

# Thomas Walter

## List of Publications by Citations

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

49 papers	3,544 citations	25 h-index	59 g-index
60 ext. papers	4,360 ext. citations	10.9 avg, IF	4.9 L-index

#	Paper	IF	Citations
49	Phenotypic profiling of the human genome by time-lapse microscopy reveals cell division genes. <i>Nature</i> , <b>2010</b> , 464, 721-7	50.4	668
48	A contribution of image processing to the diagnosis of diabetic retinopathy--detection of exudates in color fundus images of the human retina. <i>IEEE Transactions on Medical Imaging</i> , <b>2002</b> , 21, 1236-43	11.7	477
47	CellCognition: time-resolved phenotype annotation in high-throughput live cell imaging. <i>Nature Methods</i> , <b>2010</b> , 7, 747-54	21.6	256
46	Assessment of algorithms for mitosis detection in breast cancer histopathology images. <i>Medical Image Analysis</i> , <b>2015</b> , 20, 237-48	15.4	245
45	Automatic detection of microaneurysms in color fundus images. <i>Medical Image Analysis</i> , <b>2007</b> , 11, 555-66	15.4	212
44	Visualization of image data from cells to organisms. <i>Nature Methods</i> , <b>2010</b> , 7, S26-41	21.6	189
43	Reverse transfection on cell arrays for high content screening microscopy. <i>Nature Protocols</i> , <b>2007</b> , 2, 392-9	18.8	162
42	Segmentation of Nuclei in Histopathology Images by Deep Regression of the Distance Map. <i>IEEE Transactions on Medical Imaging</i> , <b>2019</b> , 38, 448-459	11.7	158
41	smiFISH and FISH-quant - a flexible single RNA detection approach with super-resolution capability. <i>Nucleic Acids Research</i> , <b>2016</b> , 44, e165	20.1	158
40	Micropilot: automation of fluorescence microscopy-based imaging for systems biology. <i>Nature Methods</i> , <b>2011</b> , 8, 246-9	21.6	107
39	Segmentation of Color Fundus Images of the Human Retina: Detection of the Optic Disc and the Vascular Tree Using Morphological Techniques. <i>Lecture Notes in Computer Science</i> , <b>2001</b> , 282-287	0.9	98
38	Visualizing biological data-now and in the future. <i>Nature Methods</i> , <b>2010</b> , 7, S2-4	21.6	85
37	Prediction of human population responses to toxic compounds by a collaborative competition. <i>Nature Biotechnology</i> , <b>2015</b> , 33, 933-40	44.5	70
36	Nuclei segmentation in histopathology images using deep neural networks <b>2017</b> ,		67
35	Waterpixels. <i>IEEE Transactions on Image Processing</i> , <b>2015</b> , 24, 3707-16	8.7	66
34	Evaluation of automated fundus photograph analysis algorithms for detecting microaneurysms, haemorrhages and exudates, and of a computer-assisted diagnostic system for grading diabetic retinopathy. <i>Diabetes and Metabolism</i> , <b>2010</b> , 36, 213-20	5.4	65
33	Mitotic lamin disassembly is triggered by lipid-mediated signaling. <i>Journal of Cell Biology</i> , <b>2012</b> , 198, 981-90	7.3	50

32	Automatic identification and clustering of chromosome phenotypes in a genome wide RNAi screen by time-lapse imaging. <i>Journal of Structural Biology</i> , <b>2010</b> , 170, 1-9	3.4	42
31	Human lymphoid organ cDC2 and macrophages play complementary roles in T follicular helper responses. <i>Journal of Experimental Medicine</i> , <b>2019</b> , 216, 1561-1581	16.6	36
30	Integration of biological data by kernels on graph nodes allows prediction of new genes involved in mitotic chromosome condensation. <i>Molecular Biology of the Cell</i> , <b>2014</b> , 25, 2522-36	3.5	36
29	EML3 is a nuclear microtubule-binding protein required for the correct alignment of chromosomes in metaphase. <i>Journal of Cell Science</i> , <b>2008</b> , 121, 1718-26	5.3	35
28	A Dual Protein-mRNA Localization Screen Reveals Compartmentalized Translation and Widespread Co-translational RNA Targeting. <i>Developmental Cell</i> , <b>2020</b> , 54, 773-791.e5	10.2	33
27	A computational framework to study sub-cellular RNA localization. <i>Nature Communications</i> , <b>2018</b> , 9, 4584	7.4	29
26	A genomic Multiprocess survey of machineries that control and link cell shape, microtubule organization, and cell-cycle progression. <i>Developmental Cell</i> , <b>2014</b> , 31, 227-239	10.2	26
25	Automatic Detection of Microaneurysms in Color Fundus Images of the Human Retina by Means of the Bounding Box Closing. <i>Lecture Notes in Computer Science</i> , <b>2002</b> , 210-220	0.9	26
24	Anti-angiogenic properties of myo-inositol trispyrophosphate in ovo and growth reduction of implanted glioma. <i>FEBS Letters</i> , <b>2007</b> , 581, 962-6	3.8	24
23	ARHGEF17 is an essential spindle assembly checkpoint factor that targets Mps1 to kinetochores. <i>Journal of Cell Biology</i> , <b>2016</b> , 212, 647-59	7.3	14
22	A choreography of centrosomal mRNAs reveals a conserved localization mechanism involving active polysome transport. <i>Nature Communications</i> , <b>2021</b> , 12, 1352	17.4	14
21	A generic methodological framework for studying single cell motility in high-throughput time-lapse data. <i>Bioinformatics</i> , <b>2015</b> , 31, i320-8	7.2	13
20	MAP1S controls microtubule stability throughout the cell cycle in human cells. <i>Journal of Cell Science</i> , <b>2014</b> , 127, 5007-13	5.3	11
19	Dynamical modelling of phenotypes in a genome-wide RNAi live-cell imaging assay. <i>BMC Bioinformatics</i> , <b>2013</b> , 14, 308	3.6	10
18	New general features based on superpixels for image segmentation learning <b>2016</b> ,		9
17	Waterpixels: Superpixels based on the watershed transformation <b>2014</b> ,		9
16	Automatic Analysis of Color Fundus Photographs and Its Application to the Diagnosis of Diabetic Retinopathy <b>2005</b> , 315-368		8
15	The kinesin KIF1C transports APC-dependent mRNAs to cell protrusions. <i>Rna</i> , <b>2021</b> , 27, 1528-1544	5.8	6

14	High-throughput microscopy using live mammalian cells. <i>Cold Spring Harbor Protocols</i> , <b>2010</b> , 2010, pdb.top84	5
13	Domain-invariant features for mechanism of action prediction in a multi-cell-line drug screen. <i>Bioinformatics</i> , <b>2020</b> , 36, 1607-1613	7.2 4
12	Spatial Repulsion Between Markers Improves Watershed Performance. <i>Lecture Notes in Computer Science</i> , <b>2015</b> , 194-202	0.9 3
11	Kernel Multitask Regression for Toxicogenetics. <i>Molecular Informatics</i> , <b>2017</b> , 36, 1700053	3.8 2
10	A Deep Learning Approach To Identify mRNA Localization Patterns <b>2019</b> ,	2
9	Redistribution of the neurosensory retina in inferior limited macular translocation: an evaluation using image registration. <i>Graefes Archive for Clinical and Experimental Ophthalmology</i> , <b>2007</b> , 245, 437-42 <sup>3.8</sup>	2
8	The kinesin KIF1C transports APC-dependent mRNAs to cell protrusions	2
7	Spatial transcriptomics for respiratory research and medicine. <i>European Respiratory Journal</i> , <b>2021</b> , 58,	13.6 2
6	FISH-quant v2: a scalable and modular tool for smFISH image analysis.. <i>Rna</i> , <b>2022</b> ,	5.8 2
5	Infering an ontology of single cell motions from high-throughput microscopy data <b>2015</b> ,	1
4	Predicting Residual Cancer Burden In A Triple Negative Breast Cancer Cohort <b>2019</b> ,	1
3	A Localization Screen Reveals Translation Factories and Widespread Co-Translational Protein Targeting. <i>SSRN Electronic Journal</i> ,	1 1
2	FISH-quant v2: a scalable and modular analysis tool for smFISH image analysis	1
1	<b>2018</b> ,	1