Axel Mosig

List of Publications by Year in descending order

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52	1,590	23	38
papers	citations	h-index	g-index
62	62	62	2332
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Evolution of Vault RNAs. Molecular Biology and Evolution, 2009, 26, 1975-1991.	3.5	130
2	Label-free imaging of drug distribution and metabolism in colon cancer cells by Raman microscopy. Analyst, The, 2014, 139, 1155.	1.7	126
3	Immunohistochemistry, histopathology and infrared spectral histopathology of colon cancer tissue sections. Journal of Biophotonics, 2013, 6, 88-100.	1.1	101
4	Structure and Function of the Smallest Vertebrate Telomerase RNA from Teleost Fish. Journal of Biological Chemistry, 2008, 283, 2049-2059.	1.6	78
5	The common ancestral core of vertebrate and fungal telomerase RNAs. Nucleic Acids Research, 2013, 41, 450-462.	6.5	70
6	Evolutionary patterns of non-coding RNAs. Theory in Biosciences, 2005, 123, 301-369.	0.6	64
7	Characterization of statistical features for plant microRNA prediction. BMC Genomics, 2011, 12, 108.	1.2	63
8	Label-free classification of colon cancer grading using infrared spectral histopathology. Faraday Discussions, 2016, 187, 105-118.	1.6	56
9	Omnisphero: a high-content image analysis (HCA) approach for phenotypic developmental neurotoxicity (DNT) screenings of organoid neurosphere cultures in vitro. Archives of Toxicology, 2017, 91, 2017-2028.	1.9	56
10	Evolution of the vertebrate Y RNA cluster. Theory in Biosciences, 2007, 126, 9-14.	0.6	54
11	Spectral histopathology of colon cancer tissue sections by Raman imaging with 532 nm excitation provides label free annotation of lymphocytes, erythrocytes and proliferating nuclei of cancer cells. Analyst, The, 2013, 138, 4035.	1.7	51
12	Spatial and molecular resolution of diffuse malignant mesothelioma heterogeneity by integrating label-free FTIR imaging, laser capture microdissection and proteomics. Scientific Reports, 2017, 7, 44829.	1.6	49
13	RNAs everywhere: genome-wide annotation of structured RNAs. Journal of Experimental Zoology Part B: Molecular and Developmental Evolution, 2007, 308B, 1-25.	0.6	46
14	Arthropod 7SK RNA. Molecular Biology and Evolution, 2008, 25, 1923-1930.	3.5	45
15	Automated Identification of Subcellular Organelles by Coherent Anti-Stokes Raman Scattering. Biophysical Journal, 2014, 106, 1910-1920.	0.2	43
16	Circulation times of prostate cancer and hepatocellular carcinoma cells by in vivo flow cytometry. Cytometry Part A: the Journal of the International Society for Analytical Cytology, 2011, 79A, 848-854.	1,1	41
17	Fast and Noninvasive Diagnosis of Cervical Cancer by Coherent Anti-Stokes Raman Scattering. Analytical Chemistry, 2019, 91, 13900-13906.	3.2	39
18	Noninvasive Diagnosis of High-Grade Urothelial Carcinoma in Urine by Raman Spectral Imaging. Analytical Chemistry, 2017, 89, 6893-6899.	3.2	38

#	Article	IF	CITATIONS
19	Invertebrate 7SK snRNAs. Journal of Molecular Evolution, 2008, 66, 107-115.	0.8	37
20	Establishment of an a priori protocol for the implementation and interpretation of an inâ€vitro testing battery for the assessment of developmental neurotoxicity. EFSA Supporting Publications, 2020, 17, 1938E.	0.3	36
21	Tracking cells in Life Cell Imaging videos using topological alignments. Algorithms for Molecular Biology, 2009, 4, 10.	0.3	33
22	Approximately matching polygonal curves with respect to the Fr \tilde{A} © chet distance. Computational Geometry: Theory and Applications, 2005, 30, 113-127.	0.3	29
23	Hierarchical deep convolutional neural networks combine spectral and spatial information for highly accurate Ramanâ€microscopyâ€based cytopathology. Journal of Biophotonics, 2018, 11, e201800022.	1.1	29
24	Colocalization of fluorescence and Raman microscopic images for the identification of subcellular compartments: a validation study. Analyst, The, 2015, 140, 2360-2368.	1.7	24
25	Lead-seq: transcriptome-wide structure probing in vivo using lead(II) ions. Nucleic Acids Research, 2020, 48, e71-e71.	6.5	24
26	Fragrep: An Efficient Search Tool for Fragmented Patterns in Genomic Sequences. Genomics, Proteomics and Bioinformatics, 2006, 4, 56-60.	3.0	21
27	Prevalent and distinct spliceosomal 3′-end processing mechanisms for fungal telomerase RNA. Nature Communications, 2015, 6, 6105.	5.8	21
28	Deep representation learning for domain adaptable classification of infrared spectral imaging data. Bioinformatics, 2020, 36, 287-294.	1.8	19
29	U7 snRNAs: A Computational Survey. Genomics, Proteomics and Bioinformatics, 2007, 5, 187-195.	3.0	17
30	Customized strategies for discovering distant ncRNA homologs. Briefings in Functional Genomics & Proteomics, 2009, 8, 451-460.	3.8	17
31	<i>Ct3d:</i> tracking microglia motility in 3D using a novel cosegmentation approach. Bioinformatics, 2011, 27, 564-571.	1.8	17
32	Homology Search with Fragmented Nucleic Acid Sequence Patterns. Lecture Notes in Computer Science, 2007, , 335-345.	1.0	13
33	A computational evaluation of over-representation of regulatory motifs in the promoter regions of differentially expressed genes. BMC Bioinformatics, 2010, 11 , 267 .	1.2	11
34	A Propagated Skeleton Approach to High Throughput Screening of Neurite Outgrowth for In Vitro Parkinson's Disease Modelling. Cells, 2021, 10, 931.	1.8	10
35	Fully automated registration of vibrational microspectroscopic images in histologically stained tissue sections. BMC Bioinformatics, 2015, 16, 396.	1.2	9
36	Integrating spatial, morphological, and textural information for improved cell type differentiation using Raman microscopy. Journal of Chemometrics, 2018, 32, e2973.	0.7	9

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37	Similarity maps and hierarchical clustering for annotating FT-IR spectral images. BMC Bioinformatics, 2013, 14, 333.	1.2	8
38	An Image Skeletonizationâ€Based Tool for Pollen Tube Morphology Analysis and Phenotyping ^F . Journal of Integrative Plant Biology, 2013, 55, 131-141.	4.1	8
39	Automatic counting and positioning of 5-bromo-2-deoxyuridine (BrdU) positive cells in cortical layers of rat brain slices. NeuroToxicology, 2014, 43, 127-133.	1.4	8
40	Grayscale representation of infrared microscopy images by extended multiplicative signal correction for registration with histological images. Journal of Biophotonics, 2020, 13, e201960223.	1.1	8
41	Reliable Identification and Quantification of Neural Cells in Microscopic Images of Neurospheres. Cytometry Part A: the Journal of the International Society for Analytical Cytology, 2021, , .	1.1	6
42	Signaling pathways of heat- and hypersalinity-induced polyp bailout in Pocillopora acuta. Coral Reefs, 2021, 40, 1713-1728.	0.9	5
43	Cell Counting for In Vivo Flow Cytometer Signals Using Wavelet-Based Dynamic Peak Picking. , 2009, , .		3
44	A representation learning approach for recovering scatterâ€corrected spectra from Fourierâ€transform infrared spectra of tissue samples. Journal of Biophotonics, 2021, 14, e202000385.	1.1	3
45	Shape decomposition algorithms for laser capture microdissection. Algorithms for Molecular Biology, 2021, 16, 15.	0.3	3
46	Regulatory Signals in Genomic Sequences. , 2007, , 189-216.		1
47	Association of high microsatellite instability (MSI-H) with a high immunoscore (IS) compared to PD-L1 expression and increased survival in patients (pts) with metastatic colorectal cancer (mCRC) treated with oxaliplatin (Ox) and fluoropyrimidine (FP): A pooled analysis of the AIO KRK 0207 and RO91 trials Journal of Clinical Oncology, 2017, 35, 3595-3595.	0.8	1
48	Dynamic Programming Algorithms for Efficiently Computing Cosegmentations between Biological Images. Lecture Notes in Computer Science, 2011, , 339-350.	1.0	1
49	Inverse folding based pre-training for the reliable identification of intrinsic transcription terminators. PLoS Computational Biology, 2022, 18, e1010240.	1.5	1
50	Discovering cis-regulatory modules by optimizing barbecues. Discrete Applied Mathematics, 2009, 157, 2458-2468.	0.5	0
51	Identifying minimally redundant wavenumbers for vibrational microspectroscopic image analysis. , 2014, , .		0
52	Letter to the Editor regarding the recent contribution by Roussel etÂal., SARS-CoV-2: Fear versus data. International Journal of Antimicrobial Agents, 2020, 56, 106074.	1.1	0