

# Raffaele Raucci

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12155137/publications.pdf>

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9  
papers

150  
citations

1163117  
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1474206  
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g-index

9  
all docs

9  
docs citations

9  
times ranked

298  
citing authors

#	ARTICLE	IF	CITATIONS
1	Conformational analysis of the human chemokine receptor CXCR3. <i>Molecular Immunology</i> , 2017, 92, 76-86.	2.2	1
2	Structure-fluctuation-function relationships of seven pro-angiogenic isoforms of VEGFA, important mediators of tumorigenesis. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2015, 1854, 410-425.	2.3	14
3	An overview of the sequence features of N- and C-terminal segments of the human chemokine receptors. <i>Cytokine</i> , 2014, 70, 141-150.	3.2	14
4	Structure-function relationship and evolutionary history of the human selenoprotein M (SelM) found over-expressed in hepatocellular carcinoma. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2014, 1844, 447-456.	2.3	17
5	Peptides targeting chemokine receptor CXCR4: structural behavior and biological binding studies. <i>Journal of Peptide Science</i> , 2014, 20, 270-278.	1.4	8
6	Peptide Folding Problem: A Molecular Dynamics Study on Polyalanines Using Different Force Fields. <i>International Journal of Peptide Research and Therapeutics</i> , 2013, 19, 117-123.	1.9	13
7	Genealogy of an ancient protein family: the Sirtuins, a family of disordered members. <i>BMC Evolutionary Biology</i> , 2013, 13, 60.	3.2	47
8	Common structural interactions between the receptors CXCR3, CXCR4 and CXCR7 complexed with their natural ligands, CXCL11 and CXCL12, by a modeling approach. <i>Cytokine</i> , 2013, 64, 316-321.	3.2	22
9	The N-terminal Region of CXCL11 as Structural Template for CXCR3 Molecular Recognition: Synthesis, Conformational Analysis, and Binding Studies. <i>Chemical Biology and Drug Design</i> , 2012, 80, 254-265.	3.2	14