

# Volker Knecht

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/482348/publications.pdf>

Version: 2024-02-01

12  
papers

107  
citations

1307594

7  
h-index

1372567

10  
g-index

12  
all docs

12  
docs citations

12  
times ranked

175  
citing authors

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | Long-Range Conformational Response of a PDZ Domain to Ligand Binding and Release: A Molecular Dynamics Study. <i>Journal of Chemical Theory and Computation</i> , 2016, 12, 870-878.  | 5.3 | 23        |
| 2  | Conformational Diversity of the Fibrillogenic Fusion Peptide B18 in Different Environments from Molecular Dynamics Simulations. <i>Journal of Physical Chemistry B</i> , 2007, 111, 4161-4170.                              | 2.6 | 16        |
| 3  | Multiscale Molecular Dynamics Studies Reveal Different Modes of Receptor Clustering by Gb3-Binding Lectins. <i>Journal of Chemical Theory and Computation</i> , 2021, 17, 2488-2501.  | 5.3 | 15        |
| 4  | $\beta$ -Hairpin Folding by a Model Amyloid Peptide in Solution and at an Interface. <i>Journal of Physical Chemistry B</i> , 2008, 112, 9476-9483.   | 2.6 | 12        |
| 5  | Fusion-Relevant Changes in Lipid Shape of Hydrated Cholesterol Hemisuccinate Induced by pH and Counterion Species. <i>Journal of Physical Chemistry B</i> , 2010, 114, 14941-14946.   | 2.6 | 12        |
| 6  | Binding of SV40's Viral Capsid Protein VP1 to Its Glycosphingolipid Receptor GM1 Induces Negative Membrane Curvature: A Molecular Dynamics Study. <i>Langmuir</i> , 2019, 35, 3534-3544.                                    | 3.5 | 10        |
| 7  | Effect of Sodium and Chloride Binding on a Lecithin Bilayer. A Molecular Dynamics Study. <i>Membranes</i> , 2017, 7, 5.   | 3.0 | 7         |
| 8  | Mechanisms for allosteric activation of protease DegS by ligand binding and oligomerization as revealed from molecular dynamics simulations. <i>Proteins: Structure, Function and Bioinformatics</i> , 2016, 84, 1690-1705. | 2.6 | 6         |
| 9  | Structure Formation in Langmuir Peptide Films As Revealed from Coarse-Grained Molecular Dynamics Simulations. <i>Langmuir</i> , 2017, 33, 6492-6502.  | 3.5 | 5         |
| 10 | Common Force Field Thermodynamics of Cholesterol. <i>Scientific World Journal</i> , The, 2013, 2013, 1-7.   | 2.1 | 1         |
| 11 | Cover Image, Volume 84, Issue 11. <i>Proteins: Structure, Function and Bioinformatics</i> , 2016, 84, C1.   | 2.6 | 0         |
| 12 | Thermodynamic Origin of Multilayer Structures in Langmuir Polymer Films. <i>Langmuir</i> , 2017, 33, 11399-11405.   | 3.5 | 0         |