Diana Isolda Clotilde Fernandez

List of Publications by Year in descending order

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DIANA ISOLDA CLOTILDE

| # | Article | IF | CITATIONS |
|----|--|-------------------|-------------|
| 1 | Coffee resistance to the main diseases: leaf rust and coffee berry disease. Brazilian Journal of Plant Physiology, 2006, 18, 119-147. | 0.5 | 179 |
| 2 | Differentiation ofFusarium oxysporumf. sp.vasinfectumRaces on Cotton by Random Amplified Polymorphic DNA (RAPD) Analysis. Phytopathology, 1994, 84, 622. | 2.2 | 151 |
| 3 | Plant-Parasitic Nematode Infections in Rice: Molecular and Cellular Insights. Annual Review of Phytopathology, 2014, 52, 135-153. | 7.8 | 123 |
| 4 | Trifling variation in truffles. Nature, 1998, 394, 734-734. | 27.8 | 92 |
| 5 | 454â€pyrosequencing of <i>Coffea arabica</i> leaves infected by the rust fungus <i>Hemileia vastatrix</i> reveals <i>in planta</i> â€expressed pathogenâ€secreted proteins and plant functions in a late compatible plant–rust interaction. Molecular Plant Pathology, 2012, 13, 17-37. | 4.2 | 81 |
| 6 | Systematic Numbering of Vegetative Compatibility Groups in the Plant Pathogenic Fungus Fusarium oxysporum. Phytopathology, 1998, 88, 30-32. | 2.2 | 76 |
| 7 | Coffee (Coffea arabica L.) genes early expressed during infection by the rust fungus (Hemileia) Tj ETQq1 1 0.7843 | 314 rgBT / 4.2 | Overlock 10 |
| 8 | Population genetics and dynamics of the black truffle in a man-made truffle field. Heredity, 2001, 86, 451-458. | 2.6 | 65 |
| 9 | Characterization of a Single Clonal Lineage ofFusarium oxysporumf. sp.albedinisCausing Bayoud Disease of Date Palm in Morocco. Phytopathology, 1996, 86, 787. | 2.2 | 65 |
| 10 | Transcriptomic and histological responses of African rice (Oryza glaberrima) to Meloidogyne graminicola provide new insights into root-knot nematode resistance in monocots. Annals of Botany, 2017, 119, 885-899. | 2.9 | 54 |
| 11 | Review: Potential biotechnological assets related to plant immunity modulation applicable in engineering disease-resistant crops. Plant Science, 2018, 270, 72-84. | 3.6 | 52 |
| 12 | Dual RNAâ€seq reveals <i>Meloidogyne graminicola</i> transcriptome and candidate effectors during the interaction with rice plants. Molecular Plant Pathology, 2016, 17, 860-874. | 4.2 | 48 |
| 13 | Monitoring of the early molecular resistance responses of coffee (Coffea arabica L.) to the rust fungus (Hemileia vastatrix) using real-time quantitative RT-PCR. Plant Science, 2006, 170, 1045-1051. | 3.6 | 47 |
| 14 | <i>Fot</i> 1 Insertions in the <i>Fusarium oxysporum</i> f. sp. <i>albedinis</i> Genome Provide Diagnostic PCR Targets for Detection of the Date Palm Pathogen. Applied and Environmental Microbiology, 1998, 64, 633-636. | 3.1 | 47 |
| 15 | Meloidogyne incognita - rice (Oryza sativa) interaction: a new model system to study plant-root-knot nematode interactions in monocotyledons. Rice, 2014, 7, 23. | 4.0 | 45 |
| 16 | Angiosperm Gymnostoma trees produce root nodules colonized by arbuscular mycorrhizal fungi related to Glomus. New Phytologist, 2001, 149, 115-125. | 7.3 | 40 |
| 17 | Biotechnological applications for the improvement of coffee (Coffea arabica L.). In Vitro Cellular and Developmental Biology - Plant, 2002, 38, 129-138. | 2.1 | 38 |
| 18 | Resistance to Meloidogyne incognita expresses a hypersensitive-like response in Coffea arabica. European Journal of Plant Pathology, 2010, 127, 365-373. | 1.7 | 36 |

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|----|---|-----|-----------|
| 19 | Validation of RT-qPCR reference genes for in planta expression studies in Hemileia vastatrix, the causal agent of coffee leaf rust. Fungal Biology, 2011, 115, 891-901. | 2.5 | 36 |
| 20 | Identification of coffee WRKY transcription factor genes and expression profiling in resistance responses to pathogens. Tree Genetics and Genomes, 2010, 6, 767-781. | 1.6 | 34 |
| 21 | Cellular and molecular analyses of coffee resistance to Hemileia vastatrix and nonhost resistance to Uromyces vignae in the resistance-donor genotype HDT832/2. European Journal of Plant Pathology, 2012, 133, 141-157. | 1.7 | 32 |
| 22 | Rapid Detection of the Fusarium oxysporum Lineage Containing the Canary Island Date Palm Wilt Pathogen. Phytopathology, 1999, 89, 407-413. | 2.2 | 30 |
| 23 | Biphasic haustorial differentiation of coffee rust (<i>Hemileia vastatrix</i> race II) associated with defence responses in resistant and susceptible coffee cultivars. Plant Pathology, 2009, 58, 944-955. | 2.4 | 30 |
| 24 | Variability of nuclear and mitochondrial ribosomal DNA of a truffle species (Tuber aestivum). Mycological Research, 1996, 100, 547-550. | 2.5 | 27 |
| 25 | Sub-genomic origin and regulation patterns of a duplicated WRKY gene in the allotetraploid species Coffea arabica. Tree Genetics and Genomes, 2008, 4, 379-390. | 1.6 | 26 |
| 26 | Heat shock-induced susceptibility of green coffee leaves and berries to Colletotrichum gloeosporioides and its association to PR and hsp70 gene expression. Physiological and Molecular Plant Pathology, 2003, 63, 181-190. | 2.5 | 25 |
| 27 | Overview of the functional virulent genome of the coffee leaf rust pathogen Hemileia vastatrix with an emphasis on early stages of infection. Frontiers in Plant Science, 2014, 5, 88. | 3.6 | 25 |
| 28 | Rice susceptibility to root-knot nematodes is enhanced by the Meloidogyne incognita MSP18 effector gene. Planta, 2019, 250, 1215-1227. | 3.2 | 23 |
| 29 | Non-host resistance responses of <i>Arabidopsis thaliana</i> to the coffee leaf rust fungus (<i>Hemileia vastatrix</i>). Botany, 2010, 88, 621-629. | 1.0 | 22 |
| 30 | Identification and characterization of the Non-race specific Disease Resistance 1 (NDR1) orthologous protein in coffee. BMC Plant Biology, 2011, 11, 144. | 3.6 | 22 |
| 31 | Transgenic coffee fruits from Coffea arabica genetically modified by bombardment. In Vitro Cellular and Developmental Biology - Plant, 2009, 45, 532-539. | 2.1 | 21 |
| 32 | Promoter analysis of the WRKY transcription factors CaWRKY1a and CaWRKY1b homoeologous genes in coffee (Coffea arabica). Plant Cell Reports, 2013, 32, 1263-1276. | 5.6 | 19 |
| 33 | Title is missing!. European Journal of Plant Pathology, 1997, 103, 485-490. | 1.7 | 18 |
| 34 | Genetic diversity among isolates of Fusariumoxysporum f.sp. canariensis. Plant Pathology, 2000, 49, 155-164. | 2.4 | 16 |
| 35 | Analyses of the Root-Knot Nematode (Meloidogyne graminicola) Transcriptome during Host Infection Highlight Specific Gene Expression Profiling in Resistant Rice Plants. Pathogens, 2020, 9, 644. | 2.8 | 15 |
| 36 | Expression profiling of genes involved in the biotrophic colonisation of Coffea arabica leaves by Hemileia vastatrix. European Journal of Plant Pathology, 2012, 133, 261-277. | 1.7 | 14 |

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|----|---|------------------|--------------|
| 37 | Biological control mechanisms of D-pinitol against powdery mildew in cucumber. Physiological and Molecular Plant Pathology, 2014, 88, 52-60. | 2.5 | 14 |
| 38 | Evolutionarily conserved plant genes responsive to root-knot nematodes identified by comparative genomics. Molecular Genetics and Genomics, 2020, 295, 1063-1078. | 2.1 | 14 |
| 39 | Development of Diagnostic SCAR Markers for <i>Meloidogyne graminicola</i> , <i>M. oryzae</i> , and <i>M. salasi</i> Associated with Irrigated Rice Fields in Americas. Plant Disease, 2019, 103, 83-88. | 1.4 | 12 |
| 40 | A Chemosensory GPCR as a Potential Target to Control the Root-Knot Nematode Meloidogyne incognita Parasitism in Plants. Molecules, 2019, 24, 3798. | 3.8 | 11 |
| 41 | Differentially expressed genes in cotton plant genotypes infected with Meloidogyne incognita. Plant Science, 2009, 177, 492-497. | 3.6 | 9 |
| 42 | Recent Advances in Understanding Plant–Nematode Interactions in Monocots. Advances in Botanical Research, 2015, 73, 189-219. | 1.1 | 8 |
| 43 | The Mi-EFF1/Minc17998 effector interacts with the soybean GmHub6 protein to promote host plant parasitism by Meloidogyne incognita. Physiological and Molecular Plant Pathology, 2021, 114, 101630. | 2.5 | 8 |
| 44 | Early responses of coffee immunity-related genes to root-knot nematode infection. Physiological and Molecular Plant Pathology, 2017, 100, 142-150. | 2.5 | 7 |
| 45 | Minc00344 and Mj-NULG1a effectors interact with GmHub10 protein to promote the soybean parasitism by Meloidogyne incognita and M. javanica. Experimental Parasitology, 2021, 229, 108153. | 1.2 | 7 |
| 46 | Seed-Specific Stable Expression of the α-Al1 Inhibitor in Coffee Grains and the In Vivo Implications for the Development of the Coffee Berry Borer. Tropical Plant Biology, 2015, 8, 98-107. | 1.9 | 5 |
| 47 | Enzymology, Histological and Ultrastructural Effects of Ar-Turmerone on Culex pipiens pallens Larvae. Insects, 2020, 11, 336. | 2.2 | 5 |
| 48 | Assessing the transmission service risk. International Journal of Electrical Power and Energy Systems, 2006, 28, 119-126. | 5.5 | 2 |
| 49 | 12 Rust Fungi: Achievements and Future Challenges on Genomics and Host–Parasite Interactions. , 2013, , 315-341. | | 1 |
| 50 | Searching in Silico Novel Targets for Specific Coffee Rust Disease Control. Lecture Notes in Computer Science, 2020, , 109-115. | 1.3 | 1 |
| 51 | Special issue "Deepen knowledge in plant pathology for innovative agroecology― European Journal of Plant Pathology, 2018, 152, 853-854. | 1.7 | 0 |
| 52 | Transposable Elements in Fungal Pathogens: New Diagnostic Tools. , 2002, , 171-192. | | 0 |
| 53 | Meloidogyne-SP4 effector gene silencing reduces reproduction of root-knot nematodes in rice (Oryza) Tj ETQq1 | 1 0.78431 0.6 | 4 rgBT /Over |
| 54 | Proteome dataset of Hemileia vastatrix by LC–MS/MS label-free identification. Data in Brief, 2022, 43, 108433. | 1.0 | 0 |