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List of Publications by Year in descending order

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1040056 1199594 12 473 9 12 citations h-index g-index papers 16 16 16 904 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Combining Functional and Structural Genomics to Sample the Essential Burkholderia Structome. PLoS ONE, 2013, 8, e53851.	2.5	113
2	Increasing the structural coverage of tuberculosis drug targets. Tuberculosis, 2015, 95, 142-148.	1.9	103
3	Calmodulin-like proteins localized to the conoid regulate motility and cell invasion by Toxoplasma gondii. PLoS Pathogens, 2017, 13, e1006379.	4.7	89
4	Structural genomics of infectious disease drug targets: the SSGCID. Acta Crystallographica Section F: Structural Biology Communications, 2011, 67, 979-984.	0.7	55
5	A doublecortin-domain protein of Toxoplasma and its orthologues bind to and modify the structure and organization of tubulin polymers. BMC Molecular and Cell Biology, 2020, 21, 8.	2.0	32
6	Enzymatic and Structural Characterization of the <i>Naegleria fowleri</i> Glucokinase. Antimicrobial Agents and Chemotherapy, 2019, 63, .	3.2	21
7	The BID Domain of Type IV Secretion Substrates Forms a Conserved Four-Helix Bundle Topped with a Hook. Structure, 2017, 25, 203-211.	3.3	15
8	Chromatin-Associated Protein Complexes Link DNA Base J and Transcription Termination in $\mbox{\ensuremath{\mbox{\sc i}}}\mbox{\ensuremath{\mbox{\sc leish}}}\mbox{\ensuremath{\mbox{\sc leish}}}\mbox{\ensuremath}}\mbox{\ensuremath{\mbox{\sc leish}}}\mbox{\ensuremath}}\mbox{\ensuremath}}\mbox{\ensuremath}}\mbox{\ensuremath}\mbox{\ensuremath}}\mbox{\ensuremath}\mbox{\ensuremath}}\mbox{\ensuremath}\mbox{\ensuremath}}\mbox{\ensuremath}\mbox{\ensuremath}}\mbox{\ensuremath}\mbox{\ensuremath}}\mbox{\ensuremath}\mbox{\ensuremath}}$	2.9	12
9	Naegleria fowleri: Protein structures to facilitate drug discovery for the deadly, pathogenic free-living amoeba. PLoS ONE, 2021, 16, e0241738.	2.5	12
10	Evolutionary Diversification of Host-Targeted Bartonella Effectors Proteins Derived from a Conserved FicTA Toxin-Antitoxin Module. Microorganisms, 2021, 9, 1645.	3.6	5
11	Toward a structome of <scp><i>Acinetobacter baumannii</i></scp> drug targets. Protein Science, 2020, 29, 789-802.	7.6	4
12	The domain architecture of the protozoan protein J-DNA–binding protein 1 suggests synergy between base J DNA binding and thymidine hydroxylase activity. Journal of Biological Chemistry, 2019, 294, 12815-12825.	3.4	3