

Maria Ciaramella

List of Publications by Year in descending order

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56
papers

1,790
citations

279798

23
h-index

289244

40
g-index

57
all docs

57
docs citations

57
times ranked

1677
citing authors

#	ARTICLE	IF	CITATIONS
1	Crystal structure of the Î²-glycosidase from the hyperthermophilic archeon <i>Sulfolobus solfataricus</i> : resilience as a key factor in thermostability. <i>Journal of Molecular Biology</i> , 1997, 271, 789-802.	4.2	235
2	Genomic remnants of alpha-globin genes in the hemoglobinless antarctic icefishes.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1995, 92, 1817-1821.	7.1	162
3	Restoration of the Activity of Active-Site Mutants of the Hyperthermophilic Î²-Glycosidase from <i>Sulfolobus solfataricus</i> : A Dependence of the Mechanism on the Action of External Nucleophiles. <i>Biochemistry</i> , 1998, 37, 17262-17270.	2.5	110
4	An Lrp-Like Protein of the Hyperthermophilic Archaeon <i>Sulfolobus solfataricus</i> Which Binds to Its Own Promoter. <i>Journal of Bacteriology</i> , 1999, 181, 1474-1480.	2.2	75
5	RNA topoisomerase is prevalent in all domains of life and associates with polyribosomes in animals. <i>Nucleic Acids Research</i> , 2016, 44, 6335-6349.	14.5	63
6	DNA bending, compaction and negative supercoiling by the architectural protein Sso7d of <i>Sulfolobus solfataricus</i> . <i>Nucleic Acids Research</i> , 2002, 30, 2656-2662.	14.5	57
7	Identification of two glutamic acid residues essential for catalysis in the Î²-glycosidase from the thermoacidophilic archaeon <i>Sulfolobus solfataricus</i> . <i>Protein Engineering, Design and Selection</i> , 1996, 9, 1191-1195.	2.1	50
8	Another extreme genome: how to live at pH 0. <i>Trends in Microbiology</i> , 2005, 13, 49-51.	7.7	47
9	Annealing of complementary DNA strands above the melting point of the duplex promoted by an archaeal protein. <i>Journal of Molecular Biology</i> , 1997, 267, 841-848.	4.2	46
10	Reverse Gyrase Recruitment to DNA after UV Light Irradiation in <i>Sulfolobus solfataricus</i> . <i>Journal of Biological Chemistry</i> , 2004, 279, 33192-33198.	3.4	46
11	Reverse gyrase and genome stability in hyperthermophilic organisms. <i>Biochemical Society Transactions</i> , 2009, 37, 69-73.	3.4	41
12	Selective degradation of reverse gyrase and DNA fragmentation induced by alkylating agent in the archaeon <i>Sulfolobus solfataricus</i> . <i>Nucleic Acids Research</i> , 2006, 34, 2098-2108.	14.5	38
13	Activity and Regulation of Archaeal DNA Alkyltransferase. <i>Journal of Biological Chemistry</i> , 2012, 287, 4222-4231.	3.4	37
14	Thermostable Î²-Glycosidase from <i>Sulfolobus Solfataricus</i> . <i>Biocatalysis</i> , 1994, 11, 89-103.	0.9	34
15	Transcriptional response to DNA damage in the archaeon <i>Sulfolobus solfataricus</i> . <i>Nucleic Acids Research</i> , 2003, 31, 6127-6138.	14.5	33
16	Molecular biology of extremophiles. <i>World Journal of Microbiology and Biotechnology</i> , 1995, 11, 71-84.	3.6	32
17	Activity and stability of hyperthermophilic enzymes: a comparative study on two archaeal Î²-glycosidases. <i>Extremophiles</i> , 2000, 4, 157-164.	2.3	32
18	Do the hemoglobinless icefishes have globin genes?. <i>Comparative Biochemistry and Physiology A, Comparative Physiology</i> , 1997, 118, 1027-1030.	0.6	29

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19	Biochemical and Structural Studies of the Mycobacterium tuberculosis <i>oxy-Methylguanine Methyltransferase</i> and Mutated Variants. <i>Journal of Bacteriology</i> , 2013, 195, 2728-2736.	2.2	29
20	A Novel Member of the Bacterial-Archaeal Regulator Family Is a Nonspecific DNA-binding Protein and Induces Positive Supercoiling. <i>Journal of Biological Chemistry</i> , 2001, 276, 10745-10752.	3.4	27
21	Dissection of reverse gyrase activities: insight into the evolution of a thermostable molecular machine. <i>Nucleic Acids Research</i> , 2008, 36, 4587-4597.	14.5	26
22	Structure-function relationships governing activity and stability of a DNA alkylation damage repair thermostable protein. <i>Nucleic Acids Research</i> , 2015, 43, 8801-8816.	14.5	26
23	Functional interaction of reverse gyrase with single-strand binding protein of the archaeon <i>Sulfolobus</i> . <i>Nucleic Acids Research</i> , 2005, 33, 564-576.	14.5	25
24	Structure and Properties of DNA Molecules Over The Full Range of Biologically Relevant Supercoiling States. <i>Scientific Reports</i> , 2018, 8, 6163.	3.3	25
25	Molecular biology of extremophiles: recent progress on the hyperthermophilic archaeon <i>Sulfolobus</i> . <i>Antonie Van Leeuwenhoek</i> , 2002, 81, 85-97.	1.7	23
26	Inhibition of translesion DNA polymerase by archaeal reverse gyrase. <i>Nucleic Acids Research</i> , 2009, 37, 4287-4295.	14.5	23
27	New control elements of bacteriophage T4 pre-replicative transcription. <i>Journal of Molecular Biology</i> , 1985, 182, 249-263.	4.2	22
28	β -Glycosidase from <i>Sulfolobus solfataricus</i> . <i>Methods in Enzymology</i> , 2001, 330, 201-215.	1.0	21
29	A novel thermostable protein-tag: optimization of the <i>Sulfolobus solfataricus</i> DNA-alkyl-transferase by protein engineering. <i>Extremophiles</i> , 2016, 20, 1-13.	2.3	21
30	Structure, evolution and properties of a novel repetitive DNA family in <i>Caenorhabditis elegans</i> . <i>Nucleic Acids Research</i> , 1988, 16, 8213-8231.	14.5	20
31	Enzymatic synthesis of oligosaccharides by two glycosyl hydrolases of <i>Sulfolobus solfataricus</i> . <i>Extremophiles</i> , 2001, 5, 145-152.	2.3	20
32	Lack of Strand-specific Repair of UV-induced DNA Lesions in Three Genes of the Archaeon <i>Sulfolobus solfataricus</i> . <i>Journal of Molecular Biology</i> , 2007, 365, 921-929.	4.2	20
33	Ionic network at the C-terminus of the β -glycosidase from the hyperthermophilic archaeon <i>Sulfolobus solfataricus</i> : Functional role in the quaternary structure thermal stabilization. <i>Proteins: Structure, Function and Bioinformatics</i> , 2002, 48, 98-106.	2.6	19
34	Positive supercoiling in thermophiles and mesophiles: of the good and evil. <i>Biochemical Society Transactions</i> , 2011, 39, 58-63.	3.4	19
35	In vivo and in vitro protein imaging in thermophilic archaea by exploiting a novel protein tag. <i>PLoS ONE</i> , 2017, 12, e0185791.	2.5	19
36	Chromatin Structure and Dynamics in Hot Environments: Architectural Proteins and DNA Topoisomerases of Thermophilic Archaea. <i>International Journal of Molecular Sciences</i> , 2014, 15, 17162-17187.	4.1	18

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37	Crystal structure of <i>Mycobacterium tuberculosis</i> O ⁶ -methylguanine-DNA methyltransferase protein clusters assembled on to damaged DNA. <i>Biochemical Journal</i> , 2016, 473, 123-133.	3.7	18
38	Interdomain interactions rearrangements control the reaction steps of a thermostable DNA alkyltransferase. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2017, 1861, 86-96.	2.4	18
39	PCR amplification and cloning of metallothionein complementary DNAs in temperate and Antarctic sea urchin characterized by a large difference in egg metallothionein content. <i>Cellular and Molecular Life Sciences</i> , 1997, 53, 472-477.	5.4	16
40	The Reverse Gyrase from <i>Pyrobaculum calidifontis</i> , a Novel Extremely Thermophilic DNA Topoisomerase Endowed with DNA Unwinding and Annealing Activities. <i>Journal of Biological Chemistry</i> , 2014, 289, 3231-3243.	3.4	15
41	Genome stability: recent insights in the topoisomerase reverse gyrase and thermophilic DNA alkyltransferase. <i>Extremophiles</i> , 2014, 18, 895-904.	2.3	14
42	Every OGT Is Illuminated by Fluorescent and Synchrotron Lights. <i>International Journal of Molecular Sciences</i> , 2017, 18, 2613.	4.1	14
43	Synergic and Opposing Activities of Thermophilic RecQ-like Helicase and Topoisomerase 3 Proteins in Holliday Junction Processing and Replication Fork Stabilization. <i>Journal of Biological Chemistry</i> , 2012, 287, 30282-30295.	3.4	13
44	Activity and regulation of archaeal DNA alkyltransferase. CONSERVED PROTEIN INVOLVED IN REPAIR OF DNA ALKYLATION DAMAGE.. <i>Journal of Biological Chemistry</i> , 2015, 290, 885.	3.4	12
45	NurA Is Endowed with Endo- and Exonuclease Activities that Are Modulated by HerA: New Insight into Their Role in DNA-End Processing. <i>PLoS ONE</i> , 2015, 10, e0142345.	2.5	12
46	Molecular biology of hyperthermophilic Archaea. <i>Advances in Biochemical Engineering/Biotechnology</i> , 1998, 61, 87-115.	1.1	11
47	The DNA Alkylguanine DNA Alkyltransferase-2 (AGT-2) Of <i>Caenorhabditis Elegans</i> Is Involved In Meiosis And Early Development Under Physiological Conditions. <i>Scientific Reports</i> , 2019, 9, 6889.	3.3	10
48	The Archaeal Topoisomerase Reverse Gyrase Is a Helix-destabilizing Protein That Unwinds Four-way DNA Junctions. <i>Journal of Biological Chemistry</i> , 2010, 285, 36532-36541.	3.4	8
49	New Insights into Structural and Functional Roles of Indole-3-acetic acid (IAA): Changes in DNA Topology and Gene Expression in Bacteria. <i>Biomolecules</i> , 2019, 9, 522.	4.0	8
50	Reverse gyrase: an unusual DNA manipulator of hyperthermophilic organisms. <i>Italian Journal of Biochemistry</i> , 2007, 56, 103-9.	0.3	8
51	Foreign transcriptional enhancers in yeast. I. Interactions of papovavirus transcriptional enhancers and a quiescent pseudopromoter on supercoiled plasmids. <i>Nucleic Acids Research</i> , 1988, 16, 8847-8868.	14.5	7
52	Foreign transcriptional enhancers in yeast. II. Interplay of the polyomavirus transcriptional enhancer and <i>Saccharomyces cerevisiae</i> promoter elements. <i>Nucleic Acids Research</i> , 1988, 16, 8869-8886.	14.5	4
53	The Prefoldin of the Crenarchaeon <i>Sulfolobus solfataricus</i> . <i>Protein and Peptide Letters</i> , 2008, 15, 1055-1062.	0.9	4
54	<i>Saccharomyces cerevisiae</i> multifunctional protein RAP1 binds to a conserved sequence in the Polyoma virus enhancer and is responsible for its transcriptional activity in yeast cells. <i>FEBS Letters</i> , 1993, 323, 77-82.	2.8	3

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55	Industrial-Scale Production of Thermostable Enzymes: The Model-System of the α -D-Glucosidase from <i>Sulfolobus Solfataricus</i> . , 1996, , 89-99.		0
56	Structure and Reaction Mechanism of the α -D-Glucosidase from the Archaeon <i>Sulfolobus Solfataricus</i> . , 1998, , 209-212.		0