

# Arkusz Berlicki

## List of Publications by Year in descending order

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75  
papers

2,774  
citations

218381

26  
h-index

182168

51  
g-index

81  
all docs

81  
docs citations

81  
times ranked

3220  
citing authors

#	ARTICLE	IF	CITATIONS
1	Remarkable Potential of the Î±-Aminophosphonate/Phosphinate Structural Motif in Medicinal Chemistry. <i>Journal of Medicinal Chemistry</i> , 2011, 54, 5955-5980.	2.9	529
2	Peptides Containing Î²-Amino Acid Patterns: Challenges and Successes in Medicinal Chemistry. <i>Journal of Medicinal Chemistry</i> , 2014, 57, 9718-9739.	2.9	249
3	Urease inhibitors as potential drugs for gastric and urinary tract infections: a patent review. <i>Expert Opinion on Therapeutic Patents</i> , 2011, 21, 945-957.	2.4	153
4	Peptide-based inhibitors of proteinâ€“protein interactions. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2016, 26, 707-713.	1.0	136
5	Bioactive Macrocyclic Inhibitors of the PDâ€“1/PDâ€“1 Immune Checkpoint. <i>Angewandte Chemie - International Edition</i> , 2017, 56, 13732-13735.	7.2	131
6	1,2-Benziselenazol-3(2 <i>H</i> )-one Derivatives As a New Class of Bacterial Urease Inhibitors. <i>Journal of Medicinal Chemistry</i> , 2016, 59, 8125-8133.	2.9	82
7	Design, Synthesis, and Evaluation of Novel Organophosphorus Inhibitors of Bacterial Ureases. <i>Journal of Medicinal Chemistry</i> , 2008, 51, 5736-5744.	2.9	81
8	Unique Î±,Î±- and Î±,Î²,Î²,Î²-Peptide Foldamers Based on <i>cis</i> -Î²-Aminocyclopentanecarboxylic Acid. <i>Angewandte Chemie - International Edition</i> , 2012, 51, 2208-2212.	7.2	80
9	Computer-Aided Analysis and Design of Phosphonic and Phosphinic Enzyme Inhibitors as Potential Drugs and Agrochemicals. <i>Current Organic Chemistry</i> , 2005, 9, 1829-1850.	0.9	64
10	Computer-Aided Optimization of Phosphinic Inhibitors of Bacterial Ureases. <i>Journal of Medicinal Chemistry</i> , 2010, 53, 5597-5606.	2.9	59
11	Inhibitors of Glutamine Synthetase and their Potential Application in Medicine. <i>Mini-Reviews in Medicinal Chemistry</i> , 2008, 8, 869-878.	1.1	52
12	Tailoring the Structure of Aminobisphosphonates To Target Plant P5C Reductase. <i>Journal of Agricultural and Food Chemistry</i> , 2008, 56, 3193-3199.	2.4	51
13	Synthesis and Evaluation of Effective Inhibitors of Plant Î <sup>1</sup> -Pyrroline-5-carboxylate Reductase. <i>Journal of Agricultural and Food Chemistry</i> , 2013, 61, 6792-6798.	2.4	51
14	The crystal structure of <i>Sporosarcina pasteurii</i> urease in a complex with citrate provides new hints for inhibitor design. <i>Journal of Biological Inorganic Chemistry</i> , 2013, 18, 391-399.	1.1	49
15	Structure-Guided, Single-Point Modifications in the Phosphinic Dipeptide Structure Yield Highly Potent and Selective Inhibitors of Neutral Aminopeptidases. <i>Journal of Medicinal Chemistry</i> , 2014, 57, 8140-8151.	2.9	49
16	Structural Insights into Substrate Selectivity and Activity of Bacterial Polyphosphate Kinases. <i>ACS Catalysis</i> , 2018, 8, 10746-10760.	5.5	48
17	Replacement of Thr <sup>32</sup> and Gln <sup>34</sup> in the <i>C</i> -Terminal Neuropeptide Y Fragment 25â€“36 by <i>cis</i> -Cyclobutane and <i>cis</i> -Cyclopentane Î²-Amino Acids Shifts Selectivity toward the Y <sub>4</sub> Receptor. <i>Journal of Medicinal Chemistry</i> , 2013, 56, 8422-8431.	2.9	46
18	Insight into the mechanism of three component condensation leading to aminomethylenebisphosphonates. <i>Journal of Organometallic Chemistry</i> , 2009, 694, 3806-3813.	0.8	40

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19	Plant P5C Reductase as a New Target for Aminomethylenebisphosphonates. <i>Journal of Agricultural and Food Chemistry</i> , 2007, 55, 4340-4347.	2.4	38
20	Effectiveness and mode of action of phosphonate inhibitors of plant glutamine synthetase. <i>Pest Management Science</i> , 2010, 66, 51-58.	1.7	38
21	N-substituted aminomethanephosphonic and aminomethane-P-methylphosphinic acids as inhibitors of ureases. <i>Amino Acids</i> , 2012, 42, 1937-1945.	1.2	38
22	Design, Synthesis, and Activity of Analogues of Phosphinothricin as Inhibitors of Glutamine Synthetase. <i>Journal of Medicinal Chemistry</i> , 2005, 48, 6340-6349.	2.9	34
23	Bis(aminomethyl)phosphinic Acid, a Highly Promising Scaffold for the Development of Bacterial Urease Inhibitors. <i>ACS Medicinal Chemistry Letters</i> , 2015, 6, 146-150.	1.3	31
24	Enantiodifferentiation of aminophosphonic and aminophosphinic acids with $\beta$ - and $\gamma$ -cyclodextrins. <i>Tetrahedron: Asymmetry</i> , 2003, 14, 1535-1539.	1.8	29
25	Herbicidal Pyridyl Derivatives of Aminomethylene-bisphosphonic Acid Inhibit Plant Glutamine Synthetase. <i>Journal of Agricultural and Food Chemistry</i> , 2004, 52, 3337-3344.	2.4	28
26	Enantiodifferentiation of $\alpha$ -hydroxyalkanephosphonic acids in $^{31}\text{P}$ NMR with application of $\beta$ -cyclodextrin as chiral discriminating agent. <i>Chirality</i> , 2010, 22, 63-68.	1.3	24
27	A structural insight into the P1 S1 binding mode of diaminoethylphosphonic and phosphinic acids, selective inhibitors of alanine aminopeptidases. <i>European Journal of Medicinal Chemistry</i> , 2016, 117, 187-196.	2.6	24
28	Phytotoxicity of aminobisphosphonates targeting both $\beta$ -pyrroline-5-carboxylate reductase and glutamine synthetase. <i>Pest Management Science</i> , 2017, 73, 435-443.	1.7	22
29	Cinchona alkaloids as privileged chiral solvating agents for the enantiodiscrimination of N-protected aminoalkanephosphonates – a comparative NMR study. <i>Tetrahedron: Asymmetry</i> , 2009, 20, 2709-2714.	1.8	21
30	Phosphinothricin Analogues as Inhibitors of Plant Glutamine Synthetases. <i>Journal of Agricultural and Food Chemistry</i> , 2006, 54, 796-802.	2.4	20
31	An integrated approach to the ligand binding specificity of <i>Neisseria meningitidis</i> M1 alanine aminopeptidase by fluorogenic substrate profiling, inhibitory studies and molecular modeling. <i>Biochimie</i> , 2013, 95, 419-428.	1.3	20
32	$\beta$ -Pyrroline-5-carboxylate reductase as a new target for therapeutics: inhibition of the enzyme from <i>Streptococcus pyogenes</i> and effects in vivo. <i>Amino Acids</i> , 2012, 42, 2283-2291.	1.2	18
33	Discovery of new leads against <i>Mycobacterium tuberculosis</i> using scaffold hopping and shape based similarity. <i>Bioorganic and Medicinal Chemistry</i> , 2017, 25, 4835-4844.	1.4	18
34	Covalent Inhibition of Bacterial Urease by Bifunctional Catechol-Based Phosphonates and Phosphinates. <i>Journal of Medicinal Chemistry</i> , 2021, 64, 404-416.	2.9	18
35	The use of molecular modelling for comparison of three possible modes of action of herbicidally active derivatives of aminomethylenebisphosphonic acid. <i>Pesticide Biochemistry and Physiology</i> , 2002, 73, 94-103.	1.6	17
36	Potent covalent inhibitors of bacterial urease identified by activity-reactivity profiling. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2017, 27, 1346-1350.	1.0	17

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37	Controlling the Helix Handedness of $\pm$ Peptide Foldamers through Sequence Shifting. <i>Angewandte Chemie - International Edition</i> , 2017, 56, 2087-2091.	7.2	16
38	Novel organophosphorus scaffolds of urease inhibitors obtained by substitution of Morita-Baylis-Hillman adducts with phosphorus nucleophiles. <i>European Journal of Medicinal Chemistry</i> , 2017, 133, 107-120.	2.6	16
39	Phosphorylation as a method of tuning the enantiodiscrimination potency of quinine – An NMR study. <i>Chirality</i> , 2012, 24, 318-328.	1.3	15
40	Bisphosphonic acids as effective inhibitors of <i>Mycobacterium tuberculosis</i> glutamine synthetase. <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2016, 31, 931-938.	2.5	15
41	Autofluorescence of Amyloids Determined by Enantiomeric Composition of Peptides. <i>Journal of Physical Chemistry B</i> , 2021, 125, 5502-5510.	1.2	15
42	Structural exploration of cinnamate-based phosphonic acids as inhibitors of bacterial ureases. <i>European Journal of Medicinal Chemistry</i> , 2018, 159, 307-316.	2.6	14
43	Computer-aided analysis of the interactions of glutamine synthetase with its inhibitors. <i>Bioorganic and Medicinal Chemistry</i> , 2006, 14, 4578-4585.	1.4	13
44	Bioactive Macrocyclic Inhibitors of the PD-1/PD-L1 Immune Checkpoint. <i>Angewandte Chemie</i> , 2017, 129, 13920-13923.	1.6	13
45	Cyclodextrins as NMR probes in the study of the enantiomeric compositions of N-benzyloxycarbonylamino-phosphonic and phosphinic acids. <i>Tetrahedron: Asymmetry</i> , 2004, 15, 1597-1602.	1.8	12
46	Organophosphorus Supramolecular Chemistry Part 1. Receptors for Organophosphorus Compounds. <i>Current Organic Chemistry</i> , 2006, 10, 2285-2306.	0.9	12
47	Organophosphorus Supramolecular Chemistry. Part 2. Organophosphorus Receptors. <i>Current Organic Chemistry</i> , 2007, 11, 1593-1609.	0.9	12
48	Chiral discrimination of ethyl and phenyl N-benzyloxycarbonylamino-phosphonates by cyclodextrins. <i>Tetrahedron: Asymmetry</i> , 2007, 18, 1579-1584.	1.8	12
49	A three-component synthesis of aminomethylenebis-H-phosphinates. <i>Tetrahedron Letters</i> , 2014, 55, 219-223.	0.7	12
50	Zwitterionic Phosphorylated Quinines as Chiral Solvating Agents for NMR Spectroscopy. <i>Chirality</i> , 2015, 27, 752-760.	1.3	12
51	Catechol-based inhibitors of bacterial urease. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2019, 29, 1085-1089.	1.0	12
52	Enantiodifferentiation of N-benzyloxycarbonylamino-phosphonic and phosphinic acids and their esters using cyclodextrins by means of capillary electrophoresis. <i>Journal of Chromatography A</i> , 2007, 1138, 284-290.	1.8	11
53	Systematic $\alpha$ -foldamerization of peptide inhibiting p53-MDM2/X interactions by the incorporation of trans- or cis-2-aminocyclopentanecarboxylic acid residues. <i>European Journal of Medicinal Chemistry</i> , 2020, 208, 112814.	2.6	11
54	Rational Development of Bacterial Ureases Inhibitors. <i>Chemical Record</i> , 2022, 22, e202200026.	2.9	11

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55	Helix-loop-helix peptide foldamers and their use in the construction of hydrolase mimetics. <i>Bioorganic Chemistry</i> , 2018, 81, 356-361.	2.0	10
56	Aminophosphinates against <i>Helicobacter pylori</i> ureolysis – Biochemical and whole-cell inhibition characteristics. <i>PLoS ONE</i> , 2017, 12, e0182437.	1.1	9
57	Toward very potent, non-covalent organophosphonate inhibitors of cathepsin C and related enzymes by 2-amino-1-hydroxy-alkanephosphonates dipeptides. <i>Biochimie</i> , 2013, 95, 1640-1649.	1.3	8
58	Bisphosphonic acids and related compounds as inhibitors of nucleotide and polyphosphate processing enzymes: A PPK1 and PPK2 case study. <i>Chemical Biology and Drug Design</i> , 2019, 93, 1197-1206.	1.5	8
59	Hierarchical approach for the rational construction of helix-containing nanofibrils using $\alpha$ , $\beta$ -peptides. <i>Nanoscale</i> , 2021, 13, 4000-4015.	2.8	8
60	Sequence Engineering to Control the Helix Handedness of Peptide Foldamers. <i>Chemistry - A European Journal</i> , 2017, 23, 14980-14986.	1.7	8
61	Analysis of pH-Dependent complexation of $\alpha$ -benzyloxycarbonylamino phosphonic acids by $\beta$ -cyclodextrin. Enantiodifferentiation of phosphonic acid $pK_a$ values. <i>Chirality</i> , 2007, 19, 764-768.	1.3	7
62	Three component Kabachnik-Fields condensation leading to substituted aminomethane-P-hydroxymethylphosphonic acids as a tool for screening of bacterial urease inhibitors. <i>Arkivoc</i> , 2012, 2012, 33-43.	0.3	6
63	A Conformationally Stable Acyclic $\alpha$ -Hairpin Scaffold Tolerating the Incorporation of Poorly $\beta$ -Sheet-Prone Amino Acids. <i>ChemBioChem</i> , 2022, 23, .	1.3	6
64	Miniproteins in medicinal chemistry. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2022, 71, 128806.	1.0	6
65	A computationally designed $\beta$ -amino acid-containing miniprotein. <i>Chemical Communications</i> , 2021, 57, 6015-6018.	2.2	5
66	Constrained beta-amino acid-containing miniproteins. <i>Organic and Biomolecular Chemistry</i> , 2021, 19, 4272-4278.	1.5	5
67	Covalent and noncovalent constraints yield a figure eight-like conformation of a peptide inhibiting the menin-MLL interaction. <i>European Journal of Medicinal Chemistry</i> , 2020, 207, 112748.	2.6	4
68	Towards Foldameric Miniproteins: A Helix-Turn-Helix Motif. <i>ChemPlusChem</i> , 2021, 86, 646-649.	1.3	4
69	Controlling the Helix Handedness of $\alpha$ -Peptide Foldamers through Sequence Shifting. <i>Angewandte Chemie</i> , 2017, 129, 2119-2123.	1.6	3
70	Nuclear immunophilin FKBP39 from <i>Drosophila melanogaster</i> drives spontaneous liquid-liquid phase separation. <i>International Journal of Biological Macromolecules</i> , 2020, 163, 108-119.	3.6	3
71	Design and Engineering of Miniproteins. <i>ACS Bio &amp; Med Chem Au</i> , 2022, 2, 316-327.	1.7	3
72	From Inhibitors of Lap to Inhibitors of Pal. Challenges and Advances in Computational Chemistry and Physics, 2007, , 365-398.	0.6	1

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73	Frontispiece: Sequence Engineering to Control the Helix Handedness of Peptide Foldamers. Chemistry - A European Journal, 2017, 23, .	1.7	0
74	Controlling the conformational stability of coiled-coil peptides with a single stereogenic center of a peripheral $\beta^2$ -amino acid residue. RSC Advances, 2022, 12, 4640-4647.	1.7	0
75	Phosphinic acid-based enzyme inhibitors. Phosphorus, Sulfur and Silicon and the Related Elements, 0, , 1-6.	0.8	0