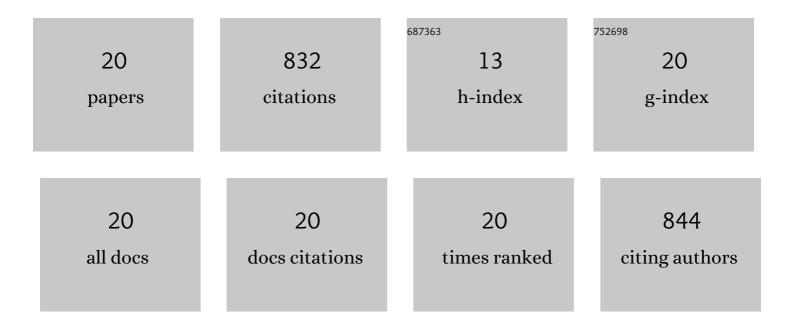
Vesna Memisevic

List of Publications by Year in descending order

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VESNA MEMISEVIC

#	Article	IF	CITATIONS
1	Mechanisms of action of Coxiella burnetii effectors inferred from host-pathogen protein interactions. PLoS ONE, 2017, 12, e0188071.	2.5	12
2	DBSecSys 2.0: a database of Burkholderia mallei and Burkholderia pseudomallei secretion systems. BMC Bioinformatics, 2016, 17, 387.	2.6	4
3	Using host-pathogen protein interactions to identify and characterize Francisella tularensis virulence factors. BMC Genomics, 2015, 16, 1106.	2.8	33
4	A reverse-phase protein microarray-based screen identifies host signaling dynamics upon Burkholderia spp. infection. Frontiers in Microbiology, 2015, 6, 683.	3.5	11
5	Mining Host-Pathogen Protein Interactions to Characterize Burkholderia mallei Infectivity Mechanisms. PLoS Computational Biology, 2015, 11, e1004088.	3.2	34
6	DBSecSys: a database of Burkholderia malleisecretion systems. BMC Bioinformatics, 2014, 15, 244.	2.6	9
7	Reconstituting protein interaction networks using parameter-dependent domain-domain interactions. BMC Bioinformatics, 2013, 14, 154.	2.6	17
8	Novel Burkholderia mallei Virulence Factors Linked to Specific Host-Pathogen Protein Interactions. Molecular and Cellular Proteomics, 2013, 12, 3036-3051.	3.8	38
9	C-GRAAL: Common-neighbors-based global GRAph ALignment of biological networks. Integrative Biology (United Kingdom), 2012, 4, 734.	1.3	79
10	Tutorial on biological networks. Wiley Interdisciplinary Reviews: Data Mining and Knowledge Discovery, 2012, 2, 298-325.	6.8	11
11	Dominating Biological Networks. PLoS ONE, 2011, 6, e23016.	2.5	85
12	Categorizing Biases in High-Confidence High-Throughput Protein-Protein Interaction Data Sets. Molecular and Cellular Proteomics, 2011, 10, M111.012500.	3.8	25
13	Protein interaction network topology uncovers melanogenesis regulatory network components within functional genomics datasets. BMC Systems Biology, 2010, 4, 84.	3.0	32
14	Complementarity of network and sequence information in homologous proteins. Journal of Integrative Bioinformatics, 2010, 7, 275-289.	1.5	20
15	An integrative approach to modeling biological networks. Journal of Integrative Bioinformatics, 2010, 7, .	1.5	14
16	Systems-level cancer gene identification from protein interaction network topology applied to melanogenesis-related functional genomics data. Journal of the Royal Society Interface, 2010, 7, 423-437.	3.4	95
17	Topological network alignment uncovers biological function and phylogeny. Journal of the Royal Society Interface, 2010, 7, 1341-1354.	3.4	281
18	An integrative approach to modeling biological networks. Journal of Integrative Bioinformatics, 2010, 7	1.5	8

#	Article	IF	CITATIONS
19	Complementarity of network and sequence information in homologous proteins. Journal of Integrative Bioinformatics, 2010, 7, .	1.5	17
20	Topological network alignment uncovers biological function and phylogeny. Nature Precedings, 2009, , .	0.1	7