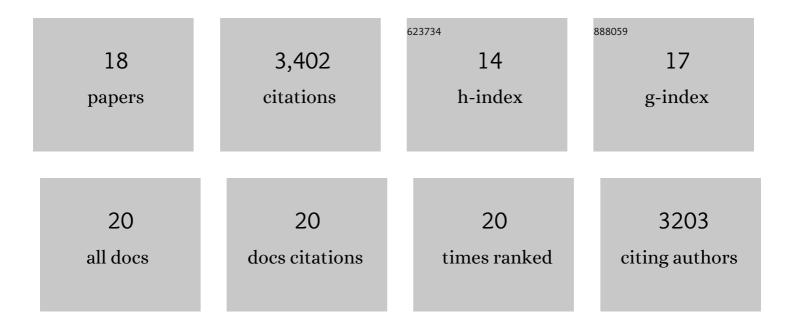
Régine Delourme

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8308179/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Early allopolyploid evolution in the post-Neolithic <i>Brassica napus</i> oilseed genome. Science, 2014, 345, 950-953.	12.6	2,089
2	Chromosome-scale assemblies of plant genomes using nanopore long reads and optical maps. Nature Plants, 2018, 4, 879-887.	9.3	316
3	Title is missing!. European Journal of Plant Pathology, 2003, 109, 871-881.	1.7	218
4	High-density SNP-based genetic map development and linkage disequilibrium assessment in Brassica napus L. BMC Genomics, 2013, 14, 120.	2.8	198
5	Quantitative Resistance to Plant Pathogens in Pyramiding Strategies for Durable Crop Protection. Frontiers in Plant Science, 2017, 8, 1838.	3.6	182
6	Homoeologous duplicated regions are involved in quantitative resistance of Brassica napus to stem canker. BMC Genomics, 2014, 15, 498.	2.8	69
7	Long-read assembly of the <i>Brassica napus</i> reference genome Darmor-bzh. GigaScience, 2020, 9, .	6.4	64
8	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
9	Stable Quantitative Resistance Loci to Blackleg Disease in Canola (Brassica napus L.) Over Continents. Frontiers in Plant Science, 2018, 9, 1622.	3.6	48
10	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
11	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
12	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
13	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
14	A geneâ€forâ€gene interaction involving a â€~late' effector contributes to quantitative resistance to the stem canker disease in <i>Brassica napus</i> . New Phytologist, 2021, 231, 1510-1524.	7.3	19
15	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
16	A Modified Meiotic Recombination in Brassica napus Largely Improves Its Breeding Efficiency. Biology, 2021, 10, 771.	2.8	7
17	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
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