

Marna D Yandeu-Nelson

List of Publications by Year in descending order

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Version: 2024-02-01

26
papers

888
citations

567281

15
h-index

580821

25
g-index

31
all docs

31
docs citations

31
times ranked

1453
citing authors

#	ARTICLE	IF	CITATIONS
1	Phylogenetic and experimental characterization of an acyl-ACP thioesterase family reveals significant diversity in enzymatic specificity and activity. <i>BMC Biochemistry</i> , 2011, 12, 44.	4.4	142
2	Subcellular-level resolution MALDI-MS imaging of maize leaf metabolites by MALDI-linear ion trap-Orbitrap mass spectrometer. <i>Analytical and Bioanalytical Chemistry</i> , 2015, 407, 2301-2309.	3.7	113
3	Characterization of two GL8 paralogs reveals that the 3-ketoacyl reductase component of fatty acid elongase is essential for maize (<i>Zea mays</i> L.) development. <i>Plant Journal</i> , 2005, 42, 844-861.	5.7	82
4	Nearly Identical Paralogs: Implications for Maize (<i>Zea mays</i> L.) Genome Evolution. <i>Genetics</i> , 2007, 175, 429-439.	2.9	60
5	Deficiency of maize starch-branching enzyme i results in altered starch fine structure, decreased digestibility and reduced coleoptile growth during germination. <i>BMC Plant Biology</i> , 2011, 11, 95.	3.6	55
6	High spatial resolution mass spectrometry imaging reveals the genetically programmed, developmental modification of the distribution of thylakoid membrane lipids among individual cells of maize leaf. <i>Plant Journal</i> , 2017, 89, 825-838.	5.7	52
7	Spatial Mapping and Profiling of Metabolite Distributions during Germination. <i>Plant Physiology</i> , 2017, 174, 2532-2548.	4.8	50
8	Biological origins of normal-chain hydrocarbons: a pathway model based on cuticular wax analyses of maize silks. <i>Plant Journal</i> , 2010, 64, 618-632.	5.7	40
9	Starch-Branched Enzyme IIa Is Required for Proper Diurnal Cycling of Starch in Leaves of Maize. <i>Plant Physiology</i> , 2011, 156, 479-490.	4.8	36
10	Temperature gradient capillary electrophoresis (TGCE) – a tool for the high-throughput discovery and mapping of SNPs and IDPs. <i>Theoretical and Applied Genetics</i> , 2005, 111, 218-225.	3.6	33
11	Unequal Sister Chromatid and Homolog Recombination at a Tandem Duplication of the <i>a1</i> Locus in Maize. <i>Genetics</i> , 2006, 173, 2211-2226.	2.9	31
12	MuDR Transposase Increases the Frequency of Meiotic Crossovers in the Vicinity of a Mu Insertion in the Maize <i>a1</i> Gene. <i>Genetics</i> , 2005, 169, 917-929.	2.9	28
13	Two distinct domains contribute to the substrate acyl chain length selectivity of plant acyl-ACP thioesterase. <i>Nature Communications</i> , 2018, 9, 860.	12.8	28
14	Effects of trans-acting Genetic Modifiers on Meiotic Recombination Across the <i>a1-sh2</i> Interval of Maize. <i>Genetics</i> , 2006, 174, 101-112.	2.9	27
15	Altering the Substrate Specificity of Acetyl-CoA Synthetase by Rational Mutagenesis of the Carboxylate Binding Pocket. <i>ACS Synthetic Biology</i> , 2019, 8, 1325-1336.	3.8	27
16	A robust and efficient method for the extraction of plant extracellular surface lipids as applied to the analysis of silks and seedling leaves of maize. <i>PLoS ONE</i> , 2017, 12, e0180850.	2.5	19
17	Maize <i>Glossy2</i> and <i>Glossy2-like</i> Genes Have Overlapping and Distinct Functions in Cuticular Lipid Deposition. <i>Plant Physiology</i> , 2020, 183, 840-853.	4.8	14
18	Microbial production of bi-functional molecules by diversification of the fatty acid pathway. <i>Metabolic Engineering</i> , 2016, 35, 9-20.	7.0	12

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19	Genetic and environmental variation impact the cuticular hydrocarbon metabolome on the stigmatic surfaces of maize. <i>BMC Plant Biology</i> , 2019, 19, 430.	3.6	11
20	Genetic control of kernel compositional variation in a maize diversity panel. <i>Plant Genome</i> , 2021, 14, e20115.	2.8	7
21	A multigenotype maize silk expression atlas reveals how exposure-related stresses are mitigated following emergence from husk leaves. <i>Plant Genome</i> , 2020, 13, e20040.	2.8	6
22	Identification of active site residues implies a two-step catalytic mechanism for acyl-ACP thioesterase. <i>Biochemical Journal</i> , 2018, 475, 3861-3873.	3.7	4
23	Predicting moisture content during maize nixtamalization using machine learning with NIR spectroscopy. <i>Theoretical and Applied Genetics</i> , 2021, 134, 3743-3757.	3.6	3
24	Remembering Dr. Nick Lauter (December 13, 1972 – January 7, 2021). <i>Current Plant Biology</i> , 2021, 27, 100214.	4.7	3
25	Development and application of a quantitative bioassay to evaluate maize silk resistance to corn earworm herbivory among progenies derived from Peruvian landrace Piura. <i>PLoS ONE</i> , 2019, 14, e0215414.	2.5	2
26	Substrate promiscuity of β -ketoacyl ACP Synthase III (KASIII): Understanding the structural basis for functional diversity of KASIII enzymes. <i>FASEB Journal</i> , 2013, 27, 559.4.	0.5	0