , 2013 , 15, 430-9	23.4	70

75	OMERO: flexible, model-driven data management for experimental biology. <i>Nature Methods</i> , 2012 , 9, 245-53	21.6	313
74	Software tools, data structures, and interfaces for microscope imaging. <i>Cold Spring Harbor Protocols</i> , 2012 , 2012, 50-61	1.2	8
73	The adenomatous polyposis coli protein contributes to normal compaction of mitotic chromatin. <i>PLoS ONE</i> , 2012 , 7, e38102	3.7	10
72	High-resolution live imaging of cell behavior in the developing neuroepithelium. <i>Journal of Visualized Experiments</i> , 2012 ,	1.6	10
71	Biological imaging software tools. <i>Nature Methods</i> , 2012 , 9, 697-710	21.6	377
70	Innovation in biological microscopy: current status and future directions. <i>BioEssays</i> , 2012 , 34, 333-40	4.1	26
69	Data management challenges in three-dimensional EM. <i>Nature Structural and Molecular Biology</i> , 2012 , 19, 1203-7	17.6	34
68	Fixation of <i>Drosophila</i> tissues for immunofluorescence. <i>Cold Spring Harbor Protocols</i> , 2011 , 2011, 931-4	1.2	3
67	Finding an image in a haystack: the case for public image repositories. <i>Nature Cell Biology</i> , 2011 , 13, 183	23.4	8
66	How to be a mitotic chromosome. <i>Chromosome Research</i> , 2011 , 19, 307-19	4.4	21
65	A novel reporter of notch signalling indicates regulated and random Notch activation during vertebrate neurogenesis. <i>BMC Biology</i> , 2011 , 9, 58	7.3	31
64	Immunolabeling of <i>Drosophila</i> embryos and tissues. <i>Cold Spring Harbor Protocols</i> , 2011 , 2011, 998-1002	1.2	3
63	OMX: a new platform for multimodal, multichannel wide-field imaging. <i>Cold Spring Harbor Protocols</i> , 2011 , 2011, 899-909	1.2	32
62	Sds22 regulates aurora B activity and microtubule-kinetochore interactions at mitosis. <i>Journal of Cell Biology</i> , 2010 , 191, 61-74	7.3	101
61	Kinetochore alignment within the metaphase plate is regulated by centromere stiffness and microtubule depolymerases. <i>Journal of Cell Biology</i> , 2010 , 188, 665-79	7.3	106
60	Advanced hardware and software tools for fast multidimensional imaging of living cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 16005-6	11.5	7
59	Metadata matters: access to image data in the real world. <i>Journal of Cell Biology</i> , 2010 , 189, 777-82	7.3	544
58	The Nup107-160 nucleoporin complex promotes mitotic events via control of the localization state of the chromosome passenger complex. <i>Molecular Biology of the Cell</i> , 2009 , 20, 5260-75	3.5	64

57	Open source bioimage informatics for cell biology. <i>Trends in Cell Biology</i> , 2009 , 19, 656-60	18.3	43
56	Usability and User-Centered Design in Scientific Software Development. <i>IEEE Software</i> , 2009 , 26, 96-102	1.5	24
55	In vivo imaging of mammalian cells: cell engineering and viability. <i>Cold Spring Harbor Protocols</i> , 2009 , 2009, pdb.ip69	1.2	1
54	In vivo imaging of mammalian cells: image acquisition and analysis. <i>Cold Spring Harbor Protocols</i> , 2009 , 2009, pdb.ip70	1.2	2
53	Bioimage informatics for experimental biology. <i>Annual Review of Biophysics</i> , 2009 , 38, 327-46	21.1	82
52	Building a scientific data grid with DiGS. <i>Philosophical Transactions Series A, Mathematical, Physical, and Engineering Sciences</i> , 2009 , 367, 2471-81	3	1
51	Minimum information specification for in situ hybridization and immunohistochemistry experiments (MISFISHIE). <i>Nature Biotechnology</i> , 2008 , 26, 305-12	44.5	97
50	Temporal profiling of the chromatin proteome reveals system-wide responses to replication inhibition. <i>Current Biology</i> , 2008 , 18, 838-43	6.3	32
49	Open tools for storage and management of quantitative image data. <i>Methods in Cell Biology</i> , 2008 , 85, 555-70	1.8	24
48	Open File Formats for High Content Analysis 2008 , 317-328		
47	Evaluating performance in three-dimensional fluorescence microscopy. <i>Journal of Microscopy</i> , 2007 , 228, 390-405	1.9	99
46	ELYS/MEL-28 chromatin association coordinates nuclear pore complex assembly and replication licensing. <i>Current Biology</i> , 2007 , 17, 1657-62	6.3	110
45	Phosphorylation by aurora-B negatively regulates survivin function during mitosis. <i>Cell Cycle</i> , 2007 , 6, 1220-30	4.7	45
44	Mitotic spindle orientation distinguishes stem cell and terminal modes of neuron production in the early spinal cord. <i>Development (Cambridge)</i> , 2007 , 134, 1943-54	6.6	79
43	Bod1, a novel kinetochore protein required for chromosome biorientation. <i>Journal of Cell Biology</i> , 2007 , 179, 187-97	7.3	44
42	Quantitative fluorescence microscopy and image deconvolution. <i>Methods in Cell Biology</i> , 2007 , 81, 447-65	5.8	39
41	Quantitative analysis of digital microscope images. <i>Methods in Cell Biology</i> , 2007 , 81, 365-96	1.8	39
40	The Open Microscopy Environment: A Collaborative Data Modeling and Software Development Project for Biological Image Informatics. <i>Principles and Practice</i> , 2007 , 71-92		4

39	Development of the Minimum Information Specification for In Situ Hybridization and Immunohistochemistry Experiments (MISFISHIE). <i>OMICS A Journal of Integrative Biology</i> , 2006 , 10, 205-8	3.8	16
38	Modelling data across labs, genomes, space and time. <i>Nature Cell Biology</i> , 2006 , 8, 1190-4	23.4	25
37	The Open Microscopy Environment (OME) Data Model and XML file: open tools for informatics and quantitative analysis in biological imaging. <i>Genome Biology</i> , 2005 , 6, R47	18.3	186
36	Phosphorylation regulates the dynamic interaction of RCC1 with chromosomes during mitosis. <i>Current Biology</i> , 2004 , 14, 1099-104	6.3	67
35	Chromosome condensation: DNA compaction in real time. <i>Current Biology</i> , 2004 , 14, R554-6	6.3	5
34	Optimisation of the two-dimensional gel electrophoresis protocol using the Taguchi approach. <i>Proteome Science</i> , 2004 , 2, 6	2.6	34
33	Aurora B regulates MCAK at the mitotic centromere. <i>Developmental Cell</i> , 2004 , 6, 253-68	10.2	431
32	Mitotic mechanics: the auroras come into view. <i>Current Opinion in Cell Biology</i> , 2003 , 15, 672-83	9	249
31	Quantitative fluorescence microscopy and image deconvolution. <i>Methods in Cell Biology</i> , 2003 , 72, 349-67	7.8	21
30	Human CLASP1 is an outer kinetochore component that regulates spindle microtubule dynamics. <i>Cell</i> , 2003 , 113, 891-904	56.2	177
29	The making of the mitotic chromosome: modern insights into classical questions. <i>Molecular Cell</i> , 2003 , 11, 557-69	17.6	223
28	Informatics and quantitative analysis in biological imaging. <i>Science</i> , 2003 , 300, 100-2	33.3	214
27	Time-lapse imaging reveals dynamic relocalization of PP1gamma throughout the mammalian cell cycle. <i>Molecular Biology of the Cell</i> , 2003 , 14, 107-17	3.5	122
26	Live cell imaging using wide-field microscopy and deconvolution. <i>Cell Structure and Function</i> , 2002 , 27, 335-41	2.2	75
25	Characterization and Use of Wide-Field Fluorescence Microscopy and Image Restoration in Quantitative Live Cell Imaging. <i>Microscopy and Microanalysis</i> , 2002 , 8, 266-267	0.5	
24	To 5D and beyond: quantitative fluorescence microscopy in the postgenomic era. <i>Traffic</i> , 2002 , 3, 29-36	5.7	47
23	Cajal body dynamics and association with chromatin are ATP-dependent. <i>Nature Cell Biology</i> , 2002 , 4, 502-8	23.4	232
22	Measuring tubulin content in <i>Toxoplasma gondii</i> : a comparison of laser-scanning confocal and wide-field fluorescence microscopy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002 , 99, 2014-9	11.5	119

21	A role for the Adenomatous Polyposis Coli protein in chromosome segregation. <i>Nature Cell Biology</i> , 2001 , 3, 429-32	23.4	453
20	Phosphorylation of threonine 156 of the mu2 subunit of the AP2 complex is essential for endocytosis in vitro and in vivo. <i>Current Biology</i> , 2001 , 11, 896-900	6.3	104
19	Chromatin-associated protein phosphatase 1 regulates aurora-B and histone H3 phosphorylation. <i>Journal of Biological Chemistry</i> , 2001 , 276, 26656-65	5.4	201
18	Nuclear dynamics: where genes are and how they got there. <i>Genome Biology</i> , 2001 , 2, REVIEWS0002	18.3	15
17	A workingperson's guide to deconvolution in light microscopy. <i>BioTechniques</i> , 2001 , 31, 1076-8, 1080, 1082 passim	2.5	259
16	Defining the Tools: an Analysis of Laser Scanning Confocal and Wide-Field/Restoration Fluorescence Microscope Imaging. <i>Microscopy and Microanalysis</i> , 2001 , 7, 1002-1003	0.5	
15	In vivo analysis of Cajal body movement, separation, and joining in live human cells. <i>Journal of Cell Biology</i> , 2000 , 151, 1561-74	7.3	223
14	Actin-dependent localization of an RNA encoding a cell-fate determinant in yeast. <i>Nature</i> , 1997 , 389, 90-3	50.4	335
13	Fuzzy sequences, specific attachments? Chromosome dynamics. <i>Current Biology</i> , 1996 , 6, 544-7	6.3	4
12	The SMC family: from chromosome condensation to dosage compensation. <i>Current Opinion in Cell Biology</i> , 1995 , 7, 329-36	9	88
11	The Collection, Processing, and Display of Digital Three-Dimensional Images of Biological Specimens 1995 , 197-210		37
10	Chromosome structure inside the nucleus. <i>Current Opinion in Cell Biology</i> , 1993 , 5, 412-6	9	20
9	High-resolution 3-D and 4-D imaging using wide-field CCD-based microscopy. <i>Proceedings Annual Meeting Electron Microscopy Society of America</i> , 1993 , 51, 150-151		
8	The in vivo distribution and dynamics of DNA topoisomerase II in Drosophila embryonic nuclei and chromosomes. <i>Proceedings Annual Meeting Electron Microscopy Society of America</i> , 1993 , 51, 74-75		
7	Chromosome structure and dynamics as revealed by 3-D and 4-D imaging. <i>Proceedings Annual Meeting Electron Microscopy Society of America</i> , 1992 , 50, 588-589		
6	Chromosome structure and dynamics as revealed by 3-D and 4-D imaging. <i>Proceedings Annual Meeting Electron Microscopy Society of America</i> , 1991 , 49, 396-397		
5	Focal points for chromosome condensation and decondensation revealed by three-dimensional in vivo time-lapse microscopy. <i>Nature</i> , 1989 , 342, 293-6	50.4	99
4	Community Standards for Open Cell Migration Data		3

3	OME Files - An open source reference library for the OME-XML metadata model and the OME-TIFF file format	3
2	The Image Data Resource: A Scalable Platform for Biological Image Data Access, Integration, and Dissemination	
1	OME-NGFF: scalable format strategies for interoperable bioimaging data	10