

# Dhia Bouktila

## List of Publications by Year in descending order

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Version: 2024-02-01

19  
papers

123  
citations

1307366

7  
h-index

1372474

10  
g-index

19  
all docs

19  
docs citations

19  
times ranked

136  
citing authors

#	ARTICLE	IF	CITATIONS
1	Full-genome identification and characterization of NBS-encoding disease resistance genes in wheat. <i>Molecular Genetics and Genomics</i> , 2015, 290, 257-271.	1.0	21
2	Large-scale analysis of NBS domain-encoding resistance gene analogs in Triticeae. <i>Genetics and Molecular Biology</i> , 2014, 37, 598-610.	0.6	14
3	Development of New Polymorphic Microsatellite Loci for the Barley Stem Gall Midge, <i>Mayetiola hordei</i> (Diptera: Cecidomyiidae) from an Enriched Library. <i>International Journal of Molecular Sciences</i> , 2012, 13, 14446-14450.	1.8	12
4	Genetic Variability of the Tomato Leaf Miner ( <i>Tuta absoluta</i> Meirick; Lepidoptera: Gelechiidae), in Tunisia, Inferred from RAPD-PCR. <i>Chilean Journal of Agricultural Research</i> , 2012, 72, 212-216.	0.4	11
5	A rapid diagnostic technique of <i>Bactrocera cucurbitae</i> and <i>Bactrocera zonata</i> (Diptera: Tj ETQq1 1 0,784314 rgBT /Overlock 10	1.7	10
6	Genome-wide identification, characterization, and evolutionary analysis of NBS-encoding resistance genes in barley. <i>3 Biotech</i> , 2018, 8, 453.	1.1	9
7	Genome-wide analysis of NBS-encoding resistance genes in the Mediterranean olive tree ( <i>Olea</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 function. <i>Tree Genetics and Genomes</i> , 2020, 16, 1.	0.6	8
8	Characterization of novel wheat NBS domain-containing sequences and their utilization, in silico, for genome-scale R-gene mining. <i>Molecular Genetics and Genomics</i> , 2014, 289, 599-613.	1.0	7
9	Genetic variability of green citrus aphid populations from Tunisia, assessed by RAPD markers and mitochondrial DNA sequences. <i>Entomological Science</i> , 2012, 15, 171-179.	0.3	5
10	A novel method for molecular targeting of insecticide resistance in <i>Rhopalosiphum padi</i> L. (Homoptera: Aphididae). <i>International Journal of Pest Management</i> , 2016, 62, 284-287.	0.9	5
11	Genome-wide survey of sugar beet ( <i>Beta vulgaris</i> subsp. <i>vulgaris</i> ) Dof transcription factors reveals structural diversity, evolutionary expansion and involvement in taproot development and biotic stress response. <i>Biologia (Poland)</i> , 2021, 76, 2421-2436.	0.8	4
12	An overview of irritans-mariner transposons in two <i>Mayetiola</i> species (Diptera: Cecidomyiidae). <i>European Journal of Entomology</i> , 0, 114, 379-390.	1.2	4
13	Characterization of Wheat Random Amplified Polymorphic DNA Markers Associated with the H11 Hessian Fly Resistance Gene. <i>Journal of Integrative Plant Biology</i> , 2006, 48, 958-964.	4.1	3
14	Large-scale bioinformatic analysis of the regulation of the disease resistance NBS gene family by microRNAs in Poaceae. <i>Comptes Rendus - Biologies</i> , 2016, 339, 347-356.	0.1	3
15	Genome-wide analysis of HSP90 gene family in the Mediterranean olive ( <i>Olea europaea</i> subsp. <i>europaea</i> ) provides insight into structural patterns, evolution and functional diversity. <i>Physiology and Molecular Biology of Plants</i> , 2020, 26, 2301-2318.	1.4	3
16	Hessian Fly, <i>Mayetiola destructor</i> (Say), Populations in the North of Tunisia: Virulence, Yield Loss Assessment and Phenological Data. <i>Chilean Journal of Agricultural Research</i> , 2011, 71, 401-405.	0.4	3
17	In Silico Identification of MicroRNAs with B/CYDV Gene Silencing Potential. <i>Brazilian Archives of Biology and Technology</i> , 2016, 59, .	0.5	1
18	Expression Analysis of <i>Pyrenophora teres</i> f. <i>maculata</i> -Responsive Loci in <i>Hordeum vulgare</i> . <i>Brazilian Archives of Biology and Technology</i> , 0, 62, .	0.5	0

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