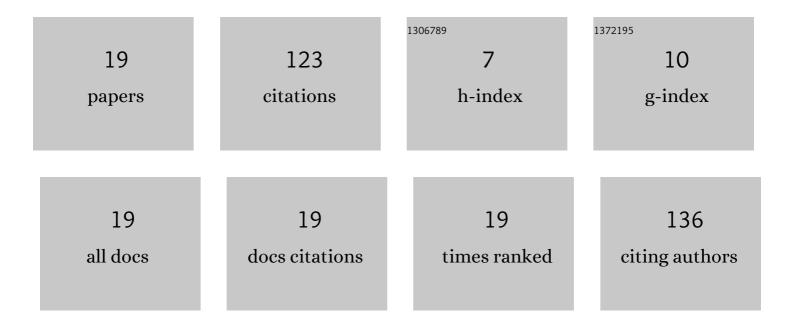
## Dhia Bouktila

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5917883/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Genome-wide survey of sugar beet (Beta vulgaris subsp. vulgaris) Dof transcription factors reveals structural diversity, evolutionary expansion and involvement in taproot development and biotic stress response. Biologia (Poland), 2021, 76, 2421-2436.	0.8	4
2	Genome-wide analysis of HSP90 gene family in the Mediterranean olive (Olea europaea subsp. europaea) provides insight into structural patterns, evolution and functional diversity. Physiology and Molecular Biology of Plants, 2020, 26, 2301-2318.	1.4	3
3	Genome-wide analysis of NBS-encoding resistance genes in the Mediterranean olive tree (Olea) Tj ETQq1 1 0.7843 function. Tree Genetics and Genomes, 2020, 16, 1.	814 rgBT / 0.6	Overlock 10 8
4	Genome-wide identification, characterization, and evolutionary analysis of NBS-encoding resistance genes in barley. 3 Biotech, 2018, 8, 453.	1.1	9
5	A novel method for molecular targeting of insecticide resistance in <i>Rhopalosiphum padi</i> L. (Homoptera: Aphididae). International Journal of Pest Management, 2016, 62, 284-287.	0.9	5
6	Large-scale bioinformatic analysis of the regulation of the disease resistance NBS gene family by microRNAs in Poaceae. Comptes Rendus - Biologies, 2016, 339, 347-356.	0.1	3
7	In Silico Identification of MicroRNAs with B/CYDV Gene Silencing Potential. Brazilian Archives of Biology and Technology, 2016, 59, .	0.5	1
8	Full-genome identification and characterization of NBS-encoding disease resistance genes in wheat. Molecular Genetics and Genomics, 2015, 290, 257-271.	1.0	21
9	Large-scale analysis of NBS domain-encoding resistance gene analogs in Triticeae. Genetics and Molecular Biology, 2014, 37, 598-610.	0.6	14
10	Characterization of novel wheat NBS domain-containing sequences and their utilization, in silico, for genome-scale R-gene mining. Molecular Genetics and Genomics, 2014, 289, 599-613.	1.0	7
11	A rapid diagnostic technique of <i>Bactrocera cucurbitae</i> and <i>Bactrocera zonata</i> (Diptera:) Tj ETQq1 1	0,784314 1.7	rgBT /Over
12	Development of New Polymorphic Microsatellite Loci for the Barley Stem Gall Midge, Mayetiola hordei (Diptera: Cecidomyiidae) from an Enriched Library. International Journal of Molecular Sciences, 2012, 13, 14446-14450.	1.8	12
13	Genetic variability of green citrus aphid populations from Tunisia, assessed by RAPD markers and mitochondrial DNA sequences. Entomological Science, 2012, 15, 171-179.	0.3	5
14	Genetic Variability of the Tomato Leaf Miner (Tuta absoluta Meirick; Lepidoptera: Gelechiidae), in Tunisia, Inferred from RAPD-PCR. Chilean Journal of Agricultural Research, 2012, 72, 212-216.	0.4	11
15	Hessian Fly, Mayetiola destructor (Say), Populations in the North of Tunisia: Virulence, Yield Loss Assessment and Phenological Data. Chilean Journal of Agricultural Research, 2011, 71, 401-405.	0.4	3
16	Characterization of Wheat Random Amplified Polymorphic DNA Markers Associated with the H11 Hessian Fly Resistance Gene. Journal of Integrative Plant Biology, 2006, 48, 958-964.	4.1	3
17	An overview of irritans-mariner transposons in two Mayetiola species (Diptera: Cecidomyiidae). European Journal of Entomology, 0, 114, 379-390.	1.2	4
18	Expression Analysis of Pyrenophora teres f. maculata-Responsive Loci in Hordeum vulgare. Brazilian Archives of Biology and Technology, 0, 62, .	0.5	0

#	Article	IF	CITATIONS
19	Large-scale identification and comparative characterization of date palm (Phoenix dactylifera L.) nucleotide-binding site (NBS) resistance genes provide insight into their structure, evolution and function. Trees - Structure and Function, 0, , .	0.9	0