

Fabrizia Fusetti

List of Publications by Year in descending order

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Version: 2024-02-01

26
papers

2,757
citations

279798

23
h-index

526287

27
g-index

27
all docs

27
docs citations

27
times ranked

3913
citing authors

#	ARTICLE	IF	CITATIONS
1	Quantitative proteomics analysis identifies MUC1 as an effect sensor of EGFR inhibition. <i>Oncogene</i> , 2019, 38, 1477-1488.	5.9	11
2	A p300 and SIRT1 Regulated Acetylation Switch of C/EBP β Controls Mitochondrial Function. <i>Cell Reports</i> , 2018, 22, 497-511.	6.4	45
3	A G β -Stimulated RapGEF Is a Receptor-Proximal Regulator of Dictyostelium Chemotaxis. <i>Developmental Cell</i> , 2016, 37, 458-472.	7.0	16
4	Non-canonical PRC1.1 Targets Active Genes Independent of H3K27me3 and Is Essential for Leukemogenesis. <i>Cell Reports</i> , 2016, 14, 332-346.	6.4	126
5	Protein costs do not explain evolution of metabolic strategies and regulation of ribosomal content: does protein investment explain an anaerobic bacterial <i>scpC</i> effect?. <i>Molecular Microbiology</i> , 2015, 97, 77-92.	2.5	57
6	Susceptibility to COPD: Differential Proteomic Profiling after Acute Smoking. <i>PLoS ONE</i> , 2014, 9, e102037.	2.5	32
7	<i>scpL</i> of <i>actococcus lactis</i> ... <i>YfiA</i> is necessary and sufficient for ribosome dimerization. <i>Molecular Microbiology</i> , 2014, 91, 394-407.	2.5	45
8	Light-harvesting complex II (LHCII) and its supramolecular organization in <i>Chlamydomonas reinhardtii</i> . <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2014, 1837, 63-72.	1.0	135
9	Proteolysin, a Novel Highly Thermostable and Cosolvent-Compatible Protease from the Thermophilic Bacterium <i>Coprothermobacter proteolyticus</i> . <i>Applied and Environmental Microbiology</i> , 2013, 79, 5625-5632.	3.1	31
10	Nonredundant and locus-specific gene repression functions of PRC1 paralog family members in human hematopoietic stem/progenitor cells. <i>Blood</i> , 2013, 121, 2452-2461.	1.4	54
11	Regulation of <i>ykrL</i> (<i>htpX</i>) by Rok and YkrK, a Novel Type of Regulator in <i>Bacillus subtilis</i> . <i>Journal of Bacteriology</i> , 2012, 194, 2837-2845.	2.2	7
12	Photosystem I of <i>Chlamydomonas reinhardtii</i> Contains Nine Light-harvesting Complexes (Lhca) Located on One Side of the Core. <i>Journal of Biological Chemistry</i> , 2011, 286, 44878-44887.	3.4	104
13	Crystal Structure of <i>Agaricus bisporus</i> Mushroom Tyrosinase: Identity of the Tetramer Subunits and Interaction with Tropolone. <i>Biochemistry</i> , 2011, 50, 5477-5486.	2.5	648
14	Differential effect of YidC depletion on the membrane proteome of <i>Escherichia coli</i> under aerobic and anaerobic growth conditions. <i>Proteomics</i> , 2010, 10, 3235-3247.	2.2	44
15	<i>Bacillus subtilis</i> SpoIIIJ and YqjG Function in Membrane Protein Biogenesis. <i>Journal of Bacteriology</i> , 2009, 191, 6749-6757.	2.2	39
16	Selenomethionine incorporation in proteins expressed in <i>Lactococcus lactis</i> . <i>Protein Science</i> , 2009, 18, 1121-1127.	7.6	39
17	The structural basis for peptide selection by the transport receptor OppA. <i>EMBO Journal</i> , 2009, 28, 1332-1340.	7.8	82
18	Functional analysis of the competence transcription factor ComK of <i>Bacillus subtilis</i> by characterization of truncation variants. <i>Microbiology (United Kingdom)</i> , 2006, 152, 473-483.	1.8	12

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19	Crystal Structure and Carbohydrate-binding Properties of the Human Cartilage Glycoprotein-39. <i>Journal of Biological Chemistry</i> , 2003, 278, 37753-37760.	3.4	183
20	Structure of Human Chitotriosidase. <i>Journal of Biological Chemistry</i> , 2002, 277, 25537-25544.	3.4	185
21	Crystal Structure of the Copper-Containing Quercetin 2,3-Dioxygenase from <i>Aspergillus japonicus</i> . <i>Structure</i> , 2002, 10, 259-268.	3.3	216
22	Structure of Tetrameric Human Phenylalanine Hydroxylase and Its Implications for Phenylketonuria. <i>Journal of Biological Chemistry</i> , 1998, 273, 16962-16967.	3.4	137
23	Crystal structure of the catalytic domain of human phenylalanine hydroxylase reveals the structural basis for phenylketonuria. <i>Nature Structural Biology</i> , 1997, 4, 995-1000.	9.7	162
24	Variability within the <i>Candida rugosa</i> Lipases family. <i>Protein Engineering, Design and Selection</i> , 1994, 7, 531-535.	2.1	97
25	Cloning and analysis of <i>Candida cylindracea</i> lipase sequences. <i>Gene</i> , 1993, 124, 45-55.	2.2	131
26	Cloning and nucleotide sequences of two lipase genes from <i>Candida cylindracea</i> . <i>Biochimica Et Biophysica Acta Gene Regulatory Mechanisms</i> , 1992, 1131, 227-232.	2.4	77