

Marie-Noëlle Rosso

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1589376/publications.pdf>

Version: 2024-02-01

21
papers

998
citations

567144

15
h-index

713332

21
g-index

22
all docs

22
docs citations

22
times ranked

1116
citing authors

#	ARTICLE	IF	CITATIONS
1	Lytic xylan oxidases from wood-decay fungi unlock biomass degradation. <i>Nature Chemical Biology</i> , 2018, 14, 306-310.	3.9	269
2	Genomic Analysis Enlightens Agaricales Lifestyle Evolution and Increasing Peroxidase Diversity. <i>Molecular Biology and Evolution</i> , 2021, 38, 1428-1446.	3.5	72
3	Enhanced degradation of softwood versus hardwood by the white-rot fungus <i>Pycnoporus coccineus</i> . <i>Biotechnology for Biofuels</i> , 2015, 8, 216.	6.2	67
4	The integrative omics of white-rot fungus <i>Pycnoporus coccineus</i> reveals co-regulated CAZymes for orchestrated lignocellulose breakdown. <i>PLoS ONE</i> , 2017, 12, e0175528.	1.1	64
5	A fungal family of lytic polysaccharide monooxygenase-like copper proteins. <i>Nature Chemical Biology</i> , 2020, 16, 345-350.	3.9	63
6	Fast solubilization of recalcitrant cellulosic biomass by the basidiomycete fungus <i>Laetisaria arvalis</i> involves successive secretion of oxidative and hydrolytic enzymes. <i>Biotechnology for Biofuels</i> , 2014, 7, 143.	6.2	53
7	The ectomycorrhizal basidiomycete <i>Laccaria bicolor</i> releases a secreted β -1,4 endoglucanase that plays a key role in symbiosis development. <i>New Phytologist</i> , 2018, 220, 1309-1321.	3.5	49
8	Fungal secretomics to probe the biological functions of lytic polysaccharide monooxygenases. <i>Carbohydrate Research</i> , 2017, 448, 155-160.	1.1	48
9	Visual Comparative Omics of Fungi for Plant Biomass Deconstruction. <i>Frontiers in Microbiology</i> , 2016, 7, 1335.	1.5	46
10	Integrative visual omics of the white-rot fungus <i>Polyporus brumalis</i> exposes the biotechnological potential of its oxidative enzymes for delignifying raw plant biomass. <i>Biotechnology for Biofuels</i> , 2018, 11, 201.	6.2	45
11	Gene family expansions and transcriptome signatures uncover fungal adaptations to wood decay. <i>Environmental Microbiology</i> , 2021, 23, 5716-5732.	1.8	44
12	Evolution of Fungal Carbohydrate-Active Enzyme Portfolios and Adaptation to Plant Cell-Wall Polymers. <i>Journal of Fungi (Basel, Switzerland)</i> , 2021, 7, 185.	1.5	38
13	Conserved white-rot enzymatic mechanism for wood decay in the Basidiomycota genus <i>Pycnoporus</i> . <i>DNA Research</i> , 2020, 27, .	1.5	32
14	Insights into an unusual Auxiliary Activity 9 family member lacking the histidine brace motif of lytic polysaccharide monooxygenases. <i>Journal of Biological Chemistry</i> , 2019, 294, 17117-17130.	1.6	30
15	Broad-specificity GH131 β -glucanases are a hallmark of fungi and oomycetes that colonize plants. <i>Environmental Microbiology</i> , 2019, 21, 2724-2739.	1.8	18
16	Large-scale phenotyping of 1,000 fungal strains for the degradation of non-natural, industrial compounds. <i>Communications Biology</i> , 2021, 4, 871.	2.0	18
17	The ectomycorrhizal basidiomycete <i>Laccaria bicolor</i> releases a GH28 polygalacturonase that plays a key role in symbiosis establishment. <i>New Phytologist</i> , 2022, 233, 2534-2547.	3.5	16
18	A Multiomic Approach to Understand How <i>Pleurotus eryngii</i> Transforms Non-Woody Lignocellulosic Material. <i>Journal of Fungi (Basel, Switzerland)</i> , 2021, 7, 426.	1.5	9

#	ARTICLE	IF	CITATIONS
19	Distribution of methionine sulfoxide reductases in fungi and conservation of the free-methionine-R-sulfoxide reductase in multicellular eukaryotes. <i>Free Radical Biology and Medicine</i> , 2021, 169, 187-215.	1.3	9
20	Plant wastes and sustainable refineries: What can we learn from fungi?. <i>Current Opinion in Green and Sustainable Chemistry</i> , 2022, 34, 100602.	3.2	5
21	Screening New Xylanase Biocatalysts from the Mangrove Soil Diversity. <i>Microorganisms</i> , 2021, 9, 1484.	1.6	3