

Giuseppina De Simone

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.

155
papers

7,529
citations

45
h-index

83
g-index

163
ext. papers

8,098
ext. citations

5.6
avg, IF

5.92
L-index

| # | Paper | IF | Citations |
|-----|---|------|-----------|
| 155 | Interaction Studies between Carbonic Anhydrase and a Sulfonamide Inhibitor by Experimental and Theoretical Approaches. <i>ACS Medicinal Chemistry Letters</i> , 2022 , 13, 271-277 | 4.3 | 1 |
| 154 | Sultam based Carbonic Anhydrase VII inhibitors for the management of neuropathic pain. <i>European Journal of Medicinal Chemistry</i> , 2022 , 227, 113956 | 6.8 | 1 |
| 153 | Inhibition of carbonic anhydrases IX/XII by SLC-0111 boosts cisplatin effects in hampering head and neck squamous carcinoma cell growth and invasion.. <i>Journal of Experimental and Clinical Cancer Research</i> , 2022 , 41, 122 | 12.8 | 2 |
| 152 | Beta-Carbonic Anhydrase 1 from <i>Trichomonas Vaginalis</i> as New Antiprotozoan Drug Target. <i>Topics in Medicinal Chemistry</i> , 2021 , 1 | 0.4 | |
| 151 | Post-translational modifications in tumor-associated carbonic anhydrases. <i>Amino Acids</i> , 2021 , | 3.5 | 5 |
| 150 | Inhibition of the β carbonic anhydrase from the protozoan pathogen with sulphonamides. <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2021 , 36, 329-334 | 5.6 | 3 |
| 149 | Benzyl alcohol inhibits carbonic anhydrases by anchoring to the zinc coordinated water molecule. <i>Biochemical and Biophysical Research Communications</i> , 2021 , 548, 217-221 | 3.4 | 4 |
| 148 | Intrinsically disordered features of carbonic anhydrase IX proteoglycan-like domain. <i>Cellular and Molecular Life Sciences</i> , 2021 , 78, 2059-2067 | 10.3 | 2 |
| 147 | Zeta-carbonic anhydrases show CS hydrolase activity: A new metabolic carbon acquisition pathway in diatoms?. <i>Computational and Structural Biotechnology Journal</i> , 2021 , 19, 3427-3436 | 6.8 | 4 |
| 146 | The crystal structures of 2-(4-benzhydrylpiperazin-1-yl)-N-(4-sulfamoylphenyl)acetamide in complex with human carbonic anhydrase II and VII provide insights into selective CA inhibitor development. <i>New Journal of Chemistry</i> , 2021 , 45, 147-152 | 3.6 | 1 |
| 145 | Design, synthesis and biochemical evaluation of novel carbonic anhydrase inhibitors triggered by structural knowledge on hCA VII. <i>Bioorganic and Medicinal Chemistry</i> , 2021 , 44, 116279 | 3.4 | 0 |
| 144 | Phenyl(thio)phosphon(amid)ate Benzenesulfonamides as Potent and Selective Inhibitors of Human Carbonic Anhydrases II and VII Counteract Allodynia in a Mouse Model of Oxaliplatin-Induced Neuropathy. <i>Journal of Medicinal Chemistry</i> , 2020 , 63, 5185-5200 | 8.3 | 6 |
| 143 | 2-Mercaptobenzoxazoles: a class of carbonic anhydrase inhibitors with a novel binding mode to the enzyme active site. <i>Chemical Communications</i> , 2020 , 56, 8297-8300 | 5.8 | 3 |
| 142 | Biochemical and structural characterisation of a protozoan beta-carbonic anhydrase from. <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2020 , 35, 1292-1299 | 5.6 | 18 |
| 141 | Looking toward the Rim of the Active Site Cavity of Druggable Human Carbonic Anhydrase Isoforms. <i>ACS Medicinal Chemistry Letters</i> , 2020 , 11, 1000-1005 | 4.3 | 1 |
| 140 | Inhibition of the newly discovered β carbonic anhydrase from the protozoan pathogen <i>Trichomonas vaginalis</i> with inorganic anions and small molecules. <i>Journal of Inorganic Biochemistry</i> , 2020 , 213, 111274 | 4.2 | 8 |
| 139 | Human carbonic anhydrases and post-translational modifications: a hidden world possibly affecting protein properties and functions. <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2020 , 35, 1450-1461 | 5.6 | 11 |

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| 138 | A Novel Inhibitor of Carbonic Anhydrases Prevents Hypoxia-Induced TNBC Cell Plasticity. <i>International Journal of Molecular Sciences</i> , 2020 , 21, | 6.3 | 5 |
| 137 | Carbonic Anhydrase Inhibitors Targeting Metabolism and Tumor Microenvironment. <i>Metabolites</i> , 2020 , 10, | 5.6 | 70 |
| 136 | Catechols: a new class of carbonic anhydrase inhibitors. <i>Chemical Communications</i> , 2020 , 56, 13033-13036 | 5.8 | 13 |
| 135 | Exploration of the residues modulating the catalytic features of human carbonic anhydrase XIII by a site-specific mutagenesis approach. <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2019 , 34, 1506-1510 | 5.6 | 6 |
| 134 | Carbonic anhydrases 2019 , 131-137 | | |
| 133 | Carbonic anhydrases 2019 , 19-54 | | 1 |
| 132 | Carbonic anhydrases 2019 , 139-148 | | |
| 131 | Biochemical and Structural Insights into Carbonic Anhydrase XII/Fab6A10 Complex. <i>Journal of Molecular Biology</i> , 2019 , 431, 4910-4921 | 6.5 | 13 |
| 130 | Exploring benzoxaborole derivatives as carbonic anhydrase inhibitors: a structural and computational analysis reveals their conformational variability as a tool to increase enzyme selectivity. <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2019 , 34, 1498-1505 | 5.6 | 5 |
| 129 | Targeted treatment of anaerobic cancer. Patent evaluation of US2016279084 and US2017056350. <i>Expert Opinion on Therapeutic Patents</i> , 2019 , 29, 1-6 | 6.8 | 4 |
| 128 | Exploring structural properties of potent human carbonic anhydrase inhibitors bearing a 4-(cycloalkylamino-1-carbonyl)benzenesulfonamide moiety. <i>European Journal of Medicinal Chemistry</i> , 2019 , 163, 443-452 | 6.8 | 16 |
| 127 | Inhibition of carbonic anhydrase IX targets primary tumors, metastases, and cancer stem cells: Three for the price of one. <i>Medicinal Research Reviews</i> , 2018 , 38, 1799-1836 | 14.4 | 159 |
| 126 | Biochemical, biophysical and molecular dynamics studies on the proteoglycan-like domain of carbonic anhydrase IX. <i>Cellular and Molecular Life Sciences</i> , 2018 , 75, 3283-3296 | 10.3 | 13 |
| 125 | The Crystal Structure of a hCA VII Variant Provides Insights into the Molecular Determinants Responsible for Its Catalytic Behavior. <i>International Journal of Molecular Sciences</i> , 2018 , 19, | 6.3 | 19 |
| 124 | Inhibition of carbonic anhydrases by a substrate analog: benzyl carbamate directly coordinates the catalytic zinc ion mimicking bicarbonate binding. <i>Chemical Communications</i> , 2018 , 54, 10312-10315 | 5.8 | 14 |
| 123 | Are Carbonic Anhydrases Suitable Targets to Fight Protozoan Parasitic Diseases?. <i>Current Medicinal Chemistry</i> , 2018 , 25, 5266-5278 | 4.3 | 18 |
| 122 | Crystal structure of the human carbonic anhydrase II adduct with 1-(4-sulfamoylphenyl-ethyl)-2,4,6-triphenylpyridinium perchlorate, a membrane-impermeant, isoform selective inhibitor. <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2018 , 33, 151-157 | 5.6 | 24 |
| 121 | Protective Role of Carbonic Anhydrases III and VII in Cellular Defense Mechanisms upon Redox Unbalance. <i>Oxidative Medicine and Cellular Longevity</i> , 2018 , 2018, 2018306 | 6.7 | 19 |

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| 120 | Disclosing the Interaction of Carbonic Anhydrase IX with Cullin-Associated NEDD8-Dissociated Protein 1 by Molecular Modeling and Integrated Binding Measurements. <i>ACS Chemical Biology</i> , 2017 , 12, 1460-1465 | 4.9 | 13 |
| 119 | Probing Molecular Interactions between Human Carbonic Anhydrases (hCAs) and a Novel Class of Benzenesulfonamides. <i>Journal of Medicinal Chemistry</i> , 2017 , 60, 4316-4326 | 8.3 | 30 |
| 118 | Insights into the role of reactive sulfhydryl groups of Carbonic Anhydrase III and VII during oxidative damage. <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2017 , 32, 5-12 | 5.6 | 26 |
| 117 | Insights into the binding mode of sulphamates and sulphamides to hCA II: crystallographic studies and binding free energy calculations. <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2017 , 32, 1002-1011 ²⁵ | 5.6 | 25 |
| 116 | Cloning, expression and purification of the complete domain of the β -carbonic anhydrase from <i>Plasmodium falciparum</i> . <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2016 , 31, 54-59 | 5.6 | 50 |
| 115 | Kinetic and X-ray crystallographic investigations of substituted 2-thio-6-oxo-1,6-dihydropyrimidine-benzenesulfonamides acting as carbonic anhydrase inhibitors. <i>Bioorganic and Medicinal Chemistry</i> , 2016 , 24, 3643-8 | 3.4 | 12 |
| 114 | L-Histidinol Dehydrogenase as a New Target for Old Diseases. <i>Current Topics in Medicinal Chemistry</i> , 2016 , 16, 2369-78 | 3 | 5 |
| 113 | A Combined Crystallographic and Theoretical Study Explains the Capability of Carboxylic Acids to Adopt Multiple Binding Modes in the Active Site of Carbonic Anhydrases. <i>Chemistry - A European Journal</i> , 2016 , 22, 97-100 | 4.8 | 34 |
| 112 | The anticonvulsant sulfamide JNJ-26990990 and its S,S-dioxide analog strongly inhibit carbonic anhydrases: solution and X-ray crystallographic studies. <i>Organic and Biomolecular Chemistry</i> , 2016 , 14, 4853-8 | 3.9 | 21 |
| 111 | Benzoxaborole as a new chemotype for carbonic anhydrase inhibition. <i>Chemical Communications</i> , 2016 , 52, 11983-11986 | 5.8 | 60 |
| 110 | X-ray crystallographic and kinetic investigations of 6-sulfamoyl-saccharin as a carbonic anhydrase inhibitor. <i>Organic and Biomolecular Chemistry</i> , 2015 , 13, 4064-9 | 3.9 | 18 |
| 109 | Discovery of 1,1SBiphenyl-4-sulfonamides as a New Class of Potent and Selective Carbonic Anhydrase XIV Inhibitors. <i>Journal of Medicinal Chemistry</i> , 2015 , 58, 8564-72 | 8.3 | 34 |
| 108 | Out of the active site binding pocket for carbonic anhydrase inhibitors. <i>Chemical Communications</i> , 2015 , 51, 302-5 | 5.8 | 96 |
| 107 | Carbonic Anhydrase VII 2015 , 151-168 | | 1 |
| 106 | Carbonic Anhydrase II as Target for Drug Design 2015 , 51-90 | | 2 |
| 105 | Human Carbonic Anhydrases: Catalytic Properties, Structural Features, and Tissue Distribution 2015 , 17-30 | | 5 |
| 104 | Thermostable Carbonic Anhydrases in Biotechnological Applications. <i>International Journal of Molecular Sciences</i> , 2015 , 16, 15456-80 | 6.3 | 50 |
| 103 | Cadmium-containing carbonic anhydrase CDCA1 in marine diatom <i>Thalassiosira weissflogii</i> . <i>Marine Drugs</i> , 2015 , 13, 1688-97 | 6 | 34 |

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| 102 | Hydroxylamine-O-sulfonamide is a versatile lead compound for the development of carbonic anhydrase inhibitors. <i>Chemical Communications</i> , 2015 , 51, 11519-22 | 5.8 | 9 |
| 101 | The zinc coordination pattern in the β -carbonic anhydrase from <i>Plasmodium falciparum</i> is different from all other carbonic anhydrase genetic families. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2015 , 25, 1385-9 | 2.9 | 95 |
| 100 | Crystal structure of the most catalytically effective carbonic anhydrase enzyme known, SazCA from the thermophilic bacterium <i>Sulfurihydrogenibium azorense</i> . <i>Bioorganic and Medicinal Chemistry Letters</i> , 2015 , 25, 2002-6 | 2.9 | 60 |
| 99 | Carbonic Anhydrases: An Overview 2015 , 3-13 | | 11 |
| 98 | CDCA1 From <i>Thalassiosira weissflogii</i> as Representative Member of β -Class CAs: General Features and Biotechnological Applications 2015 , 351-359 | | |
| 97 | <i>Sulfolobus solfataricus</i> thiol redox puzzle: characterization of an atypical protein disulfide oxidoreductase. <i>Extremophiles</i> , 2014 , 18, 219-28 | 3 | 9 |
| 96 | Biochemical characterization of the chloroplastic β -carbonic anhydrase from <i>Flaveria bidentis</i> (L.) "Kuntze". <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2014 , 29, 500-4 | 5.6 | 16 |
| 95 | Thermal-stable carbonic anhydrases: a structural overview. <i>Sub-Cellular Biochemistry</i> , 2014 , 75, 387-404 | 5.5 | 6 |
| 94 | Hydrophobic substituents of the phenylmethylsulfamide moiety can be used for the development of new selective carbonic anhydrase inhibitors. <i>BioMed Research International</i> , 2014 , 2014, 523210 | 3 | 12 |
| 93 | The structural comparison between membrane-associated human carbonic anhydrases provides insights into drug design of selective inhibitors. <i>Biopolymers</i> , 2014 , 101, 769-78 | 2.2 | 39 |
| 92 | High GADA titer increases the risk of insulin requirement in LADA patients: a 7-year follow-up (NIRAD study 7). <i>European Journal of Endocrinology</i> , 2014 , 171, 697-704 | 6.5 | 45 |
| 91 | Structural basis for the rational design of new anti- <i>Brucella</i> agents: the crystal structure of the C366S mutant of L-histidinol dehydrogenase from <i>Brucella suis</i> . <i>Biochimie</i> , 2014 , 97, 114-20 | 4.6 | 8 |
| 90 | Faox enzymes inhibited Maillard reaction development during storage both in protein glucose model system and low lactose UHT milk. <i>Amino Acids</i> , 2014 , 46, 279-88 | 3.5 | 23 |
| 89 | Amine, amino acid and oligopeptide carbonic anhydrase activators 2014 , 142-156 | | |
| 88 | Exploiting the hydrophobic and hydrophilic binding sites for designing carbonic anhydrase inhibitors. <i>Expert Opinion on Drug Discovery</i> , 2013 , 8, 793-810 | 6.2 | 215 |
| 87 | Characterization of carbonic anhydrase IX interactome reveals proteins assisting its nuclear localization in hypoxic cells. <i>Journal of Proteome Research</i> , 2013 , 12, 282-92 | 5.6 | 37 |
| 86 | Kinetic and anion inhibition studies of a β -carbonic anhydrase (FbiCA 1) from the C4 plant <i>Flaveria bidentis</i> . <i>Bioorganic and Medicinal Chemistry Letters</i> , 2013 , 23, 1626-30 | 2.9 | 33 |
| 85 | Anticancer carbonic anhydrase inhibitors: a patent review (2008 - 2013). <i>Expert Opinion on Therapeutic Patents</i> , 2013 , 23, 737-49 | 6.8 | 208 |

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| 84 | Hypoxia-targeting carbonic anhydrase IX inhibitors by a new series of nitroimidazole-sulfonamides/sulfamides/sulfamates. <i>Journal of Medicinal Chemistry</i> , 2013 , 56, 8512-20 | 8.3 | 68 |
| 83 | Human carbonic anhydrase VII protects cells from oxidative damage. <i>Biological Chemistry</i> , 2013 , 394, 1343-8 | 4.5 | 29 |
| 82 | X-ray structure of the first "extremo- β -carbonic anhydrase" a dimeric enzyme from the thermophilic bacterium <i>Sulfurihydrogenibium yellowstonense</i> YO3AOP1. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2013 , 69, 1150-9 | | 89 |
| 81 | Dithiocarbamates are strong inhibitors of the beta-class fungal carbonic anhydrases from <i>Cryptococcus neoformans</i> , <i>Candida albicans</i> and <i>Candida glabrata</i> . <i>Bioorganic and Medicinal Chemistry Letters</i> , 2012 , 22, 859-62 | 2.9 | 89 |
| 80 | Carbonic anhydrase VII is S-glutathionylated without loss of catalytic activity and affinity for sulfonamide inhibitors. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2012 , 22, 1560-4 | 2.9 | 46 |
| 79 | (In)organic anions as carbonic anhydrase inhibitors. <i>Journal of Inorganic Biochemistry</i> , 2012 , 111, 117-29 | 4.2 | 173 |
| 78 | Structural and inhibition insights into carbonic anhydrase CDCA1 from the marine diatom <i>Thalassiosira weissflogii</i> . <i>Biochimie</i> , 2012 , 94, 1232-41 | 4.6 | 88 |
| 77 | Development of potent carbonic anhydrase inhibitors incorporating both sulfonamide and sulfamide groups. <i>Journal of Medicinal Chemistry</i> , 2012 , 55, 6776-83 | 8.3 | 43 |
| 76 | Multiple binding modes of inhibitors to carbonic anhydrases: how to design specific drugs targeting 15 different isoforms?. <i>Chemical Reviews</i> , 2012 , 112, 4421-68 | 68.1 | 889 |
| 75 | Hydroxamate represents a versatile zinc binding group for the development of new carbonic anhydrase inhibitors. <i>Chemical Communications</i> , 2012 , 48, 8838-40 | 5.8 | 58 |
| 74 | Carbonic anhydrase IX as a target for designing novel anticancer drugs. <i>Current Medicinal Chemistry</i> , 2012 , 19, 821-30 | 4.3 | 44 |
| 73 | C68 from the <i>Sulfolobus islandicus</i> plasmid-virus pSSVx is a novel member of the AbrB-like transcription factor family. <i>Biochemical Journal</i> , 2011 , 435, 157-66 | 3.8 | 22 |
| 72 | Design, synthesis and characterization of a peptide able to bind proteins of the KCTD family: implications for KCTD-cullin 3 recognition. <i>Journal of Peptide Science</i> , 2011 , 17, 373-6 | 2.1 | 14 |
| 71 | Carbonic anhydrase inhibitors: X-ray crystallographic studies for the binding of N-substituted benzenesulfonamides to human isoform II. <i>Chemical Communications</i> , 2011 , 47, 11636-8 | 5.8 | 42 |
| 70 | Histone deacetylase and Cullin3-REN(KCTD11) ubiquitin ligase interplay regulates Hedgehog signalling through Gli acetylation. <i>Nature Cell Biology</i> , 2010 , 12, 132-42 | 23.4 | 252 |
| 69 | Drug design studies of the novel antitumor targets carbonic anhydrase IX and XII. <i>Current Medicinal Chemistry</i> , 2010 , 17, 1516-26 | 4.3 | 88 |
| 68 | Exploring the catalytic mechanism of the first dimeric Bcp: Functional, structural and docking analyses of Bcp4 from <i>Sulfolobus solfataricus</i> . <i>Biochimie</i> , 2010 , 92, 1435-44 | 4.6 | 20 |
| 67 | The first example of a significant active site conformational rearrangement in a carbonic anhydrase-inhibitor adduct: the carbonic anhydrase I-topiramate complex. <i>Organic and Biomolecular Chemistry</i> , 2010 , 8, 3528-33 | 3.9 | 34 |

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| 66 | Recent advances in structural studies of the carbonic anhydrase family: the crystal structure of human CA IX and CA XIII. <i>Current Pharmaceutical Design</i> , 2010 , 16, 3246-54 | 3.3 | 27 |
| 65 | Multiple catalytically active thioredoxin folds: a winning strategy for many functions. <i>Cellular and Molecular Life Sciences</i> , 2010 , 67, 3797-814 | 10.3 | 24 |
| 64 | Crystal structure of an S-formylglutathione hydrolase from <i>Pseudoalteromonas haloplanktis</i> TAC125. <i>Biopolymers</i> , 2010 , 93, 669-77 | 2.2 | 17 |
| 63 | Carbonic anhydrase IX: Biochemical and crystallographic characterization of a novel antitumor target. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2010 , 1804, 404-9 | 4 | 151 |
| 62 | Carbonic anhydrase inhibitors: crystallographic and solution binding studies for the interaction of a boron-containing aromatic sulfamide with mammalian isoforms I-XV. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2010 , 20, 3601-5 | 2.9 | 25 |
| 61 | Inhibition of the R1 fragment of the cadmium-containing zeta-class carbonic anhydrase from the diatom <i>Thalassiosira weissflogii</i> with anions. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2010 , 20, 4745-8 ^{2.9} | 2.9 | 35 |
| 60 | Crystal structure of the C183S/C217S mutant of human CA VII in complex with acetazolamide. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2010 , 20, 5023-6 | 2.9 | 73 |
| 59 | Crystal structure of the catalytic domain of the tumor-associated human carbonic anhydrase IX. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009 , 106, 16233-8 | 11.5 | 399 |
| 58 | Crystal structure of human carbonic anhydrase XIII and its complex with the inhibitor acetazolamide. <i>Proteins: Structure, Function and Bioinformatics</i> , 2009 , 74, 164-75 | 4.2 | 90 |
| 57 | Insights into the catalytic mechanism of the Bcp family: functional and structural analysis of Bcp1 from <i>Sulfolobus solfataricus</i> . <i>Proteins: Structure, Function and Bioinformatics</i> , 2009 , 76, 995-1006 | 4.2 | 22 |
| 56 | Which carbonic anhydrases are targeted by the antiepileptic sulfonamides and sulfamates?. <i>Chemical Biology and Drug Design</i> , 2009 , 74, 317-21 | 2.9 | 62 |
| 55 | The proteoglycan region of the tumor-associated carbonic anhydrase isoform IX acts as an intrinsic buffer optimizing CO ₂ hydration at acidic pH values characteristic of solid tumors. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2009 , 19, 5825-8 | 2.9 | 73 |
| 54 | Structural analysis of BldR from <i>Sulfolobus solfataricus</i> provides insights into the molecular basis of transcriptional activation in Archaea by MarR family proteins. <i>Journal of Molecular Biology</i> , 2009 , 388, 559-69 | 6.5 | 29 |
| 53 | Carbonic anhydrase inhibitors. Comparison of aliphatic sulfamate/bis-sulfamate adducts with isozymes II and IX as a platform for designing tight-binding, more isoform-selective inhibitors. <i>Journal of Medicinal Chemistry</i> , 2009 , 52, 5990-8 | 8.3 | 19 |
| 52 | Biochemical characterization of CA IX, one of the most active carbonic anhydrase isozymes. <i>Journal of Biological Chemistry</i> , 2008 , 283, 27799-27809 | 5.4 | 224 |
| 51 | Carbonic anhydrase inhibitors: bioreductive nitro-containing sulfonamides with selectivity for targeting the tumor associated isoforms IX and XII. <i>Journal of Medicinal Chemistry</i> , 2008 , 51, 3230-7 | 8.3 | 46 |
| 50 | Structure-activity relationships of C-17 cyano-substituted estratrienes as anticancer agents. <i>Journal of Medicinal Chemistry</i> , 2008 , 51, 1295-308 | 8.3 | 47 |
| 49 | Anticancer steroid sulfatase inhibitors: synthesis of a potent fluorinated second-generation agent, in vitro and in vivo activities, molecular modeling, and protein crystallography. <i>Molecular Cancer Therapeutics</i> , 2008 , 7, 2435-44 | 6.1 | 37 |

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|----|---|------|-----|
| 48 | Recent advances in research on the most novel carbonic anhydrases, CA XIII and XV. <i>Current Pharmaceutical Design</i> , 2008 , 14, 672-8 | 3.3 | 64 |
| 47 | Are carbonic anhydrase inhibitors suitable for obtaining antiobesity drugs?. <i>Current Pharmaceutical Design</i> , 2008 , 14, 655-60 | 3.3 | 137 |
| 46 | Functional and structural features of the oxyanion hole in a thermophilic esterase from <i>Alicyclobacillus acidocaldarius</i> . <i>Proteins: Structure, Function and Bioinformatics</i> , 2008 , 71, 1721-31 | 4.2 | 26 |
| 45 | Carbonic anhydrase inhibitors: binding of indanesulfonamides to the human isoform II. <i>ChemMedChem</i> , 2008 , 3, 473-7 | 3.7 | 10 |
| 44 | Investigations of the esterase, phosphatase, and sulfatase activities of the cytosolic mammalian carbonic anhydrase isoforms I, II, and XIII with 4-nitrophenyl esters as substrates. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2008 , 18, 2267-71 | 2.9 | 88 |
| 43 | Carbonic anhydrase inhibitors: the X-ray crystal structure of ethoxzolamide complexed to human isoform II reveals the importance of thr200 and gln92 for obtaining tight-binding inhibitors. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2008 , 18, 2669-74 | 2.9 | 30 |
| 42 | Carbonic anhydrase inhibitors as emerging drugs for the treatment of obesity. <i>Expert Opinion on Emerging Drugs</i> , 2008 , 13, 383-92 | 3.7 | 139 |
| 41 | Molecular modeling study for the binding of zonisamide and topiramate to the human mitochondrial carbonic anhydrase isoform VA. <i>Bioorganic and Medicinal Chemistry</i> , 2007 , 15, 4152-8 | 3.4 | 34 |
| 40 | Carbonic anhydrase inhibitors: binding of an antiglaucoma glycosyl-sulfanilamide derivative to human isoform II and its consequences for the drug design of enzyme inhibitors incorporating sugar moieties. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2007 , 17, 1726-31 | 2.9 | 35 |
| 39 | Carbonic anhydrase inhibitors: inhibition of human, bacterial, and archaeal isozymes with benzene-1,3-disulfonamides--solution and crystallographic studies. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2007 , 17, 4201-7 | 2.9 | 45 |
| 38 | Antiobesity carbonic anhydrase inhibitors. <i>Current Topics in Medicinal Chemistry</i> , 2007 , 7, 879-84 | 3 | 82 |
| 37 | Carbonic anhydrase inhibitors: Valdecoxib binds to a different active site region of the human isoform II as compared to the structurally related cyclooxygenase II "selective" inhibitor celecoxib. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2006 , 16, 437-42 | 2.9 | 89 |
| 36 | Carbonic anhydrase inhibitors: X-ray crystallographic studies for the binding of 5-amino-1,3,4-thiadiazole-2-sulfonamide and 5-(4-amino-3-chloro-5-fluorophenylsulfonamido)-1,3,4-thiadiazole-2-sulfonamide to human isoform II. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2006 , 16, 6204-8 | 2.9 | 31 |
| 35 | 2-substituted estradiol bis-sulfamates, multitargeted antitumor agents: synthesis, in vitro SAR, protein crystallography, and in vivo activity. <i>Journal of Medicinal Chemistry</i> , 2006 , 49, 7683-96 | 8.3 | 91 |
| 34 | Carbonic anhydrase inhibitors: Hypoxia-activatable sulfonamides incorporating disulfide bonds that target the tumor-associated isoform IX. <i>Journal of Medicinal Chemistry</i> , 2006 , 49, 5544-51 | 8.3 | 93 |
| 33 | Carbonic anhydrase inhibitors: X-ray and molecular modeling study for the interaction of a fluorescent antitumor sulfonamide with isozyme II and IX. <i>Journal of the American Chemical Society</i> , 2006 , 128, 8329-35 | 16.4 | 186 |
| 32 | Metal ion substitution in the catalytic site greatly affects the binding of sulfhydryl-containing compounds to leucyl aminopeptidase. <i>Biochemistry</i> , 2006 , 45, 3226-34 | 3.2 | 30 |
| 31 | Insights on a new PDI-like family: structural and functional analysis of a protein disulfide oxidoreductase from the bacterium <i>Aquifex aeolicus</i> . <i>Journal of Molecular Biology</i> , 2006 , 356, 155-64 | 6.5 | 26 |

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| 30 | A novel member of the protein disulfide oxidoreductase family from <i>Aeropyrum pernix</i> K1: structure, function and electrostatics. <i>Journal of Molecular Biology</i> , 2006 , 362, 743-52 | 6.5 | 20 |
| 29 | Carbonic anhydrase inhibitors: stacking with Phe131 determines active site binding region of inhibitors as exemplified by the X-ray crystal structure of a membrane-impermeant antitumor sulfonamide complexed with isozyme II. <i>Journal of Medicinal Chemistry</i> , 2005 , 48, 5721-7 | 8.3 | 150 |
| 28 | Carbonic anhydrase inhibitors: X-ray crystal structure of a benzenesulfonamide strong CA II and CA IX inhibitor bearing a pentafluorophenylaminothioureido tail in complex with isozyme II. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2005 , 15, 1937-42 | 2.9 | 38 |
| 27 | Carbonic anhydrase inhibitors. Zonisamide is an effective inhibitor of the cytosolic isozyme II and mitochondrial isozyme V: solution and X-ray crystallographic studies. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2005 , 15, 2315-20 | 2.9 | 166 |
| 26 | Crystallization and preliminary X-ray diffraction studies of a protein disulfide oxidoreductase from <i>Aeropyrum pernix</i> K1. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2005 , 61, 335-6 | | 4 |
| 25 | A substrate-induced switch in the reaction mechanism of a thermophilic esterase: kinetic evidences and structural basis. <i>Journal of Biological Chemistry</i> , 2004 , 279, 6815-23 | 5.4 | 37 |
| 24 | Crystallization and preliminary X-ray diffraction studies of a protein disulfide oxidoreductase from <i>Aquifex aeolicus</i> . <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2004 , 60, 2076-7 | | 6 |
| 23 | The crystal structure of an EST2 mutant unveils structural insights on the H group of the carboxylesterase/lipase family. <i>Journal of Molecular Biology</i> , 2004 , 343, 137-46 | 6.5 | 24 |
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