

# Michaela Wimmerova

## List of Publications by Year in descending order

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90  
papers

3,972  
citations

136885

32  
h-index

123376

61  
g-index

93  
all docs

93  
docs citations

93  
times ranked

3709  
citing authors

#	ARTICLE	IF	CITATIONS
1	Structures of the lectins from <i>Pseudomonas aeruginosa</i> : insights into the molecular basis for host glycan recognition. <i>Microbes and Infection</i> , 2004, 6, 221-228.	1.0	271
2	Role of LecA and LecB Lectins in <i>Pseudomonas aeruginosa</i> -Induced Lung Injury and Effect of Carbohydrate Ligands. <i>Infection and Immunity</i> , 2009, 77, 2065-2075.	1.0	262
3	Structural basis for oligosaccharide-mediated adhesion of <i>Pseudomonas aeruginosa</i> in the lungs of cystic fibrosis patients. <i>Nature Structural Biology</i> , 2002, 9, 918-921.	9.7	247
4	Structural basis of calcium and galactose recognition by the lectin PA-IL of <i>Pseudomonas aeruginosa</i> . <i>FEBS Letters</i> , 2003, 555, 297-301.	1.3	175
5	Crystal Structure of Fungal Lectin. <i>Journal of Biological Chemistry</i> , 2003, 278, 27059-27067.	1.6	164
6	The Fucose-binding Lectin from <i>Ralstonia solanacearum</i> . <i>Journal of Biological Chemistry</i> , 2005, 280, 27839-27849.	1.6	160
7	Dirigent proteins in plants: modulating cell wall metabolism during abiotic and biotic stress exposure. <i>Journal of Experimental Botany</i> , 2017, 68, 3287-3301.	2.4	159
8	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
9	Structural basis for the interaction between human milk oligosaccharides and the bacterial lectin PA-III of <i>Pseudomonas aeruginosa</i> . <i>Biochemical Journal</i> , 2005, 389, 325-332.	1.7	129
10	Selectivity among Two Lectins: Probing the Effect of Topology, Multivalency and Flexibility of Multivalent Glycoclusters. <i>Chemistry - A European Journal</i> , 2011, 17, 2146-2159.	1.7	108
11	Bambusurilins: a New Family of Macrocyclic Anion Receptors. <i>Organic Letters</i> , 2011, 13, 4000-4003.	2.4	107
12	High affinity fucose binding of <i>Pseudomonas aeruginosa</i> lectin PA-III: 1.0 Å resolution crystal structure of the complex combined with thermodynamics and computational chemistry approaches. <i>Proteins: Structure, Function and Bioinformatics</i> , 2004, 58, 735-746.	1.5	104
13	Rational Design and Synthesis of Optimized Glycoclusters for Multivalent Lectin-Carbohydrate Interactions: Influence of the Linker Arm. <i>Chemistry - A European Journal</i> , 2012, 18, 6250-6263.	1.7	100
14	Binding of different monosaccharides by lectin PA-III from <i>Pseudomonas aeruginosa</i> : Thermodynamics data correlated with X-ray structures. <i>FEBS Letters</i> , 2006, 580, 982-987.	1.3	94
15	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
16	Structural basis of high-affinity glycan recognition by bacterial and fungal lectins. <i>Current Opinion in Structural Biology</i> , 2005, 15, 525-534.	2.6	88
17	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
18	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

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19	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
20	Microbe-focused glycan array screening platform. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 1958-1967.	3.3	71
21	A new <i>Ralstonia solanacearum</i> high-affinity mannose-binding lectin RS-III structurally resembling the <i>Pseudomonas aeruginosa</i> fucose-specific lectin PA-III. <i>Molecular Microbiology</i> , 2004, 52, 691-700.	1.2	70
22	X-ray Structures and Thermodynamics of the Interaction of PA-III from <i>Pseudomonas aeruginosa</i> with Disaccharide Derivatives. <i>ChemMedChem</i> , 2007, 2, 1328-1338.	1.6	61
23	<i>Burkholderia cenocepacia</i> BC2L-C Is a Super Lectin with Dual Specificity and Proinflammatory Activity. <i>PLoS Pathogens</i> , 2011, 7, e1002238.	2.1	61
24	Stacking Interactions between Carbohydrate and Protein Quantified by Combination of Theoretical and Experimental Methods. <i>PLoS ONE</i> , 2012, 7, e46032.	1.1	54
25	Sensitive amperometric biosensor for the determination of biogenic and synthetic amines using pea seedlings amine oxidase: a novel approach for enzyme immobilisation. <i>Biosensors and Bioelectronics</i> , 1999, 14, 695-702.	5.3	48
26	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
27	Anion Binding by Bambus[6]uril Probed in the Gas Phase and in Solution. <i>Journal of Physical Chemistry A</i> , 2011, 115, 11378-11386.	1.1	45
28	FleA Expression in <i>Aspergillus fumigatus</i> Is Recognized by Fucosylated Structures on Mucins and Macrophages to Prevent Lung Infection. <i>PLoS Pathogens</i> , 2016, 12, e1005555.	2.1	44
29	Engineering of PA-III lectin from <i>Pseudomonas aeruginosa</i> – Unravelling the role of the specificity loop for sugar preference. <i>BMC Structural Biology</i> , 2007, 7, 36.	2.3	40
30	Substrate-Assisted Catalytic Mechanism of <i>O</i> -GlcNAc Transferase Discovered by Quantum Mechanics/Molecular Mechanics Investigation. <i>Journal of the American Chemical Society</i> , 2012, 134, 15563-15571.	6.6	39
31	Unusual Entropy-Driven Affinity of <i>Chromobacterium violaceum</i> Lectin CV-III toward Fucose and Mannose. <i>Biochemistry</i> , 2006, 45, 7501-7510.	1.2	36
32	In Silico Mutagenesis and Docking Study of <i>Ralstonia solanacearum</i> RSL Lectin: Performance of Docking Software To Predict Saccharide Binding. <i>Journal of Chemical Information and Modeling</i> , 2012, 52, 1250-1261.	2.5	34
33	<i>Burkholderia cenocepacia</i> lectin A binding to heptoses from the bacterial lipopolysaccharide. <i>Glycobiology</i> , 2012, 22, 1387-1398.	1.3	31
34	Combination of Several Bioinformatics Approaches for the Identification of New Putative Glycosyltransferases in <i>Arabidopsis</i> . <i>Journal of Proteome Research</i> , 2009, 8, 743-753.	1.8	30
35	New sensitive detection method for lectin hemagglutination using microscopy. <i>Microscopy Research and Technique</i> , 2014, 77, 841-849.	1.2	30
36	The CH <sub>2</sub> Interaction in Protein-Carbohydrate Binding: Bioinformatics and In Vitro Quantification. <i>Chemistry - A European Journal</i> , 2020, 26, 10769-10780.	1.7	30

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37	Development and application of a novel recombinant <i>Aleuria aurantia</i> lectin with enhanced core fucose binding for identification of glycoprotein biomarkers of hepatocellular carcinoma. <i>Proteomics</i> , 2016, 16, 3126-3136.	1.3	29
38	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
39	Architecture and Evolution of Blade Assembly in $\beta$ -propeller Lectins. <i>Structure</i> , 2019, 27, 764-775.e3.	1.6	27
40	TRITON: a graphical tool for ligand-binding protein engineering. <i>Bioinformatics</i> , 2008, 24, 1955-1956.	1.8	25
41	Biochemical characterization of broad-specificity enzymes using multivariate experimental design and a colorimetric microplate assay: characterization of the haloalkane dehalogenase mutants. <i>Journal of Microbiological Methods</i> , 2001, 44, 149-157.	0.7	23
42	Combining fold recognition and exploratory data analysis for searching for glycosyltransferases in the genome of <i>Mycobacterium tuberculosis</i> . <i>Biochimie</i> , 2003, 85, 691-700.	1.3	22
43	ValidatorDB: database of up-to-date validation results for ligands and non-standard residues from the Protein Data Bank. <i>Nucleic Acids Research</i> , 2015, 43, D369-D375.	6.5	22
44	<i>In Silico</i> Mutagenesis and Docking Studies of <i>Pseudomonas aeruginosa</i> PA-III Lectin "Predicting Binding Modes and Energies. <i>Journal of Chemical Information and Modeling</i> , 2008, 48, 2234-2242.	2.5	19
45	Terminology of bioanalytical methods (IUPAC Recommendations 2018). <i>Pure and Applied Chemistry</i> , 2018, 90, 1121-1198.	0.9	19
46	Single-Myb-histone proteins from <i>Arabidopsis thaliana</i> : a quantitative study of telomere-binding specificity and kinetics. <i>Biochemical Journal</i> , 2009, 419, 221-230.	1.7	18
47	Synergism of the Two Myb Domains of Tay1 Protein Results in High Affinity Binding to Telomeres. <i>Journal of Biological Chemistry</i> , 2012, 287, 32206-32215.	1.6	18
48	A Novel Fucose-binding Lectin from <i>Photobacterium luminescens</i> (PLL) with an Unusual Heptabladed $\beta$ -Propeller Tetrameric Structure. <i>Journal of Biological Chemistry</i> , 2016, 291, 25032-25049.	1.6	18
49	Characterization of novel bangle lectin from <i>Photobacterium asymbiotica</i> with dual sugar-binding specificity and its effect on host immunity. <i>PLoS Pathogens</i> , 2017, 13, e1006564.	2.1	18
50	Recognition of selected monosaccharides by <i>Pseudomonas aeruginosa</i> Lectin II analyzed by molecular dynamics and free energy calculations. <i>Carbohydrate Research</i> , 2010, 345, 1432-1441.	1.1	17
51	The mink as an animal model for <i>Pseudomonas aeruginosa</i> adhesion: binding of the bacterial lectins (PA-IL and PA-III) to neoglycoproteins and to sections of pancreas and lung tissues from healthy mink. <i>Microbes and Infection</i> , 2007, 9, 566-573.	1.0	16
52	A QM/MM Investigation of the Catalytic Mechanism of Metal-Independent Core 2 $\beta$ -1,6-N-Acetylglucosaminyltransferase. <i>Chemistry - A European Journal</i> , 2013, 19, 8153-8162.	1.7	15
53	Selectivity of original C-hexopyranosyl calix[4]arene conjugates towards lectins of different origin. <i>Carbohydrate Research</i> , 2018, 469, 60-72.	1.1	14
54	Tri- and tetravalent mannoclusters cross-link and aggregate BC2L-A lectin from <i>Burkholderia cenocepacia</i> . <i>Carbohydrate Research</i> , 2017, 437, 1-8.	1.1	12

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55	Investigation of Thermal Denaturation of Barley Nonspecific Lipid Transfer Protein 1 (ns-LTP1b) by Nuclear Magnetic Resonance and Differential Scanning Calorimetry. <i>Journal of Agricultural and Food Chemistry</i> , 2009, 57, 8444-8452.	2.4	11
56	MotiveValidator: interactive web-based validation of ligand and residue structure in biomolecular complexes. <i>Nucleic Acids Research</i> , 2014, 42, W227-W233.	6.5	11
57	Synthesis of Î±-Fucopyranoside-Presenting Glycoclusters and Investigation of Their Interaction with <i>Photobacterium asymbiotica</i> Lectin (PHL). <i>Chemistry - A European Journal</i> , 2018, 24, 4055-4068.	1.7	11
58	Microscopy examination of red blood and yeast cell agglutination induced by bacterial lectins. <i>PLoS ONE</i> , 2019, 14, e0220318.	1.1	11
59	Determination of haloalkane dehalogenase activity by capillary zone electrophoresis. <i>Journal of Chromatography A</i> , 2000, 895, 219-225.	1.8	10
60	Heterologous expression and molecular characterization of the NAD(P)H:acceptor oxidoreductase (FerB) of <i>Paracoccus denitrificans</i> . <i>Protein Expression and Purification</i> , 2009, 68, 233-238.	0.6	10
61	SiteBinder: An Improved Approach for Comparing Multiple Protein Structural Motifs. <i>Journal of Chemical Information and Modeling</i> , 2012, 52, 343-359.	2.5	10
62	Influence of Trp flipping on carbohydrate binding in lectins. An example on <i>Aleuria aurantia</i> lectin AAL. <i>PLoS ONE</i> , 2017, 12, e0189375.	1.1	10
63	Importance of oligomerisation on <i>Pseudomonas aeruginosa</i> Lectin-II binding affinity. In silico and in vitro mutagenesis. <i>Journal of Molecular Modeling</i> , 2009, 15, 673-679.	0.8	9
64	Conformational dynamics are a key factor in signaling mediated by the receiver domain of a sensor histidine kinase from <i>Arabidopsis thaliana</i> . <i>Journal of Biological Chemistry</i> , 2017, 292, 17525-17540.	1.6	9
65	The Five Bacterial Lectins (PA-IL, PA-IIL, RSL, RS-IIL, and CV-IIL): Interactions with Diverse Animal Cells and Glycoproteins. <i>Advances in Experimental Medicine and Biology</i> , 2011, 705, 155-211.	0.8	9
66	X-ray vs. NMR structure of N-terminal domain of Î±-subunit of RNA polymerase. <i>Journal of Structural Biology</i> , 2014, 187, 174-186.	1.3	8
67	Synthesis of Î²-d-galactopyranoside-Presenting Glycoclusters, Investigation of Their Interactions with <i>Pseudomonas aeruginosa</i> Lectin A (PA-IL) and Evaluation of Their Anti-Adhesion Potential. <i>Biomolecules</i> , 2019, 9, 686.	1.8	8
68	Visualization of hydrogen atoms in a perdeuterated lectin-fucose complex reveals key details of protein-carbohydrate interactions. <i>Structure</i> , 2021, 29, 1003-1013.e4.	1.6	8
69	Differential pulse polarographic study of the redox centres in pea amine oxidase. <i>Bioelectrochemistry</i> , 1996, 41, 173-179.	1.0	7
70	Newly identified DNA methyltransferases of <i>Ixodes ricinus</i> ticks. <i>Ticks and Tick-borne Diseases</i> , 2020, 11, 101348.	1.1	7
71	Molecular Modeling of Glycosyltransferases. , 2006, 347, 145-156.		6
72	Investigation of the Binding Affinity of a Broad Array of l-Fucosides with Six Fucose-Specific Lectins of Bacterial and Fungal Origin. <i>Molecules</i> , 2019, 24, 2262.	1.7	6

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73	Plant aminoaldehyde dehydrogenases oxidize a wide range of nitrogenous heterocyclic aldehydes. <i>Amino Acids</i> , 2012, 43, 1189-1202.	1.2	5
74	Fluorescent Cellular Assay for Screening Agents Inhibiting <i>Pseudomonas aeruginosa</i> Adherence. <i>Sensors</i> , 2015, 15, 1945-1953.	2.1	5
75	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
76	Heptabladed $\beta$ -propeller lectins PLL2 and PHL from <i>Photobacterium</i> spp. recognize O-methylated sugars and influence the host immune system. <i>FEBS Journal</i> , 2021, 288, 1343-1365.	2.2	5
77	Fucosylated inhibitors of recently identified bangle lectin from <i>Photobacterium asymbiotica</i> . <i>Scientific Reports</i> , 2019, 9, 14904.	1.6	4
78	Engineering the <i>Pseudomonas aeruginosa</i> II lectin: designing mutants with changed affinity and specificity. <i>Journal of Computer-Aided Molecular Design</i> , 2014, 28, 951-960.	1.3	3
79	Structure and properties of AB21, a novel <i>Agaricus bisporus</i> protein with structural relation to bacterial pore-forming toxins. <i>Proteins: Structure, Function and Bioinformatics</i> , 2018, 86, 897-911.	1.5	3
80	Crystallization and initial X-ray diffraction studies of the flavoenzyme NAD(P)H:(acceptor) oxidoreductase (FerB) from the soil bacterium <i>Paracoccus denitrificans</i> . <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2010, 66, 431-434.	0.7	2
81	Step-By-Step In Vitro Mutagenesis: Lessons From Fucose-Binding Lectin PA-III. <i>Methods in Molecular Biology</i> , 2017, 1498, 399-419.	0.4	2
82	Cytokinin and Ethylene Signaling. , 2018, , 165-200.		2
83	Lectin PLL3, a Novel Monomeric Member of the Seven-Bladed $\beta$ -Propeller Lectin Family. <i>Molecules</i> , 2019, 24, 4540.	1.7	2
84	Synthesis of Tetravalent Thio- and Selenogalactoside-Presenting Galactoclusters and Their Interactions with Bacterial Lectin PA-IL from <i>Pseudomonas aeruginosa</i> . <i>Molecules</i> , 2021, 26, 542.	1.7	2
85	Development of 48-condition buffer screen for protein stability assessment. <i>European Biophysics Journal</i> , 2021, 50, 461-471.	1.2	2
86	Protein engineering study of $\beta$ -mannosidase to set up a potential chemically efficient biocatalyst. <i>Glycobiology</i> , 2014, 24, 1301-1311.	1.3	1
87	Evaluation of anti-PAIIL lectin hen yolk antibody as an agent inhibiting <i>Pseudomonas aeruginosa</i> adherence to epithelial cells. <i>Monatshefte für Chemie</i> , 2016, 147, 889-896.	0.9	1
88	Purification and Some Properties of Isocitrate Dehydrogenase from <i>Paracoccus denitrificans</i> . <i>Preparative Biochemistry and Biotechnology</i> , 2004, 34, 279-289.	1.0	0
89	In Silico Engineering of Proteins That Recognize Small Molecules. , 2012, , .		0
90	Crystallization and preliminary X-ray crystallographic analysis of recombinant $\beta$ -mannosidase from <i>Aspergillus niger</i> . <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2013, 69, 288-291.	0.7	0