

Catherine Marie Breton

List of Publications by Year in descending order

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35
papers

1,071
citations

516710

16
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414414

32
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36
all docs

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docs citations

36
times ranked

1085
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | A digital catalog of high-density markers for banana germplasm collections. <i>Plants People Planet</i> , 2022, 4, 61-67. | 3.3 | 7 |
| 2 | A Protocol for Detection of Large Chromosome Variations in Banana Using Next Generation Sequencing. , 2022, , 129-148. | | 1 |
| 3 | Unravelling the complex story of intergenomic recombination in ABB allotriploid bananas. <i>Annals of Botany</i> , 2021, 127, 7-20. | 2.9 | 27 |
| 4 | Filling the gaps in gene banks: Collecting, characterizing, and phenotyping wild banana relatives of Papua New Guinea. <i>Crop Science</i> , 2021, 61, 137-149. | 1.8 | 19 |
| 5 | A Dual-Successive-Screen Model at Pollen/Stigma and Pollen Tube/Ovary Explaining Paradoxical Self-Incompatibility Diagnosis in the Olive Tree – An Interpretative Update of the Literature. <i>Plants</i> , 2021, 10, 1938. | 3.5 | 6 |
| 6 | Ten simple rules for switching from face-to-face to remote conference: An opportunity to estimate the reduction in GHG emissions. <i>PLoS Computational Biology</i> , 2021, 17, e1009321. | 3.2 | 1 |
| 7 | Reply to Saumitou-Laprade et al. (2017) – “Controlling for genetic identity of varieties, pollen contamination and stigma receptivity is essential to characterize the self-incompatibility system of <i>Olea europaea</i> L.”. Eva: https://doi.org/10.1111/eva.12498 . <i>Evolutionary Applications</i> , 2018, 11, 1465-1470. | 3.1 | 9 |
| 8 | Characterization of olive progenies derived from a Tunisian breeding program by morphological traits and SSR markers. <i>Scientia Horticulturae</i> , 2018, 236, 127-136. | 3.6 | 12 |
| 9 | Identification of olive pollen donor trees and pollinizers under controlled pollination environment using STR markers. <i>Australian Journal of Crop Science</i> , 2018, 12, 1566-1572. | 0.3 | 2 |
| 10 | The sporophytic self-incompatibility mating system is conserved in <i>Olea europaea</i> subsp. <i>cuspidata</i> and <i>O. e. europaea</i> . <i>Euphytica</i> , 2017, 213, 1. | 1.2 | 3 |
| 11 | Comment on Saumitou et al. (2017): Elucidation of the genetic architecture of self-incompatibility in olive: evolutionary consequences and perspectives for orchard management™. <i>Evolutionary Applications</i> , 2017, 10, 855-859. | 3.1 | 6 |
| 12 | Potential of combining morphometry and ancient DNA information to investigate grapevine domestication. <i>Vegetation History and Archaeobotany</i> , 2017, 26, 345-356. | 2.1 | 20 |
| 13 | A model based on S-allele dominance relationships to explain pseudo self-fertility of varieties in the olive tree. <i>Euphytica</i> , 2016, 210, 105-117. | 1.2 | 14 |
| 14 | Specific features in the olive self-incompatibility system: A method to decipher S-allele pairs based on fruit settings. <i>Scientia Horticulturae</i> , 2015, 181, 62-75. | 3.6 | 15 |
| 15 | The self-incompatibility mating system of the olive (<i>Olea europaea</i> L.) functions with dominance between S-alleles. <i>Tree Genetics and Genomes</i> , 2014, 10, 1055-1067. | 1.6 | 39 |
| 16 | Genetic and environmental features for oil composition in olive varieties. <i>OCL - Oilseeds and Fats, Crops and Lipids</i> , 2014, 21, D504. | 1.4 | 3 |
| 17 | From the Olive Flower to the Drupe: Flower Types, Pollination, Self and Inter-Compatibility and Fruit Set. , 2013, , . | | 7 |
| 18 | New hypothesis elucidates self-incompatibility in the olive tree regarding S-alleles dominance relationships as in the sporophytic model. <i>Comptes Rendus - Biologies</i> , 2012, 335, 563-572. | 0.2 | 34 |

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 19 | Transfer of architectural traits from perennial <i>Helianthus mollis</i> Lam. to sunflower (<i>H. annuus</i> L.) and localisation of introgression. <i>Euphytica</i> , 2012, 186, 557-572. | 1.2 | 8 |
| 20 | Transcriptome Analysis of <i>Sarracenia</i> , an Insectivorous Plant. <i>DNA Research</i> , 2011, 18, 253-261. | 3.4 | 28 |
| 21 | Gene transfer from wild <i>Helianthus</i> to sunflower: topicalities and limits. <i>Oleagineux Corps Gras Lipides</i> , 2010, 17, 104-114. | 0.2 | 15 |
| 22 | Genetic Relationships between Cultivated and Wild Olive Trees (<i>Olea Europaea</i> L. Var. <i>Europaea</i> and <i>Tj ETQq0 0 0 ggBT /Overlock 10 Tf</i>) | 0.4 | 22 |
| 23 | Oil accumulation kinetic along ripening in four olive cultivars varying for fruit size. <i>Oleagineux Corps Gras Lipides</i> , 2009, 16, 58-64. | 0.2 | 15 |
| 24 | Oleaster (var. <i>sylvestris</i>) and subsp. <i>cuspidata</i> are suitable genetic resources for improvement of the olive (<i>Olea europaea</i> subsp. <i>europaea</i> var. <i>europaea</i>). <i>Genetic Resources and Crop Evolution</i> , 2009, 56, 393-403. | 1.6 | 49 |
| 25 | The origins of the domestication of the olive tree. <i>Comptes Rendus - Biologies</i> , 2009, 332, 1059-1064. | 0.2 | 90 |
| 26 | Are olive cultivars distinguishable from oleaster trees based on morphology of drupes and pits, oil composition and microsatellite polymorphisms?. <i>Acta Botanica Gallica</i> , 2008, 155, 531-545. | 0.9 | 8 |
| 27 | Genetic diversity in Tunisian olive accessions and their relatedness with other Mediterranean olive genotypes. <i>Scientia Horticulturae</i> , 2008, 115, 416-419. | 3.6 | 10 |
| 28 | Differences between native and introduced olive cultivars as revealed by morphology of drupes, oil composition and SSR polymorphisms: A case study in Tunisia. <i>Scientia Horticulturae</i> , 2008, 116, 280-290. | 3.6 | 87 |
| 29 | Taming the wild and "wilding"™ the tame: Tree breeding and dispersal in Australia and the Mediterranean. <i>Plant Science</i> , 2008, 175, 197-205. | 3.6 | 37 |
| 30 | Comparison between classical and Bayesian methods to investigate the history of olive cultivars using SSR-polymorphisms. <i>Plant Science</i> , 2008, 175, 524-532. | 3.6 | 82 |
| 31 | Reply to comment on Breton et al.: "Taming the wild and "wilding"™ the tame: Tree breeding and dispersal in Australia and the Mediterranean". <i>Plant Science</i> , 2008, 175, 208-209. | 3.6 | 1 |
| 32 | Genetic diversity and gene flow between the wild olive (oleaster, <i>Olea europaea</i> L.) and the olive: several Plio-Pleistocene refuge zones in the Mediterranean basin suggested by simple sequence repeats analysis. <i>Journal of Biogeography</i> , 2006, 33, 1916-1928. | 3.0 | 138 |
| 33 | Comparative Study of Methods for DNA Preparation from Olive Oil Samples to Identify Cultivar SSR Alleles in Commercial Oil Samples: A Possible Forensic Applications. <i>Journal of Agricultural and Food Chemistry</i> , 2004, 52, 531-537. | 5.2 | 80 |
| 34 | Olive domestication from structure of oleasters and cultivars using nuclear RAPDs and mitochondrial RFLPs. <i>Genetics Selection Evolution</i> , 2001, 33, S251. | 3.0 | 79 |
| 35 | Cultivar Identification in Olive Based on RAPD Markers. <i>Journal of the American Society for Horticultural Science</i> , 2001, 126, 668-675. | 1.0 | 93 |