

# Shanteri Singh

## List of Publications by Year in descending order

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46  
papers

1,601  
citations

304743

22  
h-index

302126

39  
g-index

47  
all docs

47  
docs citations

47  
times ranked

1982  
citing authors

#	ARTICLE	IF	CITATIONS
1	Glycosyltransferase structural biology and its role in the design of catalysts for glycosylation. <i>Current Opinion in Biotechnology</i> , 2011, 22, 800-808.	6.6	136
2	Facile Chemoenzymatic Strategies for the Synthesis and Utilization of Adenosyl-L-Methionine Analogues. <i>Angewandte Chemie - International Edition</i> , 2014, 53, 3965-3969.	13.8	120
3	Solution structure of a late embryogenesis abundant protein (LEA14) from <i>Arabidopsis thaliana</i> , a cellular stress-related protein. <i>Protein Science</i> , 2005, 14, 2601-2609.	7.6	104
4	The structural biology of enzymes involved in natural product glycosylation. <i>Natural Product Reports</i> , 2012, 29, 1201.	10.3	99
5	Auto-induction medium for the production of [U-15N]- and [U-13C, U-15N]-labeled proteins for NMR screening and structure determination. <i>Protein Expression and Purification</i> , 2005, 40, 268-278.	1.3	91
6	The structure of flavin-dependent tryptophan 7-halogenase RebH. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008, 70, 289-293.	2.6	89
7	Broadening the scope of glycosyltransferase-catalyzed sugar nucleotide synthesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 7648-7653.	7.1	88
8	Structure and Mechanism of the Rebeccamycin Sugar 4-O-Methyltransferase RebM. <i>Journal of Biological Chemistry</i> , 2008, 283, 22628-22636.	3.4	57
9	Comparison of cell-based and cell-free protocols for producing target proteins from the <i>Arabidopsis thaliana</i> genome for structural studies. <i>Proteins: Structure, Function and Bioinformatics</i> , 2005, 59, 633-643.	2.6	56
10	Biochemical and Structural Insights of the Early Glycosylation Steps in Calicheamicin Biosynthesis. <i>Chemistry and Biology</i> , 2008, 15, 842-853.	6.0	51
11	Structure and specificity of a permissive bacterial C-prenyltransferase. <i>Nature Chemical Biology</i> , 2017, 13, 366-368.	8.0	50
12	Complete set of glycosyltransferase structures in the calicheamicin biosynthetic pathway reveals the origin of regiospecificity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 17649-17654.	7.1	47
13	Understanding molecular recognition of promiscuity of thermophilic methionine adenosyltransferase MAT from <i>Sulfolobus solfataricus</i> . <i>FEBS Journal</i> , 2014, 281, 4224-4239.	4.7	36
14	Functional AdoMet Isosteres Resistant to Classical AdoMet Degradation Pathways. <i>ACS Chemical Biology</i> , 2016, 11, 2484-2491.	3.4	36
15	Structural Insight into the Self-Sacrifice Mechanism of Eneidyne Resistance. <i>ACS Chemical Biology</i> , 2006, 1, 451-460.	3.4	34
16	Venturicidin C, a new 20-membered macrolide produced by <i>Streptomyces</i> sp. TS-2-2. <i>Journal of Antibiotics</i> , 2014, 67, 223-230.	2.0	33
17	Structural characterization of CalO2: A putative orsellinic acid P450 oxidase in the calicheamicin biosynthetic pathway. <i>Proteins: Structure, Function and Bioinformatics</i> , 2009, 74, 50-60.	2.6	27
18	Structural characterization of the mitomycin 7-O-methyltransferase. <i>Proteins: Structure, Function and Bioinformatics</i> , 2011, 79, 2181-2188.	2.6	26

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19	Crystal structure of SsfS6, the putative <i>C</i> -glycosyltransferase involved in SF2575 biosynthesis. <i>Proteins: Structure, Function and Bioinformatics</i> , 2013, 81, 1277-1282.	2.6	24
20	Structure-Guided Functional Characterization of Eneidyne Self-Sacrifice Resistance Proteins, CalU16 and CalU19. <i>ACS Chemical Biology</i> , 2014, 9, 2347-2358.	3.4	24
21	Functionalized Anodic Aluminum Oxide Membraneâ€“Electrode System for Enzyme Immobilization. <i>ACS Nano</i> , 2014, 8, 8104-8112.	14.6	22
22	Structural Polymorphism and Dynamism in the DNA Segment GATCTTCCCCCGGAA:â€‰% NMR Investigations of Hairpin, Dumbbell, Nicked Duplex, Parallel Strands, and i-Motif. <i>Biochemistry</i> , 1997, 36, 13214-13222.	2.5	21
23	Three-dimensional structure of the AAH26994.1 protein from <i>Mus musculus</i> , a putative eukaryotic Urm1. <i>Protein Science</i> , 2005, 14, 2095-2102.	7.6	21
24	Chemoenzymatic synthesis of daptomycin analogs active against daptomycin-resistant strains. <i>Applied Microbiology and Biotechnology</i> , 2020, 104, 7853-7865.	3.6	20
25	FgaPT2, a biocatalytic tool for alkyl-diversification of indole natural products. <i>MedChemComm</i> , 2019, 10, 1465-1475.	3.4	19
26	A Simple Strategy for Glycosyltransferaseâ€“Catalyzed Aminosugar Nucleotide Synthesis. <i>ChemBioChem</i> , 2014, 15, 647-651.	2.6	18
27	Determination of Alkylâ€“Donor Promiscuity of Tyrosineâ€“O-â€“Prenyltransferase SirD from <i>Leptosphaeria maculans</i> . <i>ChemBioChem</i> , 2017, 18, 2323-2327.	2.6	18
28	The native production of the sesquiterpene isopterocarpolone by <i>Streptomyces</i> sp. RM-14-6. <i>Natural Product Research</i> , 2014, 28, 337-339.	1.8	17
29	Rapid Transport of Protons across Membranes by Aliphatic Amines and Acids. <i>The Journal of Physical Chemistry</i> , 1995, 99, 11302-11305.	2.9	16
30	Structural characterization of CalO1: a putative orsellinic acid methyltransferase in the calicheamicin-biosynthetic pathway. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2011, 67, 197-203.	2.5	16
31	Acceptor substrate determines donor specificity of an aromatic prenyltransferase: expanding the biocatalytic potential of NphB. <i>Applied Microbiology and Biotechnology</i> , 2020, 104, 4383-4395.	3.6	14
32	Structure of <i>Arabidopsis thaliana</i> At1g77540 Protein, a Minimal Acetyltransferase from the COG2388 Family. <i>Biochemistry</i> , 2006, 45, 14325-14336.	2.5	13
33	Structural and Functional Characterization of CalS11, a TDP-Rhamnose 3â€“O-Methyltransferase Involved in Calicheamicin Biosynthesis. <i>ACS Chemical Biology</i> , 2013, 8, 1632-1639.	3.4	12
34	Structural Basis for the Stereochemical Control of Amine Installation in Nucleotide Sugar Aminotransferases. <i>ACS Chemical Biology</i> , 2015, 10, 2048-2056.	3.4	12
35	Glycosyloxyamine Neoglycosylation: A Model Study Using Calicheamicin. <i>ChemMedChem</i> , 2011, 6, 774-776.	3.2	11
36	Characterization of the Calicheamicin Orsellinate C2â€“O-Methyltransferase CalO6. <i>ChemBioChem</i> , 2014, 15, 1418-1421.	2.6	10

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37	Structural characterization of AtmS13, a putative sugar aminotransferase involved in indolocarbazole AT2433 aminopentose biosynthesis. <i>Proteins: Structure, Function and Bioinformatics</i> , 2015, 83, 1547-1554.	2.6	10
38	A General NMR-Based Strategy for the in Situ Characterization of Sugar-Nucleotide-Dependent Biosynthetic Pathways. <i>Organic Letters</i> , 2014, 16, 3220-3223.	4.6	9
39	Indole C6 Functionalization of Tryprostatin B Using Prenyltransferase CdpNPT. <i>Catalysts</i> , 2020, 10, 1247.	3.5	9
40	Novel Homologs of Isopentenyl Phosphate Kinase Reveal Class-Wide Substrate Flexibility. <i>ChemCatChem</i> , 2021, 13, 3781-3788.	3.7	8
41	Characterization of Early Enzymes Involved in TDP-Aminodideoxypentose Biosynthesis en Route to Indolocarbazole AT2433. <i>ChemBioChem</i> , 2015, 16, 2141-2146.	2.6	6
42	Structural Characterization of CalS8, a TDP-d-Glucose Dehydrogenase Involved in Calicheamicin Aminodideoxypentose Biosynthesis. <i>Journal of Biological Chemistry</i> , 2015, 290, 26249-26258.	3.4	5
43	Loop dynamics of thymidine diphosphate-rhamnose 3-O-methyltransferase (CalS11), an enzyme in calicheamicin biosynthesis. <i>Structural Dynamics</i> , 2016, 3, 012004.	2.3	5
44	Structural dynamics of a methionine $\beta$ -lyase for calicheamicin biosynthesis: Rotation of the conserved tyrosine stacking with pyridoxal phosphate. <i>Structural Dynamics</i> , 2016, 3, 034702.	2.3	4
45	Evidence for a novel $\beta$ -bend structure with prolines at the corner: 1H and 13C NMR study of cyclo(Pro-Pro-Gly) <sub>2</sub> . <i>Magnetic Resonance in Chemistry</i> , 1993, 31, 944-953.	1.9	3
46	Molecular Basis for the Substrate Promiscuity of Isopentenyl Phosphate Kinase from <i>Candidatus methanomethylophilus alvus</i> . <i>ACS Chemical Biology</i> , 2022, 17, 85-102.	3.4	2