

# Ramos, Hcc

## List of Publications by Year in descending order

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Version: 2024-02-01

32  
papers

277  
citations

933447

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h-index

1058476

14  
g-index

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docs citations

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times ranked

203  
citing authors

| #  | ARTICLE  | IF  | CITATIONS |
|----|--|-----|-----------|
| 1  | Topcross hybrids in papaya ( <i>Carica papaya</i> L.): evaluation of the potential for increasing fruit quality in new cultivars. Archives of Agronomy and Soil Science, 2022, 68, 1473-1486.                    | 2.6 | 5         |
| 2  | Genetic structure analysis of <i>Mauritia flexuosa</i> natural population from the Lençóis Maranhenses region using microsatellite markers. Scientia Agricola, 2022, 79, .                                       | 1.2 | 1         |
| 3  | Is there a possibility to improve a developed hybrid? A current demand on papaya ( <i>Carica papaya</i> L.). Euphytica, 2022, 218, .   | 1.2 | 0         |
| 4  | Combining ability of recombinant F4 papaya lines: a strategy to select hybrid combination. Scientia Agricola, 2021, 78, .  | 1.2 | 7         |
| 5  | Discovery of SNPs and InDels in papaya genotypes and its potential for marker assisted selection of fruit quality traits. Scientific Reports, 2021, 11, 292.   | 3.3 | 10        |
| 6  | Genetic diversity of papaya ( <i>Carica papaya</i> L.) F5 recombinant inbred lines using the Ward-MLM strategy. Genetic Resources and Crop Evolution, 2021, 68, 3333-3343.                                       | 1.6 | 0         |
| 7  | MOLECULAR CHARACTERIZATION OF ELITE LINES OF PAPAYA ( <i>Carica papaya</i> L.) VIA SSR MARKERS. Revista Do Especialista, 2021, 3, 49-58.   | 0.6 | 4         |
| 8  | New source of alleles for resistance to black spot and phoma spot in papaya ( <i>Carica papaya</i> L.). Euphytica, 2021, 217, 1.   | 1.2 | 1         |
| 9  | Genotype analysis by trait is a practical and efficient approach on discrimination of inbred lines and identification of papaya ( <i>Carica papaya</i> L.) ideotypes for fruit quality. Euphytica, 2021, 217, 1. | 1.2 | 2         |
| 10 | Papaya ( <i>Carica papaya</i> L.) S1 family recurrent selection: Opportunities and selection alternatives from the base population. Scientia Horticulturae, 2020, 260, 108848.                                   | 3.6 | 13        |
| 11 | Molecular sexing in papaya ( <i>Carica papaya</i> L.) seeds based on endosperm DNA. Euphytica, 2020, 216, 1.   | 1.2 | 4         |
| 12 | Topcross hybrids in papaya: Genes derived from backcrossing provide resistance to multiple diseases. Crop Protection, 2020, 137, 105240.   | 2.1 | 2         |
| 13 | Genotyping-by-sequencing technology reveals directions for coconut ( <i>Cocos nucifera</i> L.) breeding strategies for water production. Euphytica, 2020, 216, 1.  | 1.2 | 9         |
| 14 | TWENTY-TWO-YEAR PAPAYA BREEDING PROGRAM: FROM BREEDING STRATEGY ESTABLISHMENT TO CULTIVAR DEVELOPMENT. Revista Do Especialista, 2020, 1, 9-27.   | 0.6 | 9         |
| 15 | First report of a genetic map and evidence of QTL for resistance to CABMV in a segregating population of <i>Passiflora</i> . European Journal of Plant Pathology, 2019, 155, 903-915.                            | 1.7 | 7         |
| 16 | Quantification of floral abnormalities in a population generated from sexual polymorphism aiming at recurrent selection in papaya. Bragantia, 2019, 78, 158-165.   | 1.3 | 7         |
| 17 | Combining ability for fruit yield and quality in papaya recombinant inbred lines from the sexual conversion backcrossing. Euphytica, 2019, 215, 1.   | 1.2 | 5         |
| 18 | UC10: a new early Formosa papaya cultivar. Crop Breeding and Applied Biotechnology, 2019, 19, 131-134.   | 0.4 | 11        |

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|----|---|-----|-----------|
| 19 | Development of superior lines of papaya from the Formosa group using the pedigree method and REML/Blup procedure. <i>Bragantia</i> , 2019, 78, 350-360.                                       | 1.3 | 16        |
| 20 | â€UC14â€™: a new papaya cultivar with intermediate fruit size. <i>Crop Breeding and Applied Biotechnology</i> , 2019, 19, 226-229.  | 0.4 | 7         |
| 21 | Selection of legitimate dwarf coconut hybrid seedlings using DNA fingerprinting. <i>Crop Breeding and Applied Biotechnology</i> , 2018, 18, 409-416.  | 0.4 | 14        |
| 22 | A hermaphrodite genotype in dioecious papaya progeny: sex reversal or contamination?. <i>Euphytica</i> , 2018, 214, 1.  | 1.2 | 4         |
| 23 | Papaya recombinant inbred lines selection by image-based phenotyping. <i>Scientia Agricola</i> , 2018, 75, 208-215.   | 1.2 | 9         |
| 24 | Genetic diversity between papaya lines and their correlation with heterosis in hybrids for disease resistance and morpho-agronomic traits. <i>Summa Phytopathologica</i> , 2018, 44, 110-115. | 0.1 | 4         |
| 25 | Image-based phenotyping of morpho-agronomic traits in papaya fruits ( <i>Carica papaya</i> L. THB var.). <i>Australian Journal of Crop Science</i> , 2018, 12, 1750-1756.                     | 0.3 | 13        |
| 26 | Model-assisted phenotyping by digital images in papaya breeding program. <i>Scientia Agricola</i> , 2017, 74, 294-302.  | 1.2 | 22        |
| 27 | Combining ability of recombinant lines of papaya from backcrossing for sexual conversion. <i>Revista Ciencia Agronomica</i> , 2017, 48, .   | 0.3 | 6         |
| 28 | Development of a Gene-Centered SSR Atlas as a Resource for Papaya ( <i>Carica papaya</i> ) Marker-Assisted Selection and Population Genetic Studies. <i>PLoS ONE</i> , 2014, 9, e112654.      | 2.5 | 12        |
| 29 | Plant breeding with marker-assisted selection in Brazil. <i>Crop Breeding and Applied Biotechnology</i> , 2014, 14, 54-60.  | 0.4 | 29        |
| 30 | Combined Selection in Backcross Population of Papaya (&lt;i>Carica papaya&/i> L.) by the Mixed Model Methodology. <i>American Journal of Plant Sciences</i> , 2014, 05, 2973-2983.            | 0.8 | 20        |
| 31 | Seasonal and genetic influences on sex expression in a backcrossed segregating papaya population. <i>Crop Breeding and Applied Biotechnology</i> , 2011, 11, 97-105.                          | 0.4 | 23        |
| 32 | Comparison of multiallelic distances for the quantification of genetic diversity in the papaya. <i>Acta Scientiarum - Agronomy</i> , 2011, 33, .  | 0.6 | 1         |