

# Annabelle Varrot

## List of Publications by Year in descending order

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92  
papers

3,847  
citations

109137

35  
h-index

138251

58  
g-index

99  
all docs

99  
docs citations

99  
times ranked

3736  
citing authors

#	ARTICLE	IF	CITATIONS
1	Structure–function relationship of a novel fucoside-binding fruiting body lectin from <i>Coprinopsis cinerea</i> exhibiting nematotoxic activity. <i>Glycobiology</i> , 2022, , .	1.3	2
2	Characterization of the ganglioside recognition profile of <i>Escherichia coli</i> heat-labile enterotoxin LT-IIc. <i>Glycobiology</i> , 2022, 32, 391-403.	1.3	2
3	Non-Carbohydrate Glycomimetics as Inhibitors of Calcium(II)-Binding Lectins. <i>Angewandte Chemie</i> , 2021, 133, 8185-8195.	1.6	3
4	Non-Carbohydrate Glycomimetics as Inhibitors of Calcium(II)-Binding Lectins. <i>Angewandte Chemie - International Edition</i> , 2021, 60, 8104-8114.	7.2	17
5	Hexavalent thiofucosides to probe the role of the <i>Aspergillus fumigatus</i> lectin FleA in fungal pathogenicity. <i>Organic and Biomolecular Chemistry</i> , 2021, 19, 3234-3240.	1.5	3
6	Prediction and Validation of a Druggable Site on Virulence Factor of Drug Resistant <i>Burkholderia cenocepacia</i> . <i>Chemistry - A European Journal</i> , 2021, 27, 10341-10348.	1.7	6
7	Biochemical and structural studies of target lectin SapL1 from the emerging opportunistic microfungus <i>Scedosporium apiospermum</i> . <i>Scientific Reports</i> , 2021, 11, 16109.	1.6	4
8	Structure and engineering of tandem repeat lectins. <i>Current Opinion in Structural Biology</i> , 2020, 62, 39-47.	2.6	29
9	Characterization of novel lectins from <i>Burkholderia pseudomallei</i> and <i>Chromobacterium violaceum</i> with seven-bladed $\beta$ -propeller fold. <i>International Journal of Biological Macromolecules</i> , 2020, 152, 1113-1124.	3.6	5
10	Recombinant Lectin from Tepary Bean ( <i>Phaseolus acutifolius</i> ) with Specific Recognition for Cancer-Associated Glycans: Production, Structural Characterization, and Target Identification. <i>Biomolecules</i> , 2020, 10, 654.	1.8	6
11	BC2L-C N-Terminal Lectin Domain Complexed with Histo Blood Group Oligosaccharides Provides New Structural Information. <i>Molecules</i> , 2020, 25, 248.	1.7	5
12	Expression, Purification, and Applications of the Recombinant Lectin PVL from <i>Psathyrella velutina</i> Specific for Terminal N-Acetyl-Glucosamine. <i>Methods in Molecular Biology</i> , 2020, 2132, 421-436.	0.4	2
13	Expression, Purification, and Functional Characterization of Tectonin 2 from <i>Laccaria bicolor</i> : A Six-Bladed Beta-Propeller Lectin Specific for O-Methylated Glycans. <i>Methods in Molecular Biology</i> , 2020, 2132, 669-682.	0.4	1
14	LecB, a High Affinity Soluble Fucose-Binding Lectin from <i>Pseudomonas aeruginosa</i> . <i>Methods in Molecular Biology</i> , 2020, 2132, 475-482.	0.4	0
15	LecA (PA-IL): A Galactose-Binding Lectin from <i>Pseudomonas aeruginosa</i> . <i>Methods in Molecular Biology</i> , 2020, 2132, 257-266.	0.4	8
16	Anti-biofilm Agents against <i>Pseudomonas aeruginosa</i> : A Structure–Activity Relationship Study of C-Glycosidic LecB Inhibitors. <i>Journal of Medicinal Chemistry</i> , 2019, 62, 9201-9216.	2.9	45
17	Induction of rare conformation of oligosaccharide by binding to calcium-dependent bacterial lectin: X-ray crystallography and modelling study. <i>European Journal of Medicinal Chemistry</i> , 2019, 177, 212-220.	2.6	6
18	Architecture and Evolution of Blade Assembly in $\beta$ -propeller Lectins. <i>Structure</i> , 2019, 27, 764-775.e3.	1.6	27

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19	Crystal Structures of Fungal Tectonin in Complex with O-Methylated Glycans Suggest Key Role in Innate Immune Defense. <i>Structure</i> , 2018, 26, 391-402.e4.	1.6	28
20	Glycomimetic, Orally Bioavailable LecB Inhibitors Block Biofilm Formation of <i>Pseudomonas aeruginosa</i> . <i>Journal of the American Chemical Society</i> , 2018, 140, 2537-2545.	6.6	97
21	Multivalent Glycomimetics with Affinity and Selectivity toward Fucose-Binding Receptors from Emerging Pathogens. <i>Bioconjugate Chemistry</i> , 2018, 29, 83-88.	1.8	25
22	Multivalent Fucosides with Nanomolar Affinity for the <i>Aspergillus fumigatus</i> Lectin FleA Prevent Spore Adhesion to Pneumocytes. <i>Chemistry - A European Journal</i> , 2018, 24, 19243-19249.	1.7	15
23	Human Bronchial Epithelial Cells Inhibit <i>Aspergillus fumigatus</i> Germination of Extracellular Conidia via FleA Recognition. <i>Scientific Reports</i> , 2018, 8, 15699.	1.6	35
24	Effect of Noncanonical Amino Acids on Protein-Carbohydrate Interactions: Structure, Dynamics, and Carbohydrate Affinity of a Lectin Engineered with Fluorinated Tryptophan Analogs. <i>ACS Chemical Biology</i> , 2018, 13, 2211-2219.	1.6	22
25	Recognition of Complex Core-Fucosylated Glycans by a Mini Lectin. <i>Angewandte Chemie</i> , 2018, 130, 10335-10338.	1.6	2
26	Recognition of Complex Core-Fucosylated Glycans by a Mini Lectin. <i>Angewandte Chemie - International Edition</i> , 2018, 57, 10178-10181.	7.2	15
27	Biophysical characterization and structural determination of the potent cytotoxic <i>Psathyrella asperospora</i> lectin. <i>Proteins: Structure, Function and Bioinformatics</i> , 2017, 85, 969-975.	1.5	10
28	Histo-blood group antigens as mediators of infections. <i>Current Opinion in Structural Biology</i> , 2017, 44, 190-200.	2.6	72
29	Exploiting $sp^2$ -Hybridisation in the Development of Potent 1,5- $\alpha$ -Arabinanase Inhibitors. <i>ChemBioChem</i> , 2017, 18, 974-978.	1.3	1
30	Covalent Lectin Inhibition and Application in Bacterial Biofilm Imaging. <i>Angewandte Chemie - International Edition</i> , 2017, 56, 16559-16564.	7.2	56
31	Covalent Lectin Inhibition and Application in Bacterial Biofilm Imaging. <i>Angewandte Chemie</i> , 2017, 129, 16786-16791.	1.6	12
32	Recombinant fungal lectin as a new tool to investigate O-GlcNAcylation processes. <i>Glycobiology</i> , 2017, 27, 123-128.	1.3	22
33	O-Alkylated heavy atom carbohydrate probes for protein X-ray crystallography: Studies towards the synthesis of methyl 2-O-methyl-L-selenofucopyranoside. <i>Beilstein Journal of Organic Chemistry</i> , 2016, 12, 2828-2833.	1.3	6
34	Genomic Rearrangements and Functional Diversification of lecA and lecB Lectin-Coding Regions Impacting the Efficacy of Glycomimetics Directed against <i>Pseudomonas aeruginosa</i> . <i>Frontiers in Microbiology</i> , 2016, 7, 811.	1.5	39
35	Characterization of a high-affinity sialic acid-specific CBM40 from <i>Clostridium perfringens</i> and engineering of a divalent form. <i>Biochemical Journal</i> , 2016, 473, 2109-2118.	1.7	32
36	Dimerization of the fungal defense lectin CCL2 is essential for its toxicity against nematodes. <i>Glycobiology</i> , 2016, 27, 486-500.	1.3	17

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37	The virulence factor LecB varies in clinical isolates: consequences for ligand binding and drug discovery. <i>Chemical Science</i> , 2016, 7, 4990-5001.	3.7	50
38	Biochemical and structural characterization of the novel sialic acid-binding site of <i>Escherichia coli</i> heat-labile enterotoxin LT-IIb. <i>Biochemical Journal</i> , 2016, 473, 3923-3936.	1.7	9
39	The Hidden Conformation of Lewis x, a Human Histo-Blood Group Antigen, Is a Determinant for Recognition by Pathogen Lectins. <i>ACS Chemical Biology</i> , 2016, 11, 2011-2020.	1.6	37
40	Cinnamide Derivatives of $\alpha$ -Mannose as Inhibitors of the Bacterial Virulence Factor LecB from <i>Pseudomonas aeruginosa</i> . <i>ChemistryOpen</i> , 2015, 4, 756-767.	0.9	35
41	Structural insights into <i>Aspergillus fumigatus</i> lectin specificity: AFL binding sites are functionally non-equivalent. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2015, 71, 442-453.	2.5	27
42	Algal lectin binding to core $\beta$ fucosylated N-glycans: Structural basis for specificity and production of recombinant protein. <i>Glycobiology</i> , 2015, 25, 607-616.	1.3	17
43	A Recombinant Fungal Lectin for Labeling Truncated Glycans on Human Cancer Cells. <i>PLoS ONE</i> , 2015, 10, e0128190.	1.1	25
44	Membrane Deformation by Neolectins with Engineered Glycolipid Binding Sites. <i>Angewandte Chemie - International Edition</i> , 2014, 53, 9267-9270.	7.2	53
45	A LecA Ligand Identified from a Galactoside Conjugate Array Inhibits Host Cell Invasion by <i>Pseudomonas aeruginosa</i> . <i>Angewandte Chemie - International Edition</i> , 2014, 53, 8885-8889.	7.2	85
46	Monitoring Lectin Interactions with Carbohydrates. <i>Methods in Molecular Biology</i> , 2014, 1149, 403-414.	0.4	6
47	Secondary sugar binding site identified for LecA lectin from <i>Pseudomonas aeruginosa</i> . <i>Proteins: Structure, Function and Bioinformatics</i> , 2014, 82, 1060-1065.	1.5	18
48	Fungal lectins: structure, function and potential applications. <i>Current Opinion in Structural Biology</i> , 2013, 23, 678-685.	2.6	116
49	Reduction of Lectin Valency Drastically Changes Glycolipid Dynamics in Membranes but Not Surface Avidity. <i>ACS Chemical Biology</i> , 2013, 8, 1918-1924.	1.6	39
50	Discovery of Two Classes of Potent Glycomimetic Inhibitors of <i>Pseudomonas aeruginosa</i> LecB with Distinct Binding Modes. <i>ACS Chemical Biology</i> , 2013, 8, 1775-1784.	1.6	83
51	Aromatic thioglycoside inhibitors against the virulence factor LecA from <i>Pseudomonas aeruginosa</i> . <i>Organic and Biomolecular Chemistry</i> , 2013, 11, 6906.	1.5	81
52	Bacteria love our sugars: Interaction between soluble lectins and human fucosylated glycans, structures, thermodynamics and design of competing glycocompounds. <i>Comptes Rendus Chimie</i> , 2013, 16, 482-490.	0.2	28
53	Deciphering the Glycan Preference of Bacterial Lectins by Glycan Array and Molecular Docking with Validation by Microcalorimetry and Crystallography. <i>PLoS ONE</i> , 2013, 8, e71149.	1.1	25
54	A Soluble Fucose-Specific Lectin from <i>Aspergillus fumigatus</i> Conidia - Structure, Specificity and Possible Role in Fungal Pathogenicity. <i>PLoS ONE</i> , 2013, 8, e83077.	1.1	87

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55	A Lectin from <i>Platypodium elegans</i> with Unusual Specificity and Affinity for Asymmetric Complex N-Glycans. <i>Journal of Biological Chemistry</i> , 2012, 287, 26352-26364.	1.6	26
56	Fucose-binding Lectin from Opportunistic Pathogen <i>Burkholderia ambifaria</i> Binds to Both Plant and Human Oligosaccharidic Epitopes. <i>Journal of Biological Chemistry</i> , 2012, 287, 4335-4347.	1.6	92
57	1- <i>l</i> -Arabinofuranosylated pyrrolidines as arabinanase inhibitors. <i>Chemical Communications</i> , 2011, 47, 9684.	2.2	6
58	A TNF-like Trimeric Lectin Domain from <i>Burkholderia cenocepacia</i> with Specificity for Fucosylated Human Histo-Blood Group Antigens. <i>Structure</i> , 2010, 18, 59-72.	1.6	76
59	Contact-Killing Polyelectrolyte Microcapsules Based on Chitosan Derivatives. <i>Advanced Functional Materials</i> , 2010, 20, 3303-3312.	7.8	50
60	Structural basis of the affinity for oligomannosides and analogs displayed by BC2L-A, a <i>Burkholderia cenocepacia</i> soluble lectin. <i>Glycobiology</i> , 2010, 20, 87-98.	1.3	48
61	Structure and RNA recognition by the snRNA and snoRNA transport factor PHAX. <i>Rna</i> , 2010, 16, 1205-1216.	1.6	18
62	Discoidin I from <i>Dictyostelium discoideum</i> and Interactions with Oligosaccharides: Specificity, Affinity, Crystal Structures, and Comparison with Discoidin II. <i>Journal of Molecular Biology</i> , 2010, 400, 540-554.	2.0	34
63	Carbohydrate binding specificities and crystal structure of the cholera toxin-like B-subunit from <i>Citrobacter freundii</i> . <i>Biochimie</i> , 2010, 92, 482-490.	1.3	15
64	Role of Water Molecules in Structure and Energetics of <i>Pseudomonas aeruginosa</i> Lectin I Interacting with Disaccharides. <i>Journal of Biological Chemistry</i> , 2010, 285, 20316-20327.	1.6	37
65	Molecular Basis of Arabinobio-hydrolase Activity in Phytopathogenic Fungi. <i>Journal of Biological Chemistry</i> , 2009, 284, 12285-12296.	1.6	42
66	Structure determination of discoidin II from <i>Dictyostelium discoideum</i> and carbohydrate binding properties of the lectin domain. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008, 73, 43-52.	1.5	25
67	High Affinity Interaction between a Bivalve C-type Lectin and a Biantennary Complex-type N-Glycan Revealed by Crystallography and Microcalorimetry. <i>Journal of Biological Chemistry</i> , 2008, 283, 30112-30120.	1.6	35
68	Microbial recognition of human cell surface glycoconjugates. <i>Current Opinion in Structural Biology</i> , 2008, 18, 567-576.	2.6	253
69	Structural Basis of the Preferential Binding for Globo-Series Glycosphingolipids Displayed by <i>Pseudomonas aeruginosa</i> Lectin I. <i>Journal of Molecular Biology</i> , 2008, 383, 837-853.	2.0	133
70	Glycosylated asterisks are among the most potent low valency inducers of Concanavalin A aggregation. <i>Chemical Communications</i> , 2008, , 6507-6509.	2.2	28
71	Structural basis for mannose recognition by a lectin from opportunistic bacteria <i>Burkholderia cenocepacia</i> . <i>Biochemical Journal</i> , 2008, 411, 307-318.	1.7	74
72	<i>Mycobacterium tuberculosis</i> Strains Possess Functional Cellulases. <i>Journal of Biological Chemistry</i> , 2005, 280, 20181-20184.	1.6	62

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73	NAD <sup>+</sup> and Metal-ion Dependent Hydrolysis by Family 4 Glycosidases: Structural Insight into Specificity for Phospho-β <sup>2</sup> -d-glucosides. <i>Journal of Molecular Biology</i> , 2005, 346, 423-435.	2.0	52
74	Novel Catalytic Mechanism of Glycoside Hydrolysis Based on the Structure of an NAD <sup>+</sup> /Mn <sup>2+</sup> -Dependent Phospho-β <sup>2</sup> -Glucosidase from <i>Bacillus subtilis</i> . <i>Structure</i> , 2004, 12, 1619-1629.	1.6	88
75	An Unusual Mechanism of Glycoside Hydrolysis Involving Redox and Elimination Steps by a Family 4 β <sup>2</sup> -Glucosidase from <i>Thermotoga maritima</i> . <i>Journal of the American Chemical Society</i> , 2004, 126, 8354-8355.	6.6	119
76	Structural Basis for Ligand Binding and Processivity in Cellobiohydrolase Cel6A from <i>Humicola insolens</i> . <i>Structure</i> , 2003, 11, 855-864.	1.6	116
77	Direct experimental observation of the hydrogen-bonding network of a glycosidase along its reaction coordinate revealed by atomic resolution analyses of endoglucanase Cel5A. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2003, 59, 447-452.	2.5	29
78	Direct Observation of the Protonation State of an Imino Sugar Glycosidase Inhibitor upon Binding. <i>Journal of the American Chemical Society</i> , 2003, 125, 7496-7497.	6.6	84
79	Distortion of a cellobio-derived isofagomine highlights the potential conformational itinerary of inverting β <sup>2</sup> -glucosidases. Electronic supplementary information (ESI) available: details of data and structure quality for complex of cel6A with 1. See <a href="http://www.rsc.org/suppdata/cc/b3/b301592k/">http://www.rsc.org/suppdata/cc/b3/b301592k/</a> . <i>Chemical Communications</i> , 2003, 946-947.	2.2	38
80	Structure of the <i>Humicola insolens</i> cellobiohydrolase Cel6A D416A mutant in complex with a non-hydrolysable substrate analogue, methyl cellobiosyl-4-thio-β <sup>2</sup> -cellobioside, at 1.9 Å. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2002, 58, 2201-2204.	2.5	25
81	Atomic resolution structure of endoglucanase Cel5A in complex with methyl 4,4II,4III,4IV-tetrathio-β <sup>2</sup> -cellopentose highlights the alternative binding modes targeted by substrate mimics. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2001, 57, 1739-1742.	2.5	24
82	Mixed-Linkage Cellooligosaccharides: A New Class of Glycoside Hydrolase Inhibitors. <i>ChemBioChem</i> , 2001, 2, 319-325.	1.3	20
83	Structure and function of <i>Humicola insolens</i> family 6 cellulases: structure of the endoglucanase, Cel6B, at 1.6 Å. <i>Biochemical Journal</i> , 2000, 348, 201.	1.7	24
84	Structure and function of <i>Humicola insolens</i> family 6 cellulases: structure of the endoglucanase, Cel6B, at 1.6 Å. <i>Biochemical Journal</i> , 2000, 348, 201-207.	1.7	61
85	Insights into ligand-induced conformational change in Cel5A from <i>Bacillus agaradhaerens</i> revealed by a catalytically active crystal form. <i>Journal of Molecular Biology</i> , 2000, 297, 819-828.	2.0	46
86	Crystallization and preliminary X-ray analysis of the 6-phospho-β <sup>2</sup> -glucosidase from <i>Bacillus subtilis</i> . <i>Acta Crystallographica Section D: Biological Crystallography</i> , 1999, 55, 1212-1214.	2.5	10
87	Structure of the specificity domain of the Dorsal homologue Gambif1 bound to DNA. <i>Structure</i> , 1999, 7, 841-852.	1.6	26
88	Structural Changes of the Active Site Tunnel of <i>Humicola insolens</i> Cellobiohydrolase, Cel6A, upon Oligosaccharide Binding. <i>Biochemistry</i> , 1999, 38, 8884-8891.	1.2	83
89	Lateral Protonation of a Glycosidase Inhibitor. Structure of the <i>Bacillus agaradhaerens</i> Cel5A in Complex with a Cellobiose-Derived Imidazole at 0.97 Å. <i>Resolution. Journal of the American Chemical Society</i> , 1999, 121, 2621-2622.	6.6	55
90	Crystal structure of the catalytic core domain of the family 6 cellobiohydrolase II, Cel6A, from <i>Humicola insolens</i> , at 1.92 Å. <i>Biochemical Journal</i> , 1999, 337, 297.	1.7	31

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91	Crystal structure of the catalytic core domain of the family 6 cellobiohydrolase II, Cel6A, from <i>Humicola insolens</i> , at 1.92Å... resolution. <i>Biochemical Journal</i> , 1999, 337, 297-304.	1.7	74
92	Snapshots along an Enzymatic Reaction Coordinate:Â Analysis of a Retaining Î <sup>2</sup> -Glycoside Hydrolaseâ€â€i. <i>Biochemistry</i> , 1998, 37, 11707-11713.	1.2	255