Spyridoula Karamanou

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Bcl-xL acts as an inhibitor of IP3R channels, thereby antagonizing Ca2+-driven apoptosis. Cell Death and Differentiation, 2022, 29, 788-805.	5.0	41
2	Preproteins couple the intrinsic dynamics of SecA to its ATPase cycle to translocate via a catch and release mechanism. Cell Reports, 2022, 38, 110346.	2.9	5
3	Effective Small Molecule Antibacterials from a Novel Anti-Protein Secretion Screen. Microorganisms, 2021, 9, 592.	1.6	1
4	A nexus of intrinsic dynamics underlies translocase priming. Structure, 2021, 29, 846-858.e7.	1.6	17
5	Probing Universal Protein Dynamics Using Hydrogen–Deuterium Exchange Mass Spectrometry-Derived Residue-Level Gibbs Free Energy. Analytical Chemistry, 2021, 93, 12840-12847.	3.2	16
6	Structural Dynamics of the Functional Nonameric Type III Translocase Export Gate. Journal of Molecular Biology, 2021, 433, 167188.	2.0	7
7	Structural dynamics in the evolution of a bilobed protein scaffold. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	9
8	Trigger factor is a <i>bona fide</i> secretory pathway chaperone that interacts with SecB and the translocase. EMBO Reports, 2020, 21, e49054.	2.0	30
9	A Reporter System for Fast Quantitative Monitoring of Type 3 Protein Secretion in Enteropathogenic E. coli. Microorganisms, 2020, 8, 1786.	1.6	3
10	BDA-366, a putative Bcl-2 BH4 domain antagonist, induces apoptosis independently of Bcl-2 in a variety of cancer cell models. Cell Death and Disease, 2020, 11, 769.	2.7	15
11	Structural Basis of the Subcellular Topology Landscape of Escherichia coli. Frontiers in Microbiology, 2019, 10, 1670.	1.5	25
12	Inner Membrane Translocases and Insertases. Sub-Cellular Biochemistry, 2019, 92, 337-366.	1.0	14
13	Protein Transport Across the Bacterial Plasma Membrane by the Sec Pathway. Protein Journal, 2019, 38, 262-273.	0.7	30
14	The Preprotein Binding Domain of SecA Displays Intrinsic Rotational Dynamics. Structure, 2019, 27, 90-101.e6.	1.6	12
15	Secretome Dynamics in a Gram-Positive Bacterial Model. Molecular and Cellular Proteomics, 2019, 18, 423-436.	2.5	12
16	A double point mutation at residues lle14 and Val15 of Bclâ€2 uncovers a role for the BH4 domain in both protein stability and function. FEBS Journal, 2018, 285, 127-145.	2.2	16
17	Long-Lived Folding Intermediates Predominate the Targeting-Competent Secretome. Structure, 2018, 26, 695-707.e5.	1.6	44
18	Monitoring Protein Secretion in Streptomyces Using Fluorescent Proteins. Frontiers in Microbiology, 2018. 9. 3019.	1.5	11

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19	Transcriptomic and fluxomic changes in Streptomyces lividans producing heterologous protein. Microbial Cell Factories, 2018, 17, 198.	1.9	18
20	Streptomyces protein secretion and its application in biotechnology. FEMS Microbiology Letters, 2018, 365, .	0.7	22
21	Editorial: Thematic issue on bacterial protein export: from fundamentals to applications. FEMS Microbiology Letters, 2018, 365, .	0.7	0
22	Optimization of type 3 protein secretion in enteropathogenic Escherichia coli. FEMS Microbiology Letters, 2018, 365, .	0.7	5
23	Identification of influenza PA-Nter endonuclease inhibitors using pharmacophore- and docking-based virtual screening. Bioorganic and Medicinal Chemistry, 2018, 26, 4544-4550.	1.4	9
24	Multi-Omics and Targeted Approaches to Determine the Role of Cellular Proteases in Streptomyces Protein Secretion. Frontiers in Microbiology, 2018, 9, 1174.	1.5	29
25	Comprehensive subcellular topologies of polypeptides in Streptomyces. Microbial Cell Factories, 2018, 17, 43.	1.9	19
26	Preprotein mature domains contain translocase targeting signals that are essential for secretion. Journal of Cell Biology, 2017, 216, 1357-1369.	2.3	67
27	Preprotein Conformational Dynamics Drive Bivalent Translocase Docking and Secretion. Structure, 2017, 25, 1056-1067.e6.	1.6	28
28	Hierarchical protein targeting and secretion is controlled by an affinity switch in the type <scp>lll</scp> secretion system of enteropathogenic <i>Escherichia coli</i> . EMBO Journal, 2017, 36, 3517-3531.	3.5	54
29	Protein export through the bacterial Sec pathway. Nature Reviews Microbiology, 2017, 15, 21-36.	13.6	332
30	Large-scale production of a thermostable Rhodothermus marinus cellulase by heterologous secretion from Streptomyces lividans. Microbial Cell Factories, 2017, 16, 232.	1.9	40
31	Rapid labelâ€free quantitative analysis of the <i>E. coli</i> BL21(DE3) inner membrane proteome. Proteomics, 2016, 16, 85-97.	1.3	24
32	A polysulfobetaine hydrogel for immobilization of a glucose-binding protein. RSC Advances, 2016, 6, 83890-83900.	1.7	11
33	Protein folding in the cell envelope of Escherichia coli. Nature Microbiology, 2016, 1, 16107.	5.9	75
34	SecA-mediated targeting and translocation of secretory proteins. Biochimica Et Biophysica Acta - Molecular Cell Research, 2014, 1843, 1466-1474.	1.9	76
35	Quaternary Dynamics of the SecA Motor Drive Translocase Catalysis. Molecular Cell, 2013, 52, 655-666.	4.5	51
36	Breaking on through to the other side: protein export through the bacterial Sec system. Biochemical Journal, 2013, 449, 25-37.	1.7	64

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37	The Escherichia coli Peripheral Inner Membrane Proteome. Molecular and Cellular Proteomics, 2013, 12, 599-610.	2.5	79
38	Quantitative analysis of energy transfer between fluorescent proteins in CFP–GBP–YFP and its response to Ca2+. Physical Chemistry Chemical Physics, 2011, 13, 17852.	1.3	3
39	In Vitro Assays to Analyze Translocation of the Model Secretory Preprotein Alkaline Phosphatase. Methods in Molecular Biology, 2010, 619, 157-172.	0.4	22
40	Signal peptides are allosteric activators of the protein translocase. Nature, 2009, 462, 363-367.	13.7	125
41	Assembly of the translocase motor onto the preproteinâ€conducting channel. Molecular Microbiology, 2008, 70, 311-322.	1.2	26
42	The P. CÉZANNE Project: Innovative Approaches to Continuous Glucose Monitoring. Annual International Conference of the IEEE Engineering in Medicine and Biology Society, 2007, 2007, 6061-4.	0.5	4
43	Structural Basis for Signal-Sequence Recognition by the Translocase Motor SecA as Determined by NMR. Cell, 2007, 131, 756-769.	13.5	381
44	Bacterial protein secretion through the translocase nanomachine. Nature Reviews Microbiology, 2007, 5, 839-851.	13.6	210
45	Preprotein-controlled catalysis in the helicase motor of SecA. EMBO Journal, 2007, 26, 2904-2914.	3.5	56
46	Functional large-scale production of a novel Jonesia sp. xyloglucanase by heterologous secretion from Streptomyces lividans. Journal of Biotechnology, 2006, 121, 498-507.	1.9	54
47	Cloning, purification and characterization of a functional anthracycline glycosyltransferase. Journal of Biotechnology, 2006, 125, 425-433.	1.9	10
48	Disorder-order folding transitions underlie catalysis in the helicase motor of SecA. Nature Structural and Molecular Biology, 2006, 13, 594-602.	3.6	73
49	Identification of the Preprotein Binding Domain of SecA. Journal of Biological Chemistry, 2005, 280, 43209-43217.	1.6	76
50	Escherichia coliSecA truncated at its termini is functional and dimeric. FEBS Letters, 2005, 579, 1267-1271.	1.3	39
51	Purification of a functional mature region from a SecA-dependent preprotein. Protein Expression and Purification, 2005, 40, 336-339.	0.6	6
52	Global Co-ordination of Protein Translocation by the SecA IRA1 Switch. Journal of Biological Chemistry, 2004, 279, 22490-22497.	1.6	47
53	Helicase Motif III in SecA is essential for coupling preprotein binding to translocation ATPase. EMBO Reports, 2004, 5, 807-811.	2.0	24
54	Type III Protein Translocase. Journal of Biological Chemistry, 2003, 278, 25816-25824.	1.6	61

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55	Allosteric Communication between Signal Peptides and the SecA Protein DEAD Motor ATPase Domain. Journal of Biological Chemistry, 2002, 277, 13724-13731.	1.6	49
56	RPL29 codes for a non-essential protein of the 60S ribosomal subunit in Saccharomyces cerevisiae and exhibits synthetic lethality with mutations in genes for proteins required for subunit coupling. Biochimica Et Biophysica Acta Gene Regulatory Mechanisms, 2002, 1574, 255-261.	2.4	34
57	A molecular switch in SecA protein couples ATP hydrolysis to protein translocation. Molecular Microbiology, 1999, 34, 1133-1145.	1.2	124
58	QSR1, an Essential Yeast Gene with a Genetic Relationship to a Subunit of the Mitochondrial Cytochromebc 1 Complex, Codes for a 60 S Ribosomal Subunit Protein. Journal of Biological Chemistry, 1997, 272, 13372-13379.	1.6	75
59	Allosteric cross-talk between the hydrophobic cleft and the BH4 domain of Bcl-2 in control of inositol 1,4,5-trisphosphate receptor activity. Exploration of Targeted Anti-tumor Therapy, 0, , 375-391.	0.5	1