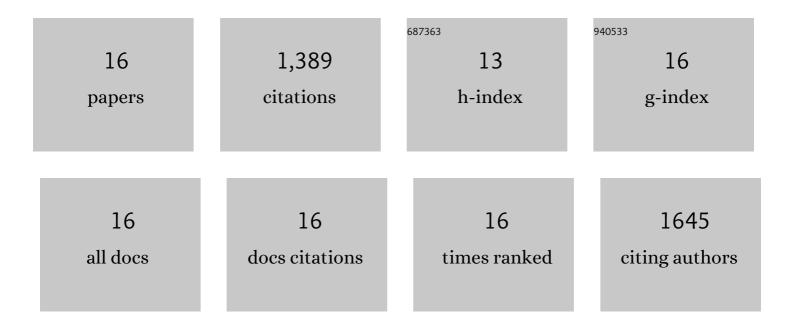
Joao M Dias

List of Publications by Year in descending order

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ΙΟΛΟ Μ ΡΙΛς

#	Article	IF	CITATIONS
1	A knottin scaffold directs the CXC-chemokine–binding specificity of tick evasins. Journal of Biological Chemistry, 2019, 294, 11199-11212.	3.4	22
2	Structural Basis for Differential Neutralization of Ebolaviruses. Viruses, 2012, 4, 447-470.	3.3	63
3	A shared structural solution for neutralizing ebolaviruses. Nature Structural and Molecular Biology, 2011, 18, 1424-1427.	8.2	113
4	Structural Basis of Chemokine Sequestration by a Tick Chemokine Binding Protein: The Crystal Structure of the Complex between Evasin-1 and CCL3. PLoS ONE, 2009, 4, e8514.	2.5	43
5	Ticks produce highly selective chemokine binding proteins with antiinflammatory activity. Journal of Experimental Medicine, 2008, 205, 2019-2031.	8.5	189
6	Molecular Cloning and Characterization of a Highly Selective Chemokine-binding Protein from the Tick Rhipicephalus sanguineus. Journal of Biological Chemistry, 2007, 282, 27250-27258.	3.4	109
7	Structural Basis for the Mechanism of Ca2+ Activation of the Di-Heme Cytochrome c Peroxidase from Pseudomonas nautica 617. Structure, 2004, 12, 961-973.	3.3	53
8	Cytochrome c Nitrite Reductase from Desulfovibrio desulfuricans ATCC 27774. Journal of Biological Chemistry, 2003, 278, 17455-17465.	3.4	98
9	Gene Sequence and the 1.8 Ã Crystal Structure of the Tungsten-Containing Formate Dehydrogenase from Desulfovibrio gigas. Structure, 2002, 10, 1261-1272.	3.3	161
10	Crystallization and preliminary X-ray diffraction analysis of two pH-dependent forms of a di-haem cytochromecperoxidase fromPseudomonas nautica. Acta Crystallographica Section D: Biological Crystallography, 2002, 58, 697-699.	2.5	6
11	Crystallization and preliminary X-ray analysis of a membrane-bound nitrite reductase from <i>Desulfovibrio desulfuricans</i> ATCC 27774. Acta Crystallographica Section D: Biological Crystallography, 2000, 56, 215-217.	2.5	6
12	Gene sequence and crystal structure of the aldehyde oxidoreductase from Desulfovibrio desulfuricans ATCC 27774. Journal of Molecular Biology, 2000, 297, 135-146.	4.2	64
13	Crystal structure of the first dissimilatory nitrate reductase at 1.9 Ã solved by MAD methods. Structure, 1999, 7, 65-79.	3.3	288
14	Crystal structure of acidic seminal fluid protein (aSFP) at 1.9 Ã resolution: a bovine polypeptide of the spermadhesin family. Journal of Molecular Biology, 1997, 274, 650-660.	4.2	42
15	The crystal structures of two spermadhesins reveal the CUB domain fold. Nature Structural Biology, 1997, 4, 783-788.	9.7	124
16	Crystallization and preliminary X-ray diffraction studies of aSFP, a bovine seminal plasma protein with a single CUB domain architecture. Protein Science, 1997, 6, 725-727.	7.6	8