

Thomas Boudier

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2465709/publications.pdf>

Version: 2024-02-01

51
papers

3,105
citations

236925

25
h-index

223800

46
g-index

61
all docs

61
docs citations

61
times ranked

5706
citing authors

#	ARTICLE	IF	CITATIONS
1	TANGO: a generic tool for high-throughput 3D image analysis for studying nuclear organization. <i>Bioinformatics</i> , 2013, 29, 1840-1841.	4.1	648
2	Three-Dimensional Architecture of Presynaptic Terminal Cytomatrix. <i>Journal of Neuroscience</i> , 2007, 27, 6868-6877.	3.6	280
3	Tomoj: tomography software for three-dimensional reconstruction in transmission electron microscopy. <i>BMC Bioinformatics</i> , 2007, 8, 288.	2.6	264
4	Electron tomography of early melanosomes: Implications for melanogenesis and the generation of fibrillar amyloid sheets. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 19726-19731.	7.1	133
5	DiAna, an ImageJ tool for object-based 3D co-localization and distance analysis. <i>Methods</i> , 2017, 115, 55-64.	3.8	132
6	Analysis of synaptic ultrastructure without fixative using high-pressure freezing and tomography. <i>European Journal of Neuroscience</i> , 2006, 24, 3463-3474.	2.6	128
7	Tangentially Migrating Neurons Assemble a Primary Cilium that Promotes Their Reorientation to the Cortical Plate. <i>Neuron</i> , 2012, 76, 1108-1122.	8.1	127
8	A-type Lamins Form Distinct Filamentous Networks with Differential Nuclear Pore Complex Associations. <i>Current Biology</i> , 2016, 26, 2651-2658.	3.9	127
9	Macrophages provide a transient muscle stem cell niche via NAMPT secretion. <i>Nature</i> , 2021, 591, 281-287.	27.8	111
10	Progerin reduces LAP2 \pm -telomere association in Hutchinson-Gilford progeria. <i>ELife</i> , 2015, 4, .	6.0	96
11	Structural basis for delta cell paracrine regulation in pancreatic islets. <i>Nature Communications</i> , 2019, 10, 3700.	12.8	80
12	Multiple binding of repressed mRNAs by the P-body protein Rck/p54. <i>Rna</i> , 2012, 18, 1702-1715.	3.5	79
13	From high-resolution AFM topographs to atomic models of supramolecular assemblies. <i>Journal of Structural Biology</i> , 2007, 159, 268-276.	2.8	70
14	Effector and stem-like memory cell fates are imprinted in distinct lymph node niches directed by CXCR3 ligands. <i>Nature Immunology</i> , 2021, 22, 434-448.	14.5	66
15	3D FISH for the quantification of methane- and sulphur-oxidizing endosymbionts in bacteriocytes of the hydrothermal vent mussel <i>Bathymodiolus azoricus</i> . <i>ISME Journal</i> , 2008, 2, 284-292.	9.8	61
16	Structural Information, Resolution, and Noise in High-Resolution Atomic Force Microscopy Topographs. <i>Biophysical Journal</i> , 2009, 96, 3822-3831.	0.5	51
17	Biom mineralization Patterns of Intracellular Carbonatogenesis in Cyanobacteria: Molecular Hypotheses. <i>Minerals (Basel, Switzerland)</i> , 2016, 6, 10.	2.0	48
18	Smart 3D-fish: Automation of distance analysis in nuclei of interphase cells by image processing. <i>Cytometry Part A: the Journal of the International Society for Analytical Cytology</i> , 2005, 67A, 18-26.	1.5	41

#	ARTICLE	IF	CITATIONS
19	Software for drift compensation, particle tracking and particle analysis of high-speed atomic force microscopy image series. <i>Journal of Molecular Recognition</i> , 2012, 25, 292-298.	2.1	39
20	A generic classification-based method for segmentation of nuclei in 3D images of early embryos. <i>BMC Bioinformatics</i> , 2014, 15, 9.	2.6	36
21	Shifting meiotic to mitotic spindle assembly in oocytes disrupts chromosome alignment. <i>EMBO Reports</i> , 2018, 19, 368-381.	4.5	30
22	A new automated 3D detection of synaptic contacts reveals the formation of cortico-striatal synapses upon cocaine treatment in vivo. <i>Brain Structure and Function</i> , 2015, 220, 2953-2966.	2.3	29
23	An Erg-driven transcriptional program controls B cell lymphopoiesis. <i>Nature Communications</i> , 2020, 11, 3013.	12.8	29
24	Organising multi-dimensional biological image information: The BioImage Database. <i>Nucleic Acids Research</i> , 1999, 27, 280-283.	14.5	28
25	High-Speed Atomic Force Microscopy: Cooperative Adhesion and Dynamic Equilibrium of Junctional Microdomain Membrane Proteins. <i>Journal of Molecular Biology</i> , 2012, 423, 249-256.	4.2	27
26	Rapid Synaptogenesis in the Nucleus Accumbens Is Induced by a Single Cocaine Administration and Stabilized by Mitogen-Activated Protein Kinase Interacting Kinase-1 Activity. <i>Biological Psychiatry</i> , 2017, 82, 806-818.	1.3	27
27	Electron tomography of biological samples. <i>Biochemistry (Moscow)</i> , 2004, 69, 1219-1225.	1.5	25
28	Functional Assessment of Genetic Variants with Outcomes Adapted to Clinical Decision-Making. <i>PLoS Genetics</i> , 2016, 12, e1006096.	3.5	24
29	Multiple-axis tomography: applications to basal bodies from <i>Paramecium tetraurelia</i> . <i>Biology of the Cell</i> , 2006, 98, 415-425.	2.0	23
30	The site of breast cancer metastases dictates their clonal composition and reversible transcriptomic profile. <i>Science Advances</i> , 2021, 7, .	10.3	23
31	Video on the Internet: An Introduction to the Digital Encoding, Compression, and Transmission of Moving Image Data. <i>Journal of Structural Biology</i> , 1999, 125, 133-155.	2.8	22
32	Simultaneous localization of MLL, AF4 and ENL genes in interphase nuclei by 3D-FISH: MLL translocation revisited. <i>BMC Cancer</i> , 2006, 6, 20.	2.6	18
33	A novel toolbox to investigate tissue spatial organization applied to the study of the islets of Langerhans. <i>Scientific Reports</i> , 2017, 7, 44261.	3.3	17
34	EM-stellar: benchmarking deep learning for electron microscopy image segmentation. <i>Bioinformatics</i> , 2021, 37, 97-106.	4.1	16
35	Use of cryo-negative staining in tomographic reconstruction of biological objects: application to T4 bacteriophage. <i>Biology of the Cell</i> , 2003, 95, 393-398.	2.0	14
36	OpenSegSPIM: a user-friendly segmentation tool for SPIM data. <i>Bioinformatics</i> , 2016, 32, 2075-2077.	4.1	14

#	ARTICLE	IF	CITATIONS
37	Analysis of Nuclear Organization with TANGO, Software for High-Throughput Quantitative Analysis of 3D Fluorescence Microscopy Images. <i>Methods in Molecular Biology</i> , 2015, 1228, 203-222.	0.9	14
38	Persistent homology for object segmentation in multidimensional grayscale images. <i>Pattern Recognition Letters</i> , 2018, 112, 277-284.	4.2	13
39	Proliferation-dependent positioning of individual centromeres in the interphase nucleus of human lymphoblastoid cell lines. <i>Molecular Biology of the Cell</i> , 2015, 26, 2550-2560.	2.1	12
40	A novel generic dictionary-based denoising method for improving noisy and densely packed nuclei segmentation in 3D time-lapse fluorescence microscopy images. <i>Scientific Reports</i> , 2019, 9, 5654.	3.3	12
41	Direct Image-Based Correlative Microscopy Technique for Coupling Identification and Structural Investigation of Bacterial Symbionts Associated with Metazoans. <i>Applied and Environmental Microbiology</i> , 2011, 77, 4172-4179.	3.1	11
42	Tomography of bacteriaâ€“mineral associations within the deep-sea hydrothermal vent shrimp <i>Rimicaris exoculata</i> . <i>Comptes Rendus Chimie</i> , 2008, 11, 268-280.	0.5	10
43	EM-net: Deep learning for electron microscopy image segmentation. , 2021, , .		9
44	Chromosomes distribute randomly to, but not within, human neutrophil nuclear lobes. <i>IScience</i> , 2021, 24, 102161.	4.1	8
45	NucleiNet: A convolutional encoder-decoder network for bio-image denoising. , 2017, 2017, 1986-1989.		7
46	VIDOS, a system for video editing and format conversion over the Internet. <i>Computer Networks</i> , 2000, 34, 931-944.	5.1	4
47	Optimal processing for gel electrophoresis images: Applying Monte Carlo Tree Search in GelApp. <i>Electrophoresis</i> , 2016, 37, 2208-2216.	2.4	3
48	A dictionary-based approach to reduce noise in fluorescent microscopy images. , 2017, , .		2
49	Detection of high-grade atypia nuclei in breast cancer imaging. , 2015, , .		1
50	3D Analysis and Reconstruction of the Chitin Secreting Gland of <i>Riftia pachyptila</i> . <i>Journal of Biological Systems</i> , 1997, 05, 445-456.	1.4	0
51	TAPAS: Towards Automated Processing and Analysis of multi-dimensional bioimage data. <i>F1000Research</i> , 2020, 9, 1278.	1.6	0