

# William C Wimley

## List of Publications by Year in descending order

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106  
papers

13,154  
citations

41258

49  
h-index

30010

103  
g-index

109  
all docs

109  
docs citations

109  
times ranked

11737  
citing authors

#	ARTICLE	IF	CITATIONS
1	Synthetic Molecular Evolution of Cell Penetrating Peptides. <i>Methods in Molecular Biology</i> , 2022, 2383, 73-89.	0.4	3
2	Ebola virus delta peptide is an enterotoxin. <i>Cell Reports</i> , 2022, 38, 110172.	2.9	3
3	Integrated Design of a Membrane-Lytic Peptide-Based Intravenous Nanotherapeutic Suppresses Triple-Negative Breast Cancer. <i>Advanced Science</i> , 2022, 9, e2105506.	5.6	7
4	The Remarkable Innate Resistance of Burkholderia bacteria to Cationic Antimicrobial Peptides: Insights into the Mechanism of AMP Resistance. <i>Journal of Membrane Biology</i> , 2022, , 1.	1.0	5
5	pH-triggered pore-forming peptides with strong composition-dependent membrane selectivity. <i>Biophysical Journal</i> , 2021, 120, 618-630.	0.2	11
6	Tuning of a Membrane-Perforating Antimicrobial Peptide to Selectively Target Membranes of Different Lipid Composition. <i>Journal of Membrane Biology</i> , 2021, 254, 75-96.	1.0	13
7	High glucose induces trafficking of prorenin receptor and stimulates profibrotic factors in the collecting duct. <i>Scientific Reports</i> , 2021, 11, 13815.	1.6	5
8	Inhibition of Streptococcus mutans biofilms with bacterial-derived outer membrane vesicles. <i>BMC Microbiology</i> , 2021, 21, 234.	1.3	18
9	Applications and evolution of melittin, the quintessential membrane active peptide. <i>Biochemical Pharmacology</i> , 2021, 193, 114769.	2.0	45
10	Membrane-selective Nanoscale Pores in Liposomes by a Synthetically Evolved Peptide: Implications for Triggered Release. <i>Nanoscale</i> , 2021, 13, 12185-12197.	2.8	11
11	Rational Modulation of pH-Triggered Macromolecular Poration by Peptide Acylation and Dimerization. <i>Journal of Physical Chemistry B</i> , 2020, 124, 8835-8843.	1.2	3
12	Broad-Spectrum Antiviral Entry Inhibition by Interfacially Active Peptides. <i>Journal of Virology</i> , 2020, 94, .	1.5	16
13	How We Came to Understand the "Tumultuous Chemical Heterogeneity" of the Lipid Bilayer Membrane. <i>Journal of Membrane Biology</i> , 2020, 253, 185-190.	1.0	0
14	Synthetic molecular evolution of host cell-compatible, antimicrobial peptides effective against drug-resistant, biofilm-forming bacteria. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 8437-8448.	3.3	43
15	Crotonylation at serine 46 impairs p53 activity. <i>Biochemical and Biophysical Research Communications</i> , 2020, 524, 730-735.	1.0	19
16	The Mechanism of Membrane Permeabilization by Peptides: Still an Enigma. <i>Australian Journal of Chemistry</i> , 2020, 73, 96.	0.5	34
17	Burkholderia thailandensis outer membrane vesicles exert antimicrobial activity against drug-resistant and competitor microbial species. <i>Journal of Microbiology</i> , 2020, 58, 550-562.	1.3	38
18	Simulation-Guided Rational <i>de Novo</i> Design of a Small Pore-Forming Antimicrobial Peptide. <i>Journal of the American Chemical Society</i> , 2019, 141, 4839-4848.	6.6	80

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19	Mechanism of Action of Peptides That Cause the pH-Triggered Macromolecular Poration of Lipid Bilayers. <i>Journal of the American Chemical Society</i> , 2019, 141, 6706-6718.	6.6	30
20	The Membrane-Active Phytopeptide Cycloviolacin O2 Simultaneously Targets HIV-1-infected Cells and Infectious Viral Particles to Potentiate the Efficacy of Antiretroviral Drugs. <i>Medicines (Basel)</i> , 2019, 10, 1650-1697.	10.0	1650-1697
21	Application of Synthetic Molecular Evolution to the Discovery of Antimicrobial Peptides. <i>Advances in Experimental Medicine and Biology</i> , 2019, 1117, 241-255.	0.8	14
22	Mechanistic Landscape of Membrane-Permeabilizing Peptides. <i>Chemical Reviews</i> , 2019, 119, 6040-6085.	23.0	173
23	How Does Melittin Permeabilize Membranes?. <i>Biophysical Journal</i> , 2018, 114, 251-253.	0.2	30
24	Potent Macromolecule-Sized Poration of Lipid Bilayers by the Macrolittins, A Synthetically Evolved Family of Pore-Forming Peptides. <i>Journal of the American Chemical Society</i> , 2018, 140, 6441-6447.	6.6	41
25	Pituitary adenylate cyclase-activating polypeptide is a potent broad-spectrum antimicrobial peptide: Structure-activity relationships. <i>Peptides</i> , 2018, 104, 35-40.	1.2	48
26	Synthetic molecular evolution of hybrid cell penetrating peptides. <i>Nature Communications</i> , 2018, 9, 2568.	5.8	65
27	Ebola Virus Delta Peptide Is a Viroporin. <i>Journal of Virology</i> , 2017, 91, .	1.5	26
28	pH-Triggered, Macromolecule-Sized Poration of Lipid Bilayers by Synthetically Evolved Peptides. <i>Journal of the American Chemical Society</i> , 2017, 139, 937-945.	6.6	61
29	Spontaneous Membrane Translocating Peptides: The Role of Leucine-Arginine Consensus Motifs. <i>Biophysical Journal</i> , 2017, 113, 835-846.	0.2	42
30	Antimicrobial peptides are degraded by the cytosolic proteases of human erythrocytes. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2017, 1859, 2319-2326.	1.4	65
31	An Outbreak of Ebola Virus Disease in the Lassa Fever Zone. <i>Journal of Infectious Diseases</i> , 2016, 214, S110-S121.	1.9	34
32	Host Cell Interactions Are a Significant Barrier to the Clinical Utility of Peptide Antibiotics. <i>ACS Chemical Biology</i> , 2016, 11, 3391-3399.	1.6	78
33	Pre-Operative Antisepsis Protocol Compliance and the Effect on Bacterial Load Reduction. <i>Surgical Infections</i> , 2016, 17, 32-37.	0.7	7
34	Mechanism Matters: A Taxonomy of Cell Penetrating Peptides. <i>Trends in Biochemical Sciences</i> , 2015, 40, 749-764.	3.7	258
35	Conformational Fine-Tuning of Pore-Forming Peptide Potency and Selectivity. <i>Journal of the American Chemical Society</i> , 2015, 137, 16144-16152.	6.6	53
36	Testing the limits of rational design by engineering pH sensitivity into membrane-active peptides. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2015, 1848, 951-957.	1.4	27

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37	The Cholesterol-dependent Cytolysin Membrane-binding Interface Discriminates Lipid Environments of Cholesterol to Support $\beta$ -Barrel Pore Insertion. <i>Journal of Biological Chemistry</i> , 2015, 290, 17733-17744.	1.6	40
38	Determining the Effects of Membrane-Interacting Peptides on Membrane Integrity. <i>Methods in Molecular Biology</i> , 2015, 1324, 89-106.	0.4	27
39	A lack of synergy between membrane-permeabilizing cationic antimicrobial peptides and conventional antibiotics. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2015, 1848, 8-15.	1.4	37
40	Novel Antiviral Agents: Design, Identification and Characterization of Interfacially Active Peptide Entry inhibitors. <i>FASEB Journal</i> , 2015, 29, 886.20.	0.2	0
41	Making the Membrane Disappear with Spontaneous Membrane Translocating Peptides. <i>FASEB Journal</i> , 2015, 29, 886.15.	0.2	0
42	Toward the de novo design of antimicrobial peptides: Lack of correlation between peptide permeabilization of lipid vesicles and antimicrobial, cytolytic, or cytotoxic activity in living cells. <i>Biopolymers</i> , 2014, 102, 1-6.	1.2	24
43	Highly Efficient Macromolecule-Sized Poration of Lipid Bilayers by a Synthetically Evolved Peptide. <i>Journal of the American Chemical Society</i> , 2014, 136, 4724-4731.	6.6	59
44	Inhibition of Arenavirus Infection by a Glycoprotein-Derived Peptide with a Novel Mechanism. <i>Journal of Virology</i> , 2014, 88, 8556-8564.	1.5	15
45	Structural Plasticity in the Topology of the Membrane-Interacting Domain of HIV-1 gp41. <i>Biophysical Journal</i> , 2014, 106, 610-620.	0.2	22
46	Interfacially active peptides and proteins. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2014, 1838, 2139.	1.4	1
47	Peptide entry inhibitors of enveloped viruses: The importance of interfacial hydrophobicity. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2014, 1838, 2180-2197.	1.4	120
48	HCV Infection Selectively Impairs Type I but Not Type III IFN Signaling. <i>American Journal of Pathology</i> , 2014, 184, 214-229.	1.9	63
49	The electrical response of bilayers to the bee venom toxin melittin: Evidence for transient bilayer permeabilization. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2013, 1828, 1357-1364.	1.4	50
50	A Membrane-Translocating Peptide Penetrates into Bilayers without Significant Bilayer Perturbations. <i>Biophysical Journal</i> , 2013, 104, 2419-2428.	0.2	42
51	Synthetic Molecular Evolution of Pore-Forming Peptides by Iterative Combinatorial Library Screening. <i>ACS Chemical Biology</i> , 2013, 8, 823-831.	1.6	27
52	pH Dependence of Microbe Sterilization by Cationic Antimicrobial Peptides. <i>Antimicrobial Agents and Chemotherapy</i> , 2013, 57, 3312-3320.	1.4	53
53	Direct Cytosolic Delivery of Polar Cargo to Cells by Spontaneous Membrane-translocating Peptides. <i>Journal of Biological Chemistry</i> , 2013, 288, 29974-29986.	1.6	52
54	Interactions of Membrane Active Peptides with Planar Supported Bilayers: An Impedance Spectroscopy Study. <i>Langmuir</i> , 2012, 28, 6088-6096.	1.6	24

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55	TMBB-DB: a transmembrane $\beta$ -barrel proteome database. <i>Bioinformatics</i> , 2012, 28, 2425-2430.	1.8	21
56	Protein folding in membranes. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2012, 1818, 925-926.	1.4	4
57	Gain-of-Function Analogues of the Pore-Forming Peptide Melittin Selected by Orthogonal High-Throughput Screening. <i>Journal of the American Chemical Society</i> , 2012, 134, 12732-12741.	6.6	86
58	Determining the mechanism of membrane permeabilizing peptides: Identification of potent, equilibrium pore-formers. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2012, 1818, 1625-1632.	1.4	57
59	A Highly Charged Voltage-Sensor Helix Spontaneously Translocates across Membranes. <i>Angewandte Chemie - International Edition</i> , 2012, 51, 7150-7153.	7.2	28
60	Release of Dengue Virus Genome Induced by a Peptide Inhibitor. <i>PLoS ONE</i> , 2012, 7, e50995.	1.1	71
61	Spontaneous Membrane-Translocating Peptides by Orthogonal High-Throughput Screening. <i>Journal of the American Chemical Society</i> , 2011, 133, 8995-9004.	6.6	173
62	Structural Plasticity in Self-Assembling Transmembrane $\beta$ -Sheets. <i>Biophysical Journal</i> , 2011, 101, 828-836.	0.2	5
63	The prediction and characterization of YshA, an unknown outer-membrane protein from <i>Salmonella typhimurium</i> . <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2011, 1808, 287-297.	1.4	18
64	High-Throughput Selection of Transmembrane Sequences That Enhance Receptor Tyrosine Kinase Activation. <i>Journal of Molecular Biology</i> , 2011, 412, 43-54.	2.0	26
65	A Look at Arginine in Membranes. <i>Journal of Membrane Biology</i> , 2011, 239, 49-56.	1.0	107
66	Antimicrobial Peptides: Successes, Challenges and Unanswered Questions. <i>Journal of Membrane Biology</i> , 2011, 239, 27-34.	1.0	406
67	FGFR3 Heterodimerization in Achondroplasia, the Most Common Form of Human Dwarfism. <i>Journal of Biological Chemistry</i> , 2011, 286, 13272-13281.	1.6	38
68	Anticancer and chemosensitizing abilities of cycloviolacin O2 from <i>Viola odorata</i> and psyllid cyclotides from <i>Psychotria leptothyrsa</i> . <i>Biopolymers</i> , 2010, 94, 617-625.	1.2	95
69	A highly accurate statistical approach for the prediction of transmembrane $\beta$ -barrels. <i>Bioinformatics</i> , 2010, 26, 1965-1974.	1.8	52
70	Energetics of Peptide and Protein Binding to Lipid Membranes. <i>Advances in Experimental Medicine and Biology</i> , 2010, 677, 14-23.	0.8	10
71	High-throughput discovery of broad-spectrum peptide antibiotics. <i>FASEB Journal</i> , 2010, 24, 3232-3238.	0.2	56
72	Describing the Mechanism of Antimicrobial Peptide Action with the Interfacial Activity Model. <i>ACS Chemical Biology</i> , 2010, 5, 905-917.	1.6	786

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73	Toward genomic identification of $\beta$ -barrel membrane proteins: Composition and architecture of known structures. <i>Protein Science</i> , 2009, 11, 301-312.	3.1	199
74	Broad-Spectrum Antimicrobial Peptides by Rational Combinatorial Design and High-Throughput Screening: The Importance of Interfacial Activity. <i>Journal of the American Chemical Society</i> , 2009, 131, 7609-7617.	6.6	262
75	Protein Folding in Membranes: Insights from Neutron Diffraction Studies of a Membrane $\beta$ -Sheet Oligomer. <i>Biophysical Journal</i> , 2008, 94, 492-505.	0.2	15
76	Characterization of antimicrobial peptide activity by electrochemical impedance spectroscopy. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2008, 1778, 2430-2436.	1.4	46
77	Biomolecular Engineering by Combinatorial Design and High-Throughput Screening: Small, Soluble Peptides That Permeabilize Membranes. <i>Journal of the American Chemical Society</i> , 2008, 130, 9849-9858.	6.6	125
78	Viroporin potential of the lentivirus lytic peptide (LLP) domains of the HIV-1 gp41 protein. <i>Virology Journal</i> , 2007, 4, 123.	1.4	33
79	$\beta$ -Sheet Pore-Forming Peptides Selected from a Rational Combinatorial Library: Mechanism of Pore Formation in Lipid Vesicles and Activity in Biological Membranes. <i>Biochemistry</i> , 2007, 46, 12124-12139.	1.2	72
80	Inhibition of severe acute respiratory syndrome-associated coronavirus (SARS-CoV) infectivity by peptides analogous to the viral spike protein. <i>Virus Research</i> , 2006, 120, 146-155.	1.1	66
81	The ERBB4/HER4 Intracellular Domain 4ICD Is a BH3-Only Protein Promoting Apoptosis of Breast Cancer Cells. <i>Cancer Research</i> , 2006, 66, 6412-6420.	0.4	189
82	Identification and Characterization of the Putative Fusion Peptide of the Severe Acute Respiratory Syndrome-Associated Coronavirus Spike Protein. <i>Journal of Virology</i> , 2005, 79, 7195-7206.	1.5	126
83	Rational combinatorial design of pore-forming $\beta$ -sheet peptides. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 10511-10515.	3.3	66
84	The Aromatic Domain of the Coronavirus Class I Viral Fusion Protein Induces Membrane Permeabilization: Putative Role during Viral Entry. <i>Biochemistry</i> , 2005, 44, 947-958.	1.2	58
85	Enhancing the Therapeutic Potential of an Anti-Leukemic Peptide. <i>Blood</i> , 2005, 106, 245-245.	0.6	0
86	Reversible Unfolding of $\beta$ -Sheets in Membranes: A Calorimetric Study. <i>Journal of Molecular Biology</i> , 2004, 342, 703-711.	2.0	33
87	The versatile $\beta$ -barrel membrane protein. <i>Current Opinion in Structural Biology</i> , 2003, 13, 404-411.	2.6	395
88	Folding of $\beta$ -sheets in membranes: specificity and promiscuity in peptide model systems. <i>Journal of Molecular Biology</i> , 2001, 309, 975-988.	2.0	51
89	A High-Throughput Screen for Identifying Transmembrane Pore-Forming Peptides. <i>Analytical Biochemistry</i> , 2001, 293, 258-263.	1.1	40
90	Designing Transmembrane $\beta$ -Helices That Insert Spontaneously. <i>Biochemistry</i> , 2000, 39, 4432-4442.	1.2	137

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91	Determining the Membrane Topology of Peptides by Fluorescence Quenching. <i>Biochemistry</i> , 2000, 39, 161-170.	1.2	80
92	MEMBRANE PROTEIN FOLDING AND STABILITY: Physical Principles. <i>Annual Review of Biophysics and Biomolecular Structure</i> , 1999, 28, 319-365.	18.3	1,595
93	An amphipathic $\alpha$ -helix at a membrane interface: a structural study using a novel X-ray diffraction method 1 Edited by D. C. Rees. <i>Journal of Molecular Biology</i> , 1999, 290, 99-117.	2.0	196
94	Hydrophobic interactions of peptides with membrane interfaces. <i>BBA - Biomembranes</i> , 1998, 1376, 339-352.	7.9	482
95	[4] Protein folding in membranes: Determining energetics of peptide-bilayer interactions. <i>Methods in Enzymology</i> , 1998, 295, 62-87.	0.4	233
96	Folding of $\beta$ -sheet membrane proteins: a hydrophobic hexapeptide model. <i>Journal of Molecular Biology</i> , 1998, 277, 1091-1110.	2.0	195
97	The Preference of Tryptophan for Membrane Interfaces. <i>Biochemistry</i> , 1998, 37, 14713-14718.	1.2	899
98	[23] Mechanism of leakage of contents of membrane vesicles determined by fluorescence reuquenching. <i>Methods in Enzymology</i> , 1997, 278, 474-486.	0.4	56
99	Solvation Energies of Amino Acid Side Chains and Backbone in a Family of Host-Guest Pentapeptides. <i>Biochemistry</i> , 1996, 35, 5109-5124.	1.2	534
100	Experimentally determined hydrophobicity scale for proteins at membrane interfaces. <i>Nature Structural and Molecular Biology</i> , 1996, 3, 842-848.	3.6	1,525
101	Structure, function, and membrane integration of defensins. <i>Current Opinion in Structural Biology</i> , 1995, 5, 521-527.	2.6	392
102	Interactions between human defensins and lipid bilayers: Evidence for formation of multimeric pores. <i>Protein Science</i> , 1994, 3, 1362-1373.	3.1	349
103	Peptides in lipid bilayers: structural and thermodynamic basis for partitioning and folding. <i>Current Opinion in Structural Biology</i> , 1994, 4, 79-86.	2.6	182
104	Membrane partitioning: Distinguishing bilayer effects from the hydrophobic effect. <i>Biochemistry</i> , 1993, 32, 6307-6312.	1.2	209
105	Transbilayer and interbilayer phospholipid exchange in dimyristoylphosphatidylcholine/dimyristoylphosphatidylethanolamine large unilamellar vesicles. <i>Biochemistry</i> , 1991, 30, 1702-1709.	1.2	76
106	Exchange and flip-flop of dimyristoyl phosphatidylcholine in liquid-crystalline, gel and two-component, two-phase large unilamellar vesicles. <i>Biochemistry</i> , 1990, 29, 1296-1303.	1.2	116