

# Rui Wu

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11162011/publications.pdf>

Version: 2024-02-01

13

papers

319

citations

933447

10

h-index

1125743

13

g-index

13

all docs

13

docs citations

13

times ranked

495

citing authors

#	ARTICLE	IF	CITATIONS
1	Crystal structure of <i>Bacillus subtilis</i> GabR, an autorepressor and transcriptional activator of <i>gabT</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 17820-17825.	7.1	66
2	The Active Site Sulfenic Acid Ligand in Nitrile Hydratases Can Function as a Nucleophile. <i>Journal of the American Chemical Society</i> , 2014, 136, 1186-1189.	13.7	54
3	Rational Design of a Transition State Analogue with Picomolar Affinity for <i>Pseudomonas aeruginosa</i> PvdQ, a Siderophore Biosynthetic Enzyme. <i>ACS Chemical Biology</i> , 2013, 8, 2192-2200.	3.4	41
4	Mechanism of Inactivation of $\beta$ -Aminobutyric Acid Aminotransferase by (1 <i>S</i> ,3 <i>S</i> )-3-Amino-4-difluoromethylene-1-cyclopentanoic Acid (CPP-115). <i>Journal of the American Chemical Society</i> , 2015, 137, 2628-2640.	13.7	29
5	PLP and GABA trigger GabR-mediated transcription regulation in <i>Bacillus subtilis</i> via external aldimine formation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 3891-3896.	7.1	26
6	The Crystal Structure of Nitrosomonas europaea Sucrose Synthase Reveals Critical Conformational Changes and Insights into Sucrose Metabolism in Prokaryotes. <i>Journal of Bacteriology</i> , 2015, 197, 2734-2746.	2.2	23
7	<i>n</i> -Alkylboronic Acid Inhibitors Reveal Determinants of Ligand Specificity in the Quorum-Quenching and Siderophore Biosynthetic Enzyme PvdQ. <i>Biochemistry</i> , 2014, 53, 6679-6686.	2.5	17
8	Design and Mechanism of Tetrahydrothiophene-Based $\beta$ -Aminobutyric Acid Aminotransferase Inactivators. <i>Journal of the American Chemical Society</i> , 2015, 137, 4525-4533.	13.7	17
9	Mechanism of Inactivation of GABA Aminotransferase by ( <i>E</i> )- and ( <i>Z</i> )-(1 <i>S</i> ,3 <i>S</i> )-3-Amino-4-fluoromethylene-1-cyclopentanoic Acid. <i>ACS Chemical Biology</i> , 2015, 10, 2087-2098.	3.4	12
10	Structural analysis reveals a pyruvate-binding activator site in the <i>Agrobacterium tumefaciens</i> ADP-“glucose” pyrophosphorylase. <i>Journal of Biological Chemistry</i> , 2019, 294, 1338-1348.	3.4	11
11	Multiple States of Nitrile Hydratase from <i>Rhodococcus equi</i> TG328-2: Structural and Mechanistic Insights from Electron Paramagnetic Resonance and Density Functional Theory Studies. <i>Biochemistry</i> , 2017, 56, 3068-3077.	2.5	9
12	Analyzing the catalytic role of active site residues in the Fe-type nitrile hydratase from <i>Comamonas testosteroni</i> Ni1. <i>Journal of Biological Inorganic Chemistry</i> , 2015, 20, 885-894.	2.6	8
13	Substrate Trapping in the Siderophore Tailoring Enzyme PvdQ. <i>ACS Chemical Biology</i> , 2017, 12, 643-647.	3.4	6