

# Andrej Perdih

## List of Publications by Year in descending order

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61  
papers

1,433  
citations

331259

21  
h-index

344852

36  
g-index

62  
all docs

62  
docs citations

62  
times ranked

1949  
citing authors

| #  | ARTICLE  | IF  | CITATIONS |
|----|--|-----|-----------|
| 1  | Structure-Based Discovery of Substituted 4,5-Bithiazoles as Novel DNA Gyrase Inhibitors. <i>Journal of Medicinal Chemistry</i> , 2012, 55, 6413-6426.  | 2.9 | 146       |
| 2  | Recent Advances in the Synthesis of Unnatural $\beta$ -Amino Acids. <i>Current Organic Chemistry</i> , 2007, 11, 801-832.  | 0.9 | 87        |
| 3  | Discovery of novel benzene 1,3-dicarboxylic acid inhibitors of bacterial MurD and MurE ligases by structure-based virtual screening approach. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2009, 19, 2668-2673. | 1.0 | 67        |
| 4  | Binding free energy calculations of N-sulphonyl-glutamic acid inhibitors of MurD ligase. <i>Journal of Molecular Modeling</i> , 2009, 15, 983-996.   | 0.8 | 57        |
| 5  | The Application of Freidinger Lactams and their Analogs in the Design of Conformationally Constrained Peptidomimetics. <i>Current Medicinal Chemistry</i> , 2006, 13, 1525-1556.                                       | 1.2 | 55        |
| 6  | In silico discovery of 2-amino-4-(2,4-dihydroxyphenyl)thiazoles as novel inhibitors of DNA gyrase B. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2010, 20, 958-962.  | 1.0 | 54        |
| 7  | Targeted molecular dynamics simulation studies of binding and conformational changes in E. coli MurD. <i>Proteins: Structure, Function and Bioinformatics</i> , 2007, 68, 243-254.                                     | 1.5 | 51        |
| 8  | Recent Developments of DNA Poisons - Human DNA Topoisomerase II $\alpha$ Inhibitors - as Anticancer Agents. <i>Current Pharmaceutical Design</i> , 2013, 19, 2474-2488.  | 0.9 | 49        |
| 9  | Recent Advances in the Development of Catalytic Inhibitors of Human DNA Topoisomerase II $\alpha$ ; As Novel Anticancer Agents. <i>Current Medicinal Chemistry</i> , 2013, 20, 694-709.                                | 1.2 | 42        |
| 10 | In silico discovery and biophysical evaluation of novel 5-(2-hydroxybenzylidene) rhodanine inhibitors of DNA gyrase B. <i>Bioorganic and Medicinal Chemistry</i> , 2012, 20, 2572-2580.                                | 1.4 | 41        |
| 11 | Flavonoids and cinnamic acid esters as inhibitors of fungal 17 $\beta$ -hydroxysteroid dehydrogenase: A synthesis, QSAR and modelling study. <i>Bioorganic and Medicinal Chemistry</i> , 2006, 14, 7404-7418.          | 1.4 | 40        |
| 12 | Small Molecule Antagonists of Integrin Receptors. <i>Current Medicinal Chemistry</i> , 2010, 17, 2371-2392.  | 1.2 | 39        |
| 13 | Benzene-1,3-dicarboxylic acid 2,5-dimethylpyrrole derivatives as multiple inhibitors of bacterial Mur ligases (MurC-MurF). <i>Bioorganic and Medicinal Chemistry</i> , 2014, 22, 4124-4134.                            | 1.4 | 34        |
| 14 | Recent Advances in the Synthesis of Unnatural $\beta$ -Amino Acids - An Updated Version. <i>Current Organic Chemistry</i> , 2011, 15, 3750-3799.   | 0.9 | 33        |
| 15 | 1,3-Diamine-Derived Bifunctional Organocatalyst Prepared from Camphor. <i>Advanced Synthesis and Catalysis</i> , 2016, 358, 3786-3796.   | 2.1 | 32        |
| 16 | Discovery of Mono- and Disubstituted 1 <i>H</i> -Pyrazolo[3,4]pyrimidines and 9 <i>H</i> -Purines as Catalytic Inhibitors of Human DNA Topoisomerase II $\alpha$ . <i>ChemMedChem</i> , 2015, 10, 345-359.             | 1.6 | 30        |
| 17 | The CWB2 Cell Wall-Anchoring Module Is Revealed by the Crystal Structures of the <i>Clostridium difficile</i> Cell Wall Proteins Cwp8 and Cwp6. <i>Structure</i> , 2017, 25, 514-521.                                  | 1.6 | 29        |
| 18 | Senescent cells as promising targets to tackle age-related diseases. <i>Ageing Research Reviews</i> , 2021, 66, 101251.  | 5.0 | 28        |

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|----|--|-----|-----------|
| 19 | 4,6-Substituted-1,3,5-triazin-2(1H)-ones as monocyclic catalytic inhibitors of human DNA topoisomerase III $\pm$ targeting the ATP binding site. <i>Bioorganic and Medicinal Chemistry</i> , 2015, 23, 4218-4229.  | 1.4 | 27        |
| 20 | Combinatorially-generated library of 6-fluoroquinolone analogs as potential novel antitubercular agents: a chemometric and molecular modeling assessment. <i>Journal of Molecular Modeling</i> , 2012, 18, 1735-1753.  | 0.8 | 26        |
| 21 | Molecular dynamics simulation and linear interaction energy study of d-Glu-based inhibitors of the MurD ligase. <i>Journal of Computer-Aided Molecular Design</i> , 2013, 27, 723-738.   | 1.3 | 25        |
| 22 | Discovery and Characterization of 2,5-Substituted Benzoic Acid Dual Inhibitors of the Anti-apoptotic Mcl-1 and Bfl-1 Proteins. <i>Journal of Medicinal Chemistry</i> , 2020, 63, 2489-2510.  | 2.9 | 23        |
| 23 | Development of a one-pot assay for screening and identification of Mur pathway inhibitors in <i>Mycobacterium tuberculosis</i> . <i>Scientific Reports</i> , 2016, 6, 35134.   | 1.6 | 22        |
| 24 | Monocyclic 4-amino-6-(phenylamino)-1,3,5-triazines as inhibitors of human DNA topoisomerase III $\pm$ . <i>Bioorganic and Medicinal Chemistry Letters</i> , 2014, 24, 5762-5768.   | 1.0 | 21        |
| 25 | Inhibitor Design Strategy Based on an Enzyme Structural Flexibility: A Case of Bacterial MurD Ligase. <i>Journal of Chemical Information and Modeling</i> , 2014, 54, 1451-1466.   | 2.5 | 21        |
| 26 | Furan-based benzene mono- and dicarboxylic acid derivatives as multiple inhibitors of the bacterial Mur ligases (MurC $\hat{=}$ MurF): experimental and computational characterization. <i>Journal of Computer-Aided Molecular Design</i> , 2015, 29, 541-560. | 1.3 | 21        |
| 27 | Cluster $\hat{=}$ based molecular docking study for <i>in silico</i> identification of novel 6 $\hat{=}$ fluoroquinolones as potential inhibitors against <i>Mycobacterium tuberculosis</i> . <i>Journal of Computational Chemistry</i> , 2013, 34, 790-801.   | 1.5 | 20        |
| 28 | In Silico Discovery of a Substituted 6-Methoxy-quinalidine with Leishmanicidal Activity in <i>Leishmania infantum</i> . <i>Molecules</i> , 2018, 23, 772.  | 1.7 | 20        |
| 29 | Structure-guided optimization of 4,6-substituted-1,3,5-triazin-2(1H)-ones as catalytic inhibitors of human DNA topoisomerase III $\pm$ . <i>European Journal of Medicinal Chemistry</i> , 2019, 175, 330-348.  | 2.6 | 20        |
| 30 | Discovery of the first inhibitors of bacterial enzyme d-aspartate ligase from <i>Enterococcus faecium</i> (Aslfm). <i>European Journal of Medicinal Chemistry</i> , 2013, 67, 208-220.   | 2.6 | 19        |
| 31 | Substituted 4,5 $\hat{=}$ -Bithiazoles as Catalytic Inhibitors of Human DNA Topoisomerase III $\pm$ . <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 3662-3678.   | 2.5 | 19        |
| 32 | MurD ligase from <i>E. coli</i> : Tetrahedral intermediate formation study by hybrid quantum mechanical/molecular mechanical replica path method. <i>Proteins: Structure, Function and Bioinformatics</i> , 2009, 74, 744-759.                                 | 1.5 | 18        |
| 33 | From carbohydrates to drug-like fragments: Rational development of novel $\hat{=}$ -amylase inhibitors. <i>Bioorganic and Medicinal Chemistry</i> , 2015, 23, 6725-6732.   | 1.4 | 18        |
| 34 | Structural Analysis of a Peptide Fragment of Transmembrane Transporter Protein Bilitranslocase. <i>PLoS ONE</i> , 2012, 7, e38967.   | 1.1 | 17        |
| 35 | The Participation of 2 $\hat{=}$ H $\hat{=}$ Pyran $\hat{=}$ ones in [4+2] Cycloadditions: An Experimental and Computational Study. <i>European Journal of Organic Chemistry</i> , 2010, 2010, 5870-5883.  | 1.2 | 16        |
| 36 | Bioassays and In Silico Methods in the Identification of Human DNA Topoisomerase III $\pm$ Inhibitors. <i>Current Medicinal Chemistry</i> , 2018, 25, 3286-3318.   | 1.2 | 15        |

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|----|--|-----|-----------|
| 37 | MurD ligase from <i>Escherichia coli</i> : C-terminal domain closing motion. <i>Computational and Theoretical Chemistry</i> , 2012, 979, 73-81.  | 1.1 | 14        |
| 38 | A Water-Assisted Catalytic Mechanism in Glycoside Hydrolases Demonstrated on the <i>Staphylococcus aureus</i> Autolysin E. <i>ACS Catalysis</i> , 2018, 8, 4334-4345.  | 5.5 | 13        |
| 39 | Discovery of Mcl-1 inhibitors from integrated high throughput and virtual screening. <i>Scientific Reports</i> , 2018, 8, 10210.   | 1.6 | 13        |
| 40 | In Silico Discovery of Novel Potent Antioxidants on the Basis of Pulvic Acid and Coumarine Derivatives and Their Experimental Evaluation. <i>PLoS ONE</i> , 2015, 10, e0140602.  | 1.1 | 12        |
| 41 | Organocatalyzed Deracemization of $\beta$ -Pyrrolinones. <i>Advanced Synthesis and Catalysis</i> , 2017, 359, 2288-2296.   | 2.1 | 11        |
| 42 | Design and synthesis of 3,5-substituted 1,2,4-oxadiazoles as catalytic inhibitors of human DNA topoisomerase II $\alpha$ . <i>Bioorganic Chemistry</i> , 2020, 99, 103828.   | 2.0 | 11        |
| 43 | Distinct labelling of fusion events in rat lactotrophs by FM 1743 and FM 4764 is associated with conformational differences. <i>Acta Physiologica</i> , 2007, 191, 35-42.  | 1.8 | 9         |
| 44 | 3,5-substituted 1,2,4-oxadiazoles as Catalytic Inhibitors of the Human DNA Topoisomerase II $\alpha$ . <i>ChemistrySelect</i> , 2017, 2, 480-488.  | 0.7 | 9         |
| 45 | Evaluation of Firefly and Renilla Luciferase Inhibition in Reporter-Gene Assays: A Case of Isoflavonoids. <i>International Journal of Molecular Sciences</i> , 2021, 22, 6927.   | 1.8 | 8         |
| 46 | Dynophore-Based Approach in Virtual Screening: A Case of Human DNA Topoisomerase II $\alpha$ . <i>International Journal of Molecular Sciences</i> , 2021, 22, 13474.   | 1.8 | 8         |
| 47 | Domain sliding of two <i>Staphylococcus aureus</i> N-acetylglucosaminidases enables their substrate-binding prior to its catalysis. <i>Communications Biology</i> , 2020, 3, 178.  | 2.0 | 7         |
| 48 | Structural elucidation of transmembrane transporter protein bilitranslocase: Conformational analysis of the second transmembrane region TM2 by molecular dynamics and NMR spectroscopy. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2013, 1828, 2609-2619. | 1.4 | 6         |
| 49 | The in vitro protective effects of the three novel nanomolar reversible inhibitors of human cholinesterases against irreversible inhibition by organophosphorous chemical warfare agents. <i>Chemico-Biological Interactions</i> , 2019, 309, 108714.                | 1.7 | 6         |
| 50 | Design, Synthesis and Evaluation of Fused Bicyclo[2.2.2]octene as a Potential Core Scaffold for the Non-Covalent Inhibitors of SARS-CoV-2 3CLpro Main Protease. <i>Pharmaceuticals</i> , 2022, 15, 539.  | 1.7 | 6         |
| 51 | Discovery of (phenylureido)piperidinyl benzamides as prospective inhibitors of bacterial autolysin E from <i>Staphylococcus aureus</i> . <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2018, 33, 1239-1247.  | 2.5 | 4         |
| 52 | Structural Analysis and Dynamic Processes of the Transmembrane Segment Inside Different Micellar Environments: Implications for the TM4 Fragment of the Bilitranslocase Protein. <i>International Journal of Molecular Sciences</i> , 2019, 20, 4172.                | 1.8 | 4         |
| 53 | All-Atom Simulations Reveal a Key Interaction Network in the HLA-E/NKG2A/CD94 Immune Complex Fine-Tuned by the Nonameric Peptide. <i>Journal of Chemical Information and Modeling</i> , 2021, 61, 3593-3603.   | 2.5 | 4         |
| 54 | Nonameric Peptide Orchestrates Signal Transduction in the Activating HLA-E/NKG2C/CD94 Immune Complex as Revealed by All-Atom Simulations. <i>International Journal of Molecular Sciences</i> , 2021, 22, 6670.   | 1.8 | 3         |

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|----|---|-----|-----------|
| 55 | Molecular recognition of acetylcholinesterase and its subnanomolar reversible inhibitor: a molecular simulations study. <i>Journal of Biomolecular Structure and Dynamics</i> , 2022, 40, 1671-1691.  | 2.0 | 2         |
| 56 | 3,5-Substituted Oxadiazoles as Catalytic Inhibitors of the Human Topoisomerase III $\beta$ Molecular Motor. <i>Proceedings (mdpi)</i> , 2019, 22, 78.   | 0.2 | 0         |
| 57 | Potential modulating effect of the <i>Ascaris suum</i> nicotinic acetylcholine receptor (nAChR) by compounds GSK575594A, diazepam and flumazenil discovered by structure-based virtual screening approach. <i>Molecular and Biochemical Parasitology</i> , 2021, 242, 111350. | 0.5 | 0         |
| 58 | Abstract 1641: Structure-based design and development of pyrazolopyridine-based inhibitors of Mcl-1. , 2014, , .  |     | 0         |
| 59 | Abstract 3658: Structure-guidedde novodesign of selective Mcl-1 Inhibitors: Synthesis, structural and biochemical characterization. , 2015, , .   |     | 0         |
| 60 | Interdomain conformational flexibility of the peptidoglycan N-acetylglucosaminidases of <i>Staphylococcus aureus</i> . <i>Acta Crystallographica Section A: Foundations and Advances</i> , 2019, 75, e125-e125.   | 0.0 | 0         |
| 61 | Abstract LB-226: Discovery of small molecule Mcl-1 and Bfl-1 inhibitors. , 2020, , .  |     | 0         |