

# Dustin J Little

## List of Publications by Year in descending order

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Version: 2024-02-01

19  
papers

807  
citations

516215

16  
h-index

794141

19  
g-index

21  
all docs

21  
docs citations

21  
times ranked

1040  
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of Poly-N-acetylglucosamine as a Major Polysaccharide Component of the <i>Bacillus subtilis</i> Biofilm Matrix. <i>Journal of Biological Chemistry</i> , 2015, 290, 19261-19272.	1.6	118
2	Sph3 Is a Glycoside Hydrolase Required for the Biosynthesis of Galactosaminogalactan in <i>Aspergillus fumigatus</i> . <i>Journal of Biological Chemistry</i> , 2015, 290, 27438-27450.	1.6	77
3	Characterization of the <i>Pseudomonas aeruginosa</i> Glycoside Hydrolase PslG Reveals That Its Levels Are Critical for Psl Polysaccharide Biosynthesis and Biofilm Formation. <i>Journal of Biological Chemistry</i> , 2015, 290, 28374-28387.	1.6	68
4	The Structure- and Metal-dependent Activity of <i>Escherichia coli</i> PgaB Provides Insight into the Partial De-N-acetylation of Poly- $\beta$ -1,6-N-acetyl-d-glucosamine. <i>Journal of Biological Chemistry</i> , 2012, 287, 31126-31137.	1.6	65
5	PgaB orthologues contain a glycoside hydrolase domain that cleaves deacetylated poly- $\beta$ (1,6)-N-acetylglucosamine and can disrupt bacterial biofilms. <i>PLoS Pathogens</i> , 2018, 14, e1006998.	2.1	59
6	<i>P. aeruginosa</i> SGNH Hydrolase-Like Proteins AlgJ and AlgX Have Similar Topology but Separate and Distinct Roles in Alginate Acetylation. <i>PLoS Pathogens</i> , 2014, 10, e1004334.	2.1	54
7	PilN Binding Modulates the Structure and Binding Partners of the <i>Pseudomonas aeruginosa</i> Type IVa Pilus Protein PilM. <i>Journal of Biological Chemistry</i> , 2016, 291, 11003-11015.	1.6	53
8	Modification and periplasmic translocation of the biofilm exopolysaccharide poly- $\beta$ -1,6-N-acetyl-d-glucosamine. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 11013-11018.	3.3	48
9	PatB1 is an O-acetyltransferase that decorates secondary cell wall polysaccharides. <i>Nature Chemical Biology</i> , 2018, 14, 79-85.	3.9	37
10	In vitro characterization of the antivirulence target of Gram-positive pathogens, peptidoglycan O-acetyltransferase A (OatA). <i>PLoS Pathogens</i> , 2017, 13, e1006667.	2.1	35
11	Functional Characterization of <i>Staphylococcus epidermidis</i> IcaB, a De-N-acetylase Important for Biofilm Formation. <i>Biochemistry</i> , 2013, 52, 5463-5471.	1.2	32
12	The Protein BpsB Is a Poly- $\beta$ -1,6-N-acetyl-d-glucosamine Deacetylase Required for Biofilm Formation in <i>Bordetella bronchiseptica</i> . <i>Journal of Biological Chemistry</i> , 2015, 290, 22827-22840.	1.6	31
13	Structural Basis for the De-N-acetylation of Poly- $\beta$ -1,6-N-acetyl-d-glucosamine in Gram-positive Bacteria. <i>Journal of Biological Chemistry</i> , 2014, 289, 35907-35917.	1.6	24
14	Synthesis and evaluation of inhibitors of <i>E. coli</i> PgaB, a polysaccharide de-N-acetylase involved in biofilm formation. <i>Organic and Biomolecular Chemistry</i> , 2012, 10, 7103.	1.5	22
15	Regulatory Evolution Drives Evasion of Host Inflammasomes by <i>Salmonella Typhimurium</i> . <i>Cell Reports</i> , 2018, 25, 825-832.e5.	2.9	22
16	Functional diversification of the NleG effector family in enterohemorrhagic <i>Escherichia coli</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 10004-10009.	3.3	19
17	Molecular basis for CesT recognition of type III secretion effectors in enteropathogenic <i>Escherichia coli</i> . <i>PLoS Pathogens</i> , 2018, 14, e1007224.	2.1	16
18	Combining <i>in situ</i> proteolysis and mass spectrometry to crystallize <i>Escherichia coli</i> PgaB. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2012, 68, 842-845.	0.7	14

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19	Direct Staudinger-Phosponite Reaction Provides Methylphosphonamidates as Inhibitors of CE4 Deacetylases. ChemBioChem, 2015, 16, 1350-1356.	1.3	11