Sebastian Daberdaku

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7441282/publications.pdf

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		1477746	1199166
15	184	6	12
papers	citations	h-index	g-index
19	19	19	205
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Hierarchical representation for PPI sites prediction. BMC Bioinformatics, 2022, 23, 96.	1.2	5
2	Predicting functional impairment trajectories in amyotrophic lateral sclerosis: a probabilistic, multifactorial model of disease progression. Journal of Neurology, 2022, 269, 3858-3878.	1.8	7
3	Exploiting mutual information for the imputation of static and dynamic mixed-type clinical data with an adaptive k-nearest neighbours approach. BMC Medical Informatics and Decision Making, 2020, 20, 174.	1.5	12
4	A Combined Interpolation and Weighted K-Nearest Neighbours Approach for the Imputation of Longitudinal ICU Laboratory Data. Journal of Healthcare Informatics Research, 2020, 4, 174-188.	5. 3	15
5	Structure-Based Antibody Paratope Prediction with 3D Zernike Descriptors and SVM. Lecture Notes in Computer Science, 2020, , 27-49.	1.0	3
6	Hierarchical Representation and Graph Convolutional Networks for the Prediction of Protein–Protein Interaction Sites. Lecture Notes in Computer Science, 2020, , 409-420.	1.0	5
7	Accelerating the computation of triangulated molecular surfaces with OpenMP. Journal of Supercomputing, 2019, 75, 3426-3470.	2.4	2
8	Interpolation and K-Nearest Neighbours Combined Imputation for Longitudinal ICU Laboratory Data. , 2019, , .		1
9	Antibody interface prediction with 3D Zernike descriptors and SVM. Bioinformatics, 2019, 35, 1870-1876.	1.8	65
10	Computing voxelised representations of macromolecular surfaces. International Journal of High Performance Computing Applications, 2018, 32, 407-432.	2.4	10
11	Fast Computation of High-resolution Solvent Excluded Protein Surface with OpenMP. , 2018, , .		1
12	Parallel Computation of Voxelised Protein Surfaces with OpenMP., 2018,,.		4
13	Exploring the potential of 3D Zernike descriptors and SVM for protein–protein interface prediction. BMC Bioinformatics, 2018, 19, 35.	1.2	42
14	Computing Discrete Fine-Grained Representations of Protein Surfaces. Lecture Notes in Computer Science, 2016, , 180-195.	1.0	7
15	Parallel Computation of Voxelized Macromolecular Surfaces by Spatial Slicing. , 2015, , .		3