

RÃ©gine Delourme

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8308179/publications.pdf>

Version: 2024-02-01

18
papers

3,402
citations

623734

14
h-index

888059

17
g-index

20
all docs

20
docs citations

20
times ranked

3203
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Large-scale transcriptomics to dissect 2Âyears of the life of a fungal phytopathogen interacting with its host plant. BMC Biology, 2021, 19, 55. | 3.8 | 21 |
| 2 | A geneâ€forâ€gene interaction involving a â€lateâ€™ effector contributes to quantitative resistance to the stem canker disease in <i>Brassica napus</i>. New Phytologist, 2021, 231, 1510-1524. | 7.3 | 19 |
| 3 | Besides stem canker severity, oilseed rape host genotype matters for the production of Leptosphaeria maculans fruit bodies. Fungal Ecology, 2021, 52, 101076. | 1.6 | 6 |
| 4 | A Modified Meiotic Recombination in Brassica napus Largely Improves Its Breeding Efficiency. Biology, 2021, 10, 771. | 2.8 | 7 |
| 5 | Long-read assembly of the <i>Brassica napus</i> reference genome Darmor-bzh. GigaScience, 2020, 9, . | 6.4 | 64 |
| 6 | Oilseed rape (Brassica napus) resistance to growth of Leptosphaeria maculans in leaves of young plants contributes to quantitative resistance in stems of adult plants. PLoS ONE, 2019, 14, e0222540. | 2.5 | 15 |
| 7 | Multi-year linkage and association mapping confirm the high number of genomic regions involved in oilseed rape quantitative resistance to blackleg. Theoretical and Applied Genetics, 2018, 131, 1627-1643. | 3.6 | 63 |
| 8 | Stable Quantitative Resistance Loci to Blackleg Disease in Canola (Brassica napus L.) Over Continents. Frontiers in Plant Science, 2018, 9, 1622. | 3.6 | 48 |
| 9 | Chromosome-scale assemblies of plant genomes using nanopore long reads and optical maps. Nature Plants, 2018, 4, 879-887. | 9.3 | 316 |
| 10 | Quantitative Resistance to Plant Pathogens in Pyramiding Strategies for Durable Crop Protection. Frontiers in Plant Science, 2017, 8, 1838. | 3.6 | 182 |
| 11 | Comparative genomic analysis of duplicated homoeologous regions involved in the resistance of Brassica napus to stem canker. Frontiers in Plant Science, 2015, 6, 772. | 3.6 | 38 |
| 12 | Homoeologous duplicated regions are involved in quantitative resistance of Brassica napus to stem canker. BMC Genomics, 2014, 15, 498. | 2.8 | 69 |
| 13 | Early allopolyploid evolution in the post-Neolithic <i>Brassica napus</i> oilseed genome. Science, 2014, 345, 950-953. | 12.6 | 2,089 |
| 14 | High-density SNP-based genetic map development and linkage disequilibrium assessment in Brassica napus L. BMC Genomics, 2013, 14, 120. | 2.8 | 198 |
| 15 | Metabotyping: A New Approach to Investigate Rapeseed (<i>Brassica napus</i> L.) Genetic Diversity in the Metabolic Response to Clubroot Infection. Molecular Plant-Microbe Interactions, 2012, 25, 1478-1491. | 2.6 | 26 |
| 16 | Assessment of a new strategy for selective phenotyping applied to complex traits in <i>Brassica napus</i>. Open Journal of Genetics, 2012, 02, 190-201. | 0.1 | 20 |
| 17 | Title is missing!. European Journal of Plant Pathology, 2003, 109, 871-881. | 1.7 | 218 |
| 18 | Differential growth of Leptosphaeria maculans in the stem of susceptible and partially resistant oilseed rape (Brassica napus L.) genotypes. Canadian Journal of Plant Pathology, 0, , 1-10. | 1.4 | 0 |