

# Kimberly A Stieglitz

## List of Publications by Year in descending order

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24  
papers

384  
citations

840776

11  
h-index

794594

19  
g-index

24  
all docs

24  
docs citations

24  
times ranked

438  
citing authors

#	ARTICLE	IF	CITATIONS
1	Characterization of recombinant fructose-1,6-bisphosphatase gene mutations: evidence of inhibition/activation of FB Pase protein by gene mutation. <i>Bioscience Reports</i> , 2019, 39, .	2.4	4
2	Structural Insights for Drugs Developed for Phospholipase D Enzymes. <i>Current Drug Discovery Technologies</i> , 2018, 15, 81-93.	1.2	9
3	Structural Analysis of Relevant Drug Targets for Alzheimer's Disease: Novel Approaches to Drug Development. <i>Current Bioactive Compounds</i> , 2017, 13, 90-100.	0.5	2
4	Identifying New Drug Targets for Potent Phospholipase D Inhibitors: Combining Sequence Alignment, Molecular Docking, and Enzyme Activity/Binding Assays. <i>Chemical Biology and Drug Design</i> , 2016, 87, 714-729.	3.2	2
5	1,3-Disubstituted 4-Aminopyrazolo [3, 4-d] Pyrimidines, a New Class of Potent Inhibitors for Phospholipase D. <i>Chemical Biology and Drug Design</i> , 2014, 84, 270-281.	3.2	14
6	The Role of Phospholipase D Enzyme(s) in Modulating Cell Signaling: Implications for Cancer Drug Development. <i>Current Bioactive Compounds</i> , 2014, 10, 124-130.	0.5	0
7	Rational Design, Synthesis, and Potency of N-Substituted Indoles, Pyrroles, and Triarylpyrazoles as Potential Fructose 1,6-Bisphosphatase Inhibitors. <i>ChemMedChem</i> , 2010, 5, 384-389.	3.2	23
8	Mobile loop mutations in an archaeal inositol monophosphatase: Modulating three-metal ion assisted catalysis and lithium inhibition. <i>Protein Science</i> , 2010, 19, 309-318.	7.6	15
9	The first high pH structure of <i>Escherichia coli</i> aspartate transcarbamoylase. <i>Proteins: Structure, Function and Bioinformatics</i> , 2009, 74, 318-327.	2.6	8
10	Novel Heteroaromatic Organofluorine Inhibitors of Fructose-1,6-bisphosphatase. <i>Journal of Medicinal Chemistry</i> , 2009, 52, 878-882.	6.4	24
11	Not so clear on oxygen. Comment on <i>Structural basis for cofactor-independent dioxygenation in vancomycin biosynthesis</i> by Widboom et al. (2007), <i>Nature (London)</i> , 447, 342-345. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2008, 64, 1000-1002.	2.5	3
12	The Structure of the R184A Mutant of the Inositol Monophosphatase Encoded by <i>suhB</i> and Implications for Its Functional Interactions in <i>Escherichia coli</i> . <i>Journal of Biological Chemistry</i> , 2007, 282, 26989-26996.	3.4	20
13	Comparison of two T-state structures of regulatory-chain mutants of <i>Escherichia coli</i> aspartate transcarbamoylase suggests that His20 and Asp19 modulate the response to heterotropic effectors. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2007, 63, 1243-1253.	2.5	1
14	Crystal structure of the tetrameric inositol 1-phosphate phosphatase (TM1415) from the hyperthermophile, <i>Thermotoga maritima</i> . <i>FEBS Journal</i> , 2007, 274, 2461-2469.	4.7	23
15	T-state Inhibitors of <i>E. coli</i> Aspartate Transcarbamoylase that Prevent the Allosteric Transition. <i>Biochemistry</i> , 2006, 45, 10062-10071.	2.5	11
16	Structural basis for ordered substrate binding and cooperativity in aspartate transcarbamoylase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 8881-8886.	7.1	46
17	Reaching for Mechanistic Consensus Across Life Kingdoms: A Structure and Insights into Catalysis of thymo-Inositol-1-phosphate Synthase (mIPS) from <i>Archaeoglobus fulgidus</i> . <i>Biochemistry</i> , 2005, 44, 213-224.	2.5	33
18	A Single Amino Acid Substitution in the Active Site of <i>Escherichia coli</i> Aspartate Transcarbamoylase Prevents the Allosteric Transition. <i>Journal of Molecular Biology</i> , 2005, 349, 413-423.	4.2	8

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19	Structure of the E.coli Aspartate Transcarbamoylase Trapped in the Middle of the Catalytic Cycle. <i>Journal of Molecular Biology</i> , 2005, 352, 478-486.	4.2	10
20	Metal Specificity Is Correlated with Two Crucial Active Site Residues in Escherichia coli Alkaline Phosphatase. <i>Biochemistry</i> , 2005, 44, 8378-8386.	2.5	47
21	240s Loop Interactions Stabilize the T State of Escherichia coli Aspartate Transcarbamoylase. <i>Journal of Biological Chemistry</i> , 2004, 279, 23302-23310.	3.4	4
22	Unexpected similarity in regulation between an archaeal inositol monophosphatase/fructose bisphosphatase and chloroplast fructose bisphosphatase. <i>Protein Science</i> , 2003, 12, 760-767.	7.6	8
23	Crystal Structure of a Dual Activity IMPase/FBPase (AF2372) from <i>Archaeoglobus fulgidus</i> . <i>Journal of Biological Chemistry</i> , 2002, 277, 22863-22874.	3.4	48
24	Binding of Proteolytically Processed Phospholipase D from <i>Streptomyces chromofuscus</i> to Phosphatidylcholine Membranes Facilitates Vesicle Aggregation and Fusion. <i>Biochemistry</i> , 2001, 40, 13954-13963.	2.5	21