

Philippe Youkharibache

List of Publications by Year in descending order

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Version: 2024-02-01

9
papers

492
citations

1478505

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h-index

1474206

9
g-index

15
all docs

15
docs citations

15
times ranked

770
citing authors

#	ARTICLE	IF	CITATIONS
1	iCn3D: From Web-Based 3D Viewer to Structural Analysis Tool in Batch Mode. <i>Frontiers in Molecular Biosciences</i> , 2022, 9, 831740.	3.5	26
2	Topological and Structural Plasticity of the Single Ig Fold and the Double Ig Fold Present in CD19. <i>Biomolecules</i> , 2021, 11, 1290.	4.0	5
3	iCn3D, a web-based 3D viewer for sharing 1D/2D/3D representations of biomolecular structures. <i>Bioinformatics</i> , 2020, 36, 131-135.	4.1	113
4	Pseudo-Symmetric Assembly of Protodomains as a Common Denominator in the Evolution of Polytopic Helical Membrane Proteins. <i>Journal of Molecular Evolution</i> , 2020, 88, 319-344.	1.8	5
5	Safety and feasibility of anti-CD19 CAR T cells with fully human binding domains in patients with B-cell lymphoma. <i>Nature Medicine</i> , 2020, 26, 270-280.	30.7	182
6	Protodomains: Symmetry-Related Supersecondary Structures in Proteins and Self-Complementarity. <i>Methods in Molecular Biology</i> , 2019, 1958, 187-219.	0.9	7
7	The Small $\hat{\beta}$ -Barrel Domain: A Survey-Based Structural Analysis. <i>Structure</i> , 2019, 27, 6-26.	3.3	51
8	Systematic Detection of Internal Symmetry in Proteins Using CE-Symm. <i>Journal of Molecular Biology</i> , 2014, 426, 2255-2268.	4.2	44
9	Sm/Lsm Genes Provide a Glimpse into the Early Evolution of the Spliceosome. <i>PLoS Computational Biology</i> , 2009, 5, e1000315.	3.2	46