

Valmir Carneiro Barbosa

List of Publications by Year in descending order

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134
papers

2,486
citations

304743

22
h-index

223800

46
g-index

137
all docs

137
docs citations

137
times ranked

4114
citing authors

#	ARTICLE	IF	CITATIONS
1	Analyzing marginal cases in differential shotgun proteomics. <i>Bioinformatics</i> , 2011, 27, 275-276.	4.1	474
2	Integrated analysis of shotgun proteomic data with PatternLab for proteomics 4.0. <i>Nature Protocols</i> , 2016, 11, 102-117.	12.0	257
3	PatternLab for proteomics: a tool for differential shotgun proteomics. <i>BMC Bioinformatics</i> , 2008, 9, 316.	2.6	127
4	Search engine processor: Filtering and organizing peptide spectrum matches. <i>Proteomics</i> , 2012, 12, 944-949.	2.2	107
5	XDIA: improving on the label-free data-independent analysis. <i>Bioinformatics</i> , 2010, 26, 847-848.	4.1	104
6	Concurrency in heavily loaded neighborhood-constrained systems. <i>ACM Transactions on Programming Languages and Systems</i> , 1989, 11, 562-584.	2.1	86
7	Identifying differences in protein expression levels by spectral counting and feature selection. <i>Genetics and Molecular Research</i> , 2008, 7, 342-356.	0.2	78
8	Improving the TFold test for differential shotgun proteomics. <i>Bioinformatics</i> , 2012, 28, 1652-1654.	4.1	73
9	SIM-XL: A powerful and user-friendly tool for peptide cross-linking analysis. <i>Journal of Proteomics</i> , 2015, 129, 51-55.	2.4	73
10	YADA: a tool for taking the most out of high-resolution spectra. <i>Bioinformatics</i> , 2009, 25, 2734-2736.	4.1	67
11	On best practices in the development of bioinformatics software. <i>Frontiers in Genetics</i> , 2014, 5, 199.	2.3	53
12	Characterization of homodimer interfaces with cross-linking mass spectrometry and isotopically labeled proteins. <i>Nature Protocols</i> , 2018, 13, 431-458.	12.0	47
13	PatternLab: From Mass Spectra to Label-Free Differential Shotgun Proteomics. <i>Current Protocols in Bioinformatics</i> , 2012, 40, Unit13.19.	25.8	39
14	GO Explorer: A gene-ontology tool to aid in the interpretation of shotgun proteomics data. <i>Proteome Science</i> , 2009, 7, 6.	1.7	35
15	Can the false-discovery rate be misleading?. <i>Proteomics</i> , 2011, 11, 4105-4108.	2.2	34
16	V-like Formations in Flocks of Artificial Birds. <i>Artificial Life</i> , 2008, 14, 179-188.	1.3	33
17	PepExplorer: A Similarity-driven Tool for Analyzing de Novo Sequencing Results. <i>Molecular and Cellular Proteomics</i> , 2014, 13, 2480-2489.	3.8	33
18	Probabilistic Heuristics for Disseminating Information in Networks. <i>IEEE/ACM Transactions on Networking</i> , 2007, 15, 425-435.	3.8	29

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19	A distributed algorithm to find k-dominating sets. <i>Discrete Applied Mathematics</i> , 2004, 141, 243-253.	0.9	27
20	Simple, efficient and thorough shotgun proteomic analysis with PatternLab V. <i>Nature Protocols</i> , 2022, 17, 1553-1578.	12.0	26
21	Analyzing Shotgun Proteomic Data with PatternLab for Proteomics. <i>Current Protocols in Bioinformatics</i> , 2010, 30, Unit 13.13.1-15.	25.8	24
22	Are Gastric Cancer Resection Margin Proteomic Profiles More Similar to Those from Controls or Tumors?. <i>Journal of Proteome Research</i> , 2012, 11, 5836-5842.	3.7	24
23	Dynamic proteomic overview of glioblastoma cells (A172) exposed to perillyl alcohol. <i>Journal of Proteomics</i> , 2010, 73, 1018-1027.	2.4	23
24	Early appraisal of the fixation probability in directed networks. <i>Physical Review E</i> , 2010, 82, 046114.	2.1	22
25	A distributed implementation of simulated annealing. <i>Journal of Parallel and Distributed Computing</i> , 1989, 6, 411-434.	4.1	21
26	On the distributed parallel simulation of Hopfield's neural networks. <i>Software - Practice and Experience</i> , 1990, 20, 967-983.	3.6	21
27	Generating all the acyclic orientations of an undirected graph. <i>Information Processing Letters</i> , 1999, 72, 71-74.	0.6	21
28	A multi-protease, multi-dissociation, bottom-up-to-top-down proteomic view of the <i>Loxosceles intermedia</i> venom. <i>Scientific Data</i> , 2017, 4, 170090.	5.3	21
29	Finding approximate palindromes in strings. <i>Pattern Recognition</i> , 2002, 35, 2581-2591.	8.1	20
30	Two Novel Evolutionary Formulations of the Graph Coloring Problem. <i>Journal of Combinatorial Optimization</i> , 2004, 8, 41-63.	1.3	20
31	Effectively addressing complex proteomic search spaces with peptide spectrum matching. <i>Bioinformatics</i> , 2013, 29, 1343-1344.	4.1	20
32	An algorithm for clock synchronization with the gradient property in sensor networks. <i>Journal of Parallel and Distributed Computing</i> , 2009, 69, 261-265.	4.1	19
33	Sharing Resources at Nonuniform Access Rates. <i>Theory of Computing Systems</i> , 2000, 34, 13-26.	1.1	18
34	Exploring the Proteomic Landscape of a Gastric Cancer Biopsy with the Shotgun Imaging Analyzer. <i>Journal of Proteome Research</i> , 2014, 13, 314-320.	3.7	18
35	A Novel Evolutionary Formulation of the Maximum Independent Set Problem. <i>Journal of Combinatorial Optimization</i> , 2004, 8, 419-437.	1.3	17
36	Dissemination strategy for immunizing scale-free networks. <i>Physical Review E</i> , 2006, 74, 056105.	2.1	15

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37	Charge Prediction Machine: Tool for Inferring Precursor Charge States of Electron Transfer Dissociation Tandem Mass Spectra. <i>Analytical Chemistry</i> , 2009, 81, 1996-2003.	6.5	15
38	On the phase transitions of graph coloring and independent sets. <i>Physica A: Statistical Mechanics and Its Applications</i> , 2004, 343, 401-423.	2.6	14
39	Distributed Breakpoint Detection in Message-Passing Programs. <i>Journal of Parallel and Distributed Computing</i> , 1996, 39, 153-167.	4.1	13
40	Learning in the combinatorial neural model. <i>IEEE Transactions on Neural Networks</i> , 1998, 9, 831-847.	4.2	13
41	A Bayesian-Network Approach to Lexical Disambiguation. <i>Cognitive Science</i> , 1993, 17, 257-283.	1.7	11
42	Pinpointing differentially expressed domains in complex protein mixtures with the cloud service of PatternLab for Proteomics. <i>Journal of Proteomics</i> , 2013, 89, 179-182.	2.4	11
43	Differential proteomic comparison of breast cancer secretome using a quantitative paired analysis workflow. <i>BMC Cancer</i> , 2019, 19, 365.	2.6	11
44	Strategies for the prevention of communication deadlocks in distributed parallel programs. <i>IEEE Transactions on Software Engineering</i> , 1990, 16, 1311-1316.	5.6	10
45	Quasispecies dynamics with network constraints. <i>Journal of Theoretical Biology</i> , 2012, 312, 114-119.	1.7	10
46	Directed cycles and related structures in random graphs: "Static properties. <i>Physica A: Statistical Mechanics and Its Applications</i> , 2003, 321, 381-397.	2.6	9
47	Scheduling links for heavy traffic on interfering routes in wireless mesh networks. <i>Computer Networks</i> , 2012, 56, 1584-1598.	5.1	9
48	A scoring model for phosphopeptide site localization and its impact on the question of whether to use MSA. <i>Journal of Proteomics</i> , 2015, 129, 42-50.	2.4	9
49	Learning Logic Programs with Neural Networks. <i>Lecture Notes in Computer Science</i> , 2001, , 15-26.	1.3	9
50	A graph model for the evolution of specificity in humoral immunity. <i>Journal of Theoretical Biology</i> , 2004, 229, 311-325.	1.7	8
51	Handling flash-crowd events to improve the performance of web applications. , 2015, , .		8
52	Mixed-Data Acquisition: Next-Generation Quantitative Proteomics Data Acquisition. <i>Journal of Proteomics</i> , 2020, 222, 103803.	2.4	8
53	The Combinatorics of Resource Sharing. <i>Applied Optimization</i> , 2002, , 27-52.	0.4	8
54	A neural system for deforestation monitoring on Landsat images of the Amazon Region. <i>International Journal of Approximate Reasoning</i> , 1994, 11, 321-359.	3.3	7

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55	Acyclic Orientations with Path Constraints. <i>RAIRO - Operations Research</i> , 2008, 42, 455-467.	1.8	7
56	DiagnoProt: a tool for discovery of new molecules by mass spectrometry. <i>Bioinformatics</i> , 2017, 33, 1883-1885.	4.1	7
57	Fast linear system solution by neural networks. <i>Operations Research Letters</i> , 1992, 11, 141-145.	0.7	6
58	Emergence of scale-free networks from local connectivity and communication trade-offs. <i>Physical Review E</i> , 2006, 74, 016113.	2.1	6
59	Network growth for enhanced natural selection. <i>Physical Review E</i> , 2009, 80, 026115.	2.1	6
60	Structured construction and simulation of nondeterministic stochastic activity networks. <i>European Journal of Operational Research</i> , 2009, 198, 266-274.	5.7	6
61	The Interleaved Multichromatic Number of a Graph. <i>Annals of Combinatorics</i> , 2002, 6, 249-256.	0.6	5
62	Local heuristics and the emergence of spanning subgraphs in complex networks. <i>Theoretical Computer Science</i> , 2006, 355, 80-95.	0.9	5
63	Descents and nodal load in scale-free networks. <i>Physical Review E</i> , 2008, 77, 046111.	2.1	5
64	Network algorithmics and the emergence of the cortical synaptic-weight distribution. <i>Physical Review E</i> , 2010, 81, 021916.	2.1	5
65	Scheduling wireless links by vertex multicoloring in the physical interference model. <i>Computer Networks</i> , 2016, 99, 125-133.	5.1	5
66	Deadlock models in distributed computation. , 2016, , .		5
67	An Occam-based evaluation of a parallel version of simulated annealing. <i>Microprocessing and Microprogramming</i> , 1990, 30, 85-92.	0.2	4
68	On reducing the complexity of matrix clocks. <i>Parallel Computing</i> , 2003, 29, 895-905.	2.1	4
69	Finding routes in anonymous sensor networks. <i>Information Processing Letters</i> , 2006, 98, 139-144.	0.6	4
70	Exploiting the distribution of distances between nodes to efficiently solve the localization problem in wireless sensor networks. , 2010, , .		4
71	Network algorithmics and the emergence of information integration in cortical models. <i>Physical Review E</i> , 2011, 84, 011904.	2.1	4
72	Local heuristic for the refinement of multi-path routing in wireless mesh networks. <i>Computer Networks</i> , 2013, 57, 273-285.	5.1	4

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73	Using PepExplorer to Filter and Organize <i>De Novo</i> Peptide Sequencing Results. Current Protocols in Bioinformatics, 2015, 51, 13.27.1-13.27.9.	25.8	4
74	Early detection of epilepsy seizures based on a weightless neural network. , 2015, 2015, 4470-4.		4
75	Information Integration from Distributed Threshold-Based Interactions. Complexity, 2017, 2017, 1-14.	1.6	4
76	Top-Down Garbage Collector: a tool for selecting high-quality top-down proteomics mass spectra. Bioinformatics, 2019, 35, 3489-3490.	4.1	4
77	Using SIM-XL to identify and annotate cross-linked peptides analyzed by mass spectrometry. Protocol Exchange, 0, , .	0.3	4
78	Specification of a communication virtual processor for parallel processing systems. Microprocessing and Microprogramming, 1988, 24, 511-518.	0.2	3
79	Feasible directions linear programming by neural networks. , 1990, , .		3
80	Monitoring the deforestation of the Amazon region with neural networks. , 0, , .		3
81	A priority dynamics for generalized drinking philosophers. Information Processing Letters, 2001, 79, 189-195.	0.6	3
82	Partially ordered distributed computations on asynchronous point-to-point networks. Parallel Computing, 2009, 35, 12-28.	2.1	3
83	Towards a Hybrid Model of First-Order Theory Refinement. Lecture Notes in Computer Science, 2000, , 92-106.	1.3	3
84	Quasispecies dynamics on a network of interacting genotypes and idiotypes: formulation of the model. Journal of Statistical Mechanics: Theory and Experiment, 2015, 2015, P01022.	2.3	3
85	From distributed algorithms to OCCAM programs by successive refinements. Journal of Systems and Software, 1994, 26, 257-272.	4.5	2
86	Defeasible time-stepping. Parallel Computing, 1999, 25, 461-489.	2.1	2
87	Cell-centric heuristics for the classification of cellular automata. Parallel Computing, 2006, 32, 44-66.	2.1	2
88	Scheduling Cyclic Task Graphs with SCC-Map. , 2012, , .		2
89	The network structure of mathematical knowledge according to the Wikipedia, MathWorld, and DLMF online libraries. Network Science, 2014, 2, 367-386.	1.0	2
90	Quasispecies dynamics on a network of interacting genotypes and idiotypes: applications to autoimmunity and immunodeficiency. Journal of Statistical Mechanics: Theory and Experiment, 2016, 2016, 063501.	2.3	2

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91	Adaptive event sensing in networks of autonomous mobile agents. Journal of Network and Computer Applications, 2016, 71, 118-129.	9.1	2
92	Coevolution of the mitotic and meiotic modes of eukaryotic cellular division. Physical Review E, 2018, 98, .	2.1	2
93	Counting trees with random walks. , 2019, 37, 96-102.		2
94	Scheduling Wireless Links in the Physical Interference Model by Fractional Edge Coloring. IEEE Wireless Communications Letters, 2020, 9, 528-532.	5.0	2
95	MPH " A Hybrid Parallel Machine. Microprocessing and Microprogramming, 1989, 25, 229-232.	0.2	1
96	Blocking versus nonblocking interprocess communication: a note on the effect on concurrency. Information Processing Letters, 1990, 36, 171-175.	0.6	1
97	An integrated software environment for large-scale Occam programming. Microprocessing and Microprogramming, 1991, 32, 393-400.	0.2	1
98	An algorithm for FIFO message delivery among migrating tasks. Information Processing Letters, 1995, 53, 261-267.	0.6	1
99	Instruction usage and the memory gap problem. , 0, , .		1
100	Directed cycles and related structures in random graphs: "Dynamic properties. Physica A: Statistical Mechanics and Its Applications, 2004, 334, 566-582.	2.6	1
101	A graph model for the evolution of specificity in humoral immunity. Journal of Theoretical Biology, 2004, 229, 311-311.	1.7	1
102	Emergence of scale-free behavior in networks from limited-horizon linking and cost trade-offs. Physica A: Statistical Mechanics and Its Applications, 2008, 387, 1016-1024.	2.6	1
103	Network Conduciveness with Application to the Graph-Coloring and Independent-Set Optimization Transitions. PLoS ONE, 2010, 5, e11232.	2.5	1
104	The predecessor-existence problem for k-reversible processes. Theoretical Computer Science, 2015, 562, 406-418.	0.9	1
105	Power-law decay of the degree-sequence probabilities of multiple random graphs with application to graph isomorphism. ESAIM - Probability and Statistics, 2017, 21, 235-250.	0.5	1
106	A computational study of f-reversible processes on graphs. Discrete Applied Mathematics, 2018, 245, 77-93.	0.9	1
107	A quantitation module for isotope-labeled peptides integrated into PatternLab for proteomics. Journal of Proteomics, 2019, 202, 103371.	2.4	1
108	Local Symmetry in Random Graphs. IEEE Transactions on Network Science and Engineering, 2020, 7, 1913-1924.	6.4	1

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109	Interspecies evolutionary dynamics mediated by public goods in bacterial quorum sensing. <i>Physical Review E</i> , 2021, 103, 012403.	2.1	1
110	Minimal Chordal Sense of Direction and Circulant Graphs. <i>Lecture Notes in Computer Science</i> , 2006, , 670-680.	1.3	1
111	Approximate Conditional Distributions of Distances between Nodes in a Two-Dimensional Sensor Network. <i>Lecture Notes in Computer Science</i> , 2009, , 324-338.	1.3	1
112	Micro-instruction placement by simulated annealing. <i>Microprocessing and Microprogramming</i> , 1991, 32, 23-28.	0.2	0
113	Learning in analog Hopfield networks. , 0, , .		0
114	A string-matching algorithm for the CREW PRAM. <i>Information Processing Letters</i> , 1993, 47, 257-259.	0.6	0
115	A BUU Code for Parallel Computers. <i>International Journal of Modern Physics C</i> , 1998, 09, 573-583.	1.7	0
116	A distributed algorithm for k-dominating sets. <i>Electronic Notes in Discrete Mathematics</i> , 2001, 7, 130-133.	0.4	0
117	Generating all the cubic graphs that have a 6-cycle double cover. <i>Electronic Notes in Discrete Mathematics</i> , 2005, 19, 87-93.	0.4	0
118	Two-dimensional cellular automata and the analysis of correlated time series. <i>Pattern Recognition Letters</i> , 2006, 27, 1353-1360.	4.2	0
119	Modeling the Input History of Programs for Improved Instruction-Memory Performance. <i>Computer Journal</i> , 2006, 49, 744-761.	2.4	0
120	Reachability and recoverability of sink nodes in growing acyclic directed networks. <i>Physica A: Statistical Mechanics and Its Applications</i> , 2008, 387, 685-693.	2.6	0
121	Optimization of supply diversity for the self-assembly of simple objects in two and three dimensions. <i>Natural Computing</i> , 2011, 10, 551-581.	3.0	0
122	The Conduciveness of CA-Rule Graphs. <i>Artificial Life</i> , 2013, 19, 255-266.	1.3	0
123	Cooperation in Cognitive Radio Networks. , 2014, , .		0
124	Revisiting deadlock prevention: A probabilistic approach. <i>Networks</i> , 2014, 63, 203-210.	2.7	0
125	Further insights into the interareal connectivity of a cortical network. <i>Network Science</i> , 2015, 3, 526-550.	1.0	0
126	A note on counting independent terms in asymptotic expressions of computational complexity. <i>Optimization Letters</i> , 2017, 11, 1757-1765.	1.6	0

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127	Information-theoretic signatures of biodiversity in the barcoding gene. Journal of Theoretical Biology, 2018, 451, 111-116.	1.7	0
128	Sketching Data Structures for Massive Graph Problems. Lecture Notes in Computer Science, 2019, , 57-67.	1.3	0
129	Leveraging the partition selection bias to achieve a high-quality clustering of mass spectra. Journal of Proteomics, 2021, 245, 104282.	2.4	0
130	Multiple Sequence Alignment Based on Set Covers. Lecture Notes in Computer Science, 2006, , 127-137.	1.3	0
131	A Methodology for Determining Amino-Acid Substitution Matrices from Set Covers. Lecture Notes in Computer Science, 2006, , 138-148.	1.3	0
132	EVOLVED PREAMBLES FOR MAX-SAT HEURISTICS. , 2011, , .		0
133	Error-Prone Cellular Automata as Metaphors of Immunity as Computation. Complex Systems, 2015, 24, 93-112.	0.3	0
134	Integrated Optimization of Heterogeneous-Network Management and the Elusive Role of Macrocells. IEEE Access, 2021, 9, 149552-149559.	4.2	0