

Yechezkel Shai

List of Publications by Year in descending order

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53
papers

4,583
citations

172457

29
h-index

206112

48
g-index

54
all docs

54
docs citations

54
times ranked

5381
citing authors

#	ARTICLE	IF	CITATIONS
1	Mode of action of membrane active antimicrobial peptides. <i>Biopolymers</i> , 2002, 66, 236-248.	2.4	1,374
2	Mode of action of linear amphipathic α -helical antimicrobial peptides. <i>Biopolymers</i> , 1998, 47, 451-463.	2.4	729
3	Structure and organization of the human antimicrobial peptide LL-37 in phospholipid membranes: relevance to the molecular basis for its non-cell-selective activity. <i>Biochemical Journal</i> , 1999, 341, 501-513.	3.7	494
4	From Innate Immunity to de-Novo Designed Antimicrobial Peptides. <i>Current Pharmaceutical Design</i> , 2002, 8, 715-725.	1.9	128
5	Cyclization of a Cytolytic Amphipathic α -Helical Peptide and Its Diastereomer: Effect on Structure, Interaction with Model Membranes, and Biological Function. <i>Biochemistry</i> , 2000, 39, 6103-6114.	2.5	119
6	Mechanisms of biofilm inhibition and degradation by antimicrobial peptides. <i>Biochemical Journal</i> , 2015, 468, 259-270.	3.7	116
7	Transmembrane domains interactions within the membrane milieu: Principles, advances and challenges. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2012, 1818, 974-983.	2.6	112
8	Specificity in Transmembrane Helix-Helix Interactions Mediated by Aromatic Residues. <i>Journal of Biological Chemistry</i> , 2007, 282, 19753-19761.	3.4	101
9	Regulation of innate immune responses by transmembrane interactions: Lessons from the TLR family. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2014, 1838, 1586-1593.	2.6	84
10	Structure and Organization of Hemolytic and Nonhemolytic Diastereomers of Antimicrobial Peptides in Membranes. <i>Biochemistry</i> , 1999, 38, 16963-16973.	2.5	82
11	The Effect of Cyclization of Magainin 2 and Melittin Analogues on Structure, Function, and Model Membrane Interactions: A Implication to Their Mode of Action. <i>Biochemistry</i> , 2001, 40, 6388-6397.	2.5	80
12	Effect of Multiple Aliphatic Amino Acids Substitutions on the Structure, Function, and Mode of Action of Diastereomeric Membrane Active Peptides. <i>Biochemistry</i> , 2001, 40, 12591-12603.	2.5	78
13	Defensive remodeling: How bacterial surface properties and biofilm formation promote resistance to antimicrobial peptides. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2015, 1848, 3089-3100.	2.6	73
14	Host Defense Peptides and Lipopeptides: Modes of Action and Potential Candidates for the Treatment of Bacterial and Fungal Infections. <i>Current Protein and Peptide Science</i> , 2006, 7, 479-486.	1.4	72
15	Rapid entry of bitter and sweet tastants into liposomes and taste cells: implications for signal transduction. <i>American Journal of Physiology - Cell Physiology</i> , 2000, 278, C17-C25.	4.6	71
16	d-Amino acids incorporation in the frog skin-derived peptide esculentin-1a(1-21)NH ₂ is beneficial for its multiple functions. <i>Amino Acids</i> , 2015, 47, 2505-2519.	2.7	70
17	Multiparametric AFM reveals turgor-responsive net-like peptidoglycan architecture in live streptococci. <i>Nature Communications</i> , 2015, 6, 7193.	12.8	60
18	HIV-1 fusion peptide targets the TCR and inhibits antigen-specific T cell activation. <i>Journal of Clinical Investigation</i> , 2005, 115, 2149-2158.	8.2	60

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19	The Identification of a Minimal Dimerization Motif QXXS That Enables Homo- and Hetero-association of Transmembrane Helices in Vivo. <i>Journal of Biological Chemistry</i> , 2005, 280, 27449-27457.	3.4	52
20	Androctonin, a hydrophilic disulphide-bridged non-haemolytic anti-microbial peptide: a plausible mode of action. <i>Biochemical Journal</i> , 2000, 345, 653-664.	3.7	51
21	Esculentin-1a-Derived Peptides Promote Clearance of <i>Pseudomonas aeruginosa</i> Internalized in Bronchial Cells of Cystic Fibrosis Patients and Lung Cell Migration: Biochemical Properties and a Plausible Mode of Action. <i>Antimicrobial Agents and Chemotherapy</i> , 2016, 60, 7252-7262.	3.2	47
22	T-Cell inactivation and immunosuppressive activity induced by HIV gp41 via novel interacting motif. <i>FASEB Journal</i> , 2007, 21, 393-401.	0.5	40
23	Characterization of the HIV N-terminal Fusion Peptide-containing Region in Context of Key gp41 Fusion Conformations. <i>Journal of Biological Chemistry</i> , 2006, 281, 21755-21762.	3.4	37
24	Sendai Virus Internal Fusion Peptide: Structural and Functional Characterization and a Plausible Mode of Viral Entry Inhibition. <i>Biochemistry</i> , 2000, 39, 11581-11592.	2.5	35
25	The HIV-1 Envelope Transmembrane Domain Binds TLR2 through a Distinct Dimerization Motif and Inhibits TLR2-Mediated Responses. <i>PLoS Pathogens</i> , 2014, 10, e1004248.	4.7	33
26	Participation of Two Fusion Peptides in Measles Virus-Induced Membrane Fusion: Emerging Similarity with Other Paramyxoviruses. <i>Biochemistry</i> , 2001, 40, 1340-1349.	2.5	32
27	Hetero-assembly Between All-l- and All-d-Amino Acid Transmembrane Domains: Forces Involved and Implication for Inactivation of Membrane Proteins. <i>Journal of Molecular Biology</i> , 2004, 344, 855-864.	4.2	31
28	ATR-FTIR studies in pore forming and membrane induced fusion peptides. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2013, 1828, 2306-2313.	2.6	31
29	Neutralization of pro-inflammatory monocytes by targeting TLR2 dimerization ameliorates colitis. <i>EMBO Journal</i> , 2016, 35, 685-698.	7.8	30
30	HIV-1 gp41 and TCR Trans-Membrane Domains Share a Motif Exploited by the HIV Virus to Modulate T-Cell Proliferation. <i>PLoS Pathogens</i> , 2010, 6, e1001085.	4.7	28
31	Chirality-independent Protein-Protein Recognition Between Transmembrane Domains in Vivo. <i>Journal of Molecular Biology</i> , 2002, 322, 491-495.	4.2	26
32	Biohybrid Polymer-Antimicrobial Peptide Medium against <i>Enterococcus faecalis</i> . <i>PLoS ONE</i> , 2014, 9, e109413.	2.5	24
33	HIV-1 fusion protein exerts complex immunosuppressive effects. <i>Trends in Biochemical Sciences</i> , 2013, 38, 345-349.	7.5	18
34	A GxxxG-like Motif within HIV-1 Fusion Peptide Is Critical to Its Immunosuppressant Activity, Structure, and Interaction with the Transmembrane Domain of the T-cell Receptor. <i>Journal of Biological Chemistry</i> , 2012, 287, 33503-33511.	3.4	16
35	NMR structure and binding of esculentin-1a (1-21)NH ₂ and its diastereomer to lipopolysaccharide: Correlation with biological functions. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2016, 1858, 800-812.	2.6	16
36	Mode of Membrane Interaction and Fusogenic Properties of a de Novo Transmembrane Model Peptide Depend on the Length of the Hydrophobic Core. <i>Journal of Biological Chemistry</i> , 2007, 282, 18388-18396.	3.4	15

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37	Functional Domains within Fusion Proteins: Prospectives for Development of Peptide Inhibitors of Viral Cell Fusion. <i>Bioscience Reports</i> , 2000, 20, 535-555.	2.4	14
38	The Transmembrane Domain of HIV-1 gp41 Inhibits T-Cell Activation by Targeting Multiple T-Cell Receptor Complex Components through Its GxxxG Motif. <i>Biochemistry</i> , 2016, 55, 1049-1057.	2.5	14
39	Early and late HIV-1 membrane fusion events are impaired by sphinganine lipidated peptides that target the fusion site. <i>Biochemical Journal</i> , 2014, 461, 213-222.	3.7	13
40	Bacterial resistance to antimicrobial peptides. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2015, 1848, 3019-3020.	2.6	13
41	Sustained Release of Antibacterial Lipopeptides from Biodegradable Polymers against Oral Pathogens. <i>PLoS ONE</i> , 2016, 11, e0162537.	2.5	10
42	Methods for Investigating Biofilm Inhibition and Degradation by Antimicrobial Peptides. <i>Methods in Molecular Biology</i> , 2017, 1548, 309-322.	0.9	10
43	Why <i>Bacillus thuringiensis</i> insecticidal toxins are so effective: unique features of their mode of action. <i>FEMS Microbiology Letters</i> , 2001, 195, 1-8.	1.8	8
44	Interfering with the Dimerization of the ErbB Receptors by Transmembrane Domain-Derived Peptides Inhibits Tumorigenic Growth in Vitro and in Vivo. <i>Biochemistry</i> , 2016, 55, 5520-5530.	2.5	7
45	The HIV gp41 pocket binding domain enables C-terminal heptad repeat transition from mediating membrane fusion to immune modulation. <i>Biochemical Journal</i> , 2016, 473, 911-918.	3.7	7
46	Mode of action of linear amphipathic α -helical antimicrobial peptides. <i>Biopolymers</i> , 1998, 47, 451-463.	2.4	7
47	An Immunomodulating Motif of the HIV-1 Fusion Protein Is Chirality-independent. <i>Journal of Biological Chemistry</i> , 2013, 288, 32852-32860.	3.4	6
48	Mode of action of linear amphipathic α -helical antimicrobial peptides. , 1998, 47, 451.		4
49	Proline localized to the interaction interface can mediate self-association of transmembrane domains. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2014, 1838, 2313-2318.	2.6	3
50	Mechanism of Membrane Permeation and Pore Formation by Antimicrobial Peptides. , 2006, , 187-217.		2
51	Response to "Interaction between HIV gp41 fusion peptide and T cell receptor: putting the puzzle pieces back together" FASEB Journal, 2007, 21, 1635-1635.	0.5	0
52	Kinetics and Thermodynamics of the Microgene Polymerization Reaction. <i>Biophysical Journal</i> , 2009, 96, 1673-1674.	0.5	0
53	The HTLV-1 gp21 fusion peptide inhibits antigen specific T-cell activation in-vitro and in mice. <i>PLoS Pathogens</i> , 2018, 14, e1007044.	4.7	0