

# Edyta Dyguda-Kazimierowicz

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

23  
papers

484  
citations

10  
h-index

22  
g-index

23  
ext. papers

553  
ext. citations

4.4  
avg, IF

3.38  
L-index

#	Paper	IF	Citations
23	Interaction between DNA, Albumin and Apo-Transferrin and Iridium(III) Complexes with Phosphines Derived from Fluoroquinolones as a Potent Anticancer Drug. <i>Pharmaceuticals</i> , <b>2021</b> , 14,	5.2	2
22	Synthesis, structural characterization, docking simulation and in vitro antiproliferative activity of the new gold(III) complex with 2-pyridineethanol. <i>Journal of Inorganic Biochemistry</i> , <b>2021</b> , 215, 111311	4.2	4
21	Extension of an Atom-Atom Dispersion Function to Halogen Bonds and Its Use for Rational Design of Drugs and Biocatalysts. <i>Journal of Physical Chemistry A</i> , <b>2021</b> , 125, 1787-1799	2.8	2
20	Revisiting the halogen bonding between phosphodiesterase type 5 and its inhibitors. <i>Journal of Molecular Modeling</i> , <b>2019</b> , 25, 29	2	4
19	Exploiting the S4-S5 Specificity of Human Neutrophil Proteinase 3 to Improve the Potency of Peptidyl Di(chlorophenyl)-phosphonate Ester Inhibitors: A Kinetic and Molecular Modeling Analysis. <i>Journal of Medicinal Chemistry</i> , <b>2018</b> , 61, 1858-1870	8.3	11
18	Validation of approximate nonempirical scoring model for menin-mixed lineage leukemia inhibitors. <i>Theoretical Chemistry Accounts</i> , <b>2018</b> , 137, 1	1.9	2
17	Theoretical Model of EphA2-Ephrin A1 Inhibition. <i>Molecules</i> , <b>2018</b> , 23,	4.8	3
16	Theoretical models of inhibitory activity for inhibitors of protein-protein interactions: targeting menin-mixed lineage leukemia with small molecules. <i>MedChemComm</i> , <b>2017</b> , 8, 2216-2227	5	6
15	Application of a simple quantum chemical approach to ligand fragment scoring for Trypanosoma brucei pteridine reductase inhibition. <i>Journal of Computer-Aided Molecular Design</i> , <b>2017</b> , 31, 715-728	4.2	6
14	Tracking molecular charge distribution along reaction paths with atomic multipole moments. <i>Structural Chemistry</i> , <b>2016</b> , 27, 429-438	1.8	5
13	Rational Design of Orthogonal Multipolar Interactions with Fluorine in Protein-Ligand Complexes. <i>Journal of Medicinal Chemistry</i> , <b>2015</b> , 58, 7465-74	8.3	52
12	Alkaline hydrolysis of organophosphorus pesticides: the dependence of the reaction mechanism on the incoming group conformation. <i>Journal of Physical Chemistry B</i> , <b>2014</b> , 118, 7277-89	3.4	34
11	Physical Nature of Fatty Acid Amide Hydrolase Interactions with Its Inhibitors: Testing a Simple Nonempirical Scoring Model. <i>Journal of Physical Chemistry B</i> , <b>2014</b> , 118, 14727-36	3.4	8
10	Nonempirical energetic analysis of reactivity and covalent inhibition of fatty acid amide hydrolase. <i>Journal of Physical Chemistry B</i> , <b>2013</b> , 117, 6656-66	3.4	10
9	Low cost prediction of relative stabilities of hydrogen bonded complexes from atomic multipole moments for overly short intermolecular distances. <i>Journal of Computational Chemistry</i> , <b>2013</b> , 34, 1797-9	3.5	8
8	Understanding the role of carbamate reactivity in fatty acid amide hydrolase inhibition by QM/MM mechanistic modelling. <i>Chemical Communications</i> , <b>2011</b> , 47, 2517-9	5.8	21
7	Gas-phase mechanisms of degradation of hazardous organophosphorus compounds: do they follow a common pattern of alkaline hydrolysis reaction as in phosphotriesterase?. <i>Journal of Physical Chemistry B</i> , <b>2008</b> , 112, 9982-91	3.4	26

6	Physical nature of intermolecular interactions within cAMP-dependent protein kinase active site: differential transition state stabilization in phosphoryl transfer reaction. <i>Journal of Physical Chemistry B</i> , <b>2008</b> , 112, 11819-26	3.4	25
5	Non-empirical study of the phosphorylation reaction catalyzed by 4-methyl-5-beta-hydroxyethylthiazole kinase: relevance of the theory of intermolecular interactions. <i>Journal of Molecular Modeling</i> , <b>2007</b> , 13, 839-49	2	8
4	The molecular basis of urokinase inhibition: from the nonempirical analysis of intermolecular interactions to the prediction of binding affinity. <i>Journal of Molecular Modeling</i> , <b>2007</b> , 13, 677-83	2	19
3	Quantitative classification of covalent and noncovalent H-bonds. <i>Journal of Physical Chemistry B</i> , <b>2006</b> , 110, 6444-6	3.4	197
2	Origins of the activity of PAL and LAP enzyme inhibitors: toward ab initio binding affinity prediction. <i>Journal of the American Chemical Society</i> , <b>2005</b> , 127, 1658-9	16.4	19
1	The Mechanism of Phosphoryl Transfer Reaction and the Role of Active Site Residues on the Basis of Ribokinase-Like Kinases. <i>International Journal of Molecular Sciences</i> , <b>2004</b> , 5, 141-153	6.3	12