Gaurav Jerath

List of Publications by Year in descending order

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14	228	1040056	996975
papers	citations	h-index	g-index
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15	15	15	127
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Geometry encoded functional programming of tumor homing peptides for targeted drug delivery. Journal of Controlled Release, 2021, 333, 16-27.	9.9	34
2	Effect of tacticity-derived topological constraints in bactericidal peptides. Biochimica Et Biophysica Acta - Biomembranes, 2017, 1859, 1388-1395.	2.6	30
3	Syndiotactic peptides for targeted delivery. Acta Biomaterialia, 2019, 87, 130-139.	8.3	30
4	Topological effects on the designability and bactericidal potency of antimicrobial peptides. Biophysical Chemistry, 2019, 248, 1-8.	2.8	27
5	Peptide-based delivery vectors with pre-defined geometrical locks. RSC Medicinal Chemistry, 2020, 11, 1303-1313.	3.9	18
6	Peptido-mimetic Approach in the Design of Syndiotactic Antimicrobial Peptides. International Journal of Peptide Research and Therapeutics, 2018, 24, 299-307.	1.9	16
7	Molecular hybridization combining tumor homing and penetrating peptide domains for cellular targeting. Drug Delivery and Translational Research, 2022, 12, 1285-1292.	5.8	16
8	Mapping drug-target interactions and synergy in multi-molecular therapeutics for pressure-overload cardiac hypertrophy. Npj Systems Biology and Applications, 2021, 7, 11.	3.0	13
9	Delivery of Small Molecules by Syndiotactic Peptides for Breast Cancer Therapy. Molecular Pharmaceutics, 2022, 19, 2877-2887.	4.6	11
10	Conformationally constrained peptides for drug delivery. Journal of Peptide Science, 2020, 26, e3244.	1.4	10
11	Characterization of ICAM-1 biophore to design cytoadherence blocking peptides. Journal of Molecular Graphics and Modelling, 2015, 57, 27-35.	2.4	8
12	Mapping the Geometric Evolution of Protein Folding Motor. PLoS ONE, 2016, 11, e0163993.	2.5	7
13	bPE toolkit: toolkit for computational protein engineering. Systems and Synthetic Biology, 2014, 8, 337-341.	1.0	6
14	Structureâ€based barcoding of proteins. Protein Science, 2014, 23, 117-120.	7.6	1