## Patrcia M Guimares

## List of Publications by Citations

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44 1,657 21 40 g-index

46 2,097 4.3 3.87 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
44	The genome sequences of Arachis duranensis and Arachis ipaensis, the diploid ancestors of cultivated peanut. <i>Nature Genetics</i> , <b>2016</b> , 48, 438-46	36.3	498
43	An analysis of synteny of Arachis with Lotus and Medicago sheds new light on the structure, stability and evolution of legume genomes. <i>BMC Genomics</i> , <b>2009</b> , 10, 45	4.5	107
42	Identification of candidate genome regions controlling disease resistance in Arachis. <i>BMC Plant Biology</i> , <b>2009</b> , 9, 112	5.3	98
41	ESTs from a wild Arachis species for gene discovery and marker development. <i>BMC Plant Biology</i> , <b>2007</b> , 7, 7	5.3	97
40	A linkage map for the B-genome of Arachis (Fabaceae) and its synteny to the A-genome. <i>BMC Plant Biology</i> , <b>2009</b> , 9, 40	5.3	83
39	Global transcriptome analysis of two wild relatives of peanut under drought and fungi infection. <i>BMC Genomics</i> , <b>2012</b> , 13, 387	4.5	72
38	Reference genes for quantitative reverse transcription-polymerase chain reaction expression studies in wild and cultivated peanut. <i>BMC Research Notes</i> , <b>2011</b> , 4, 339	2.3	48
37	Transcriptome Profiling of Wild from Water-Limited Environments Uncovers Drought Tolerance Candidate Genes. <i>Plant Molecular Biology Reporter</i> , <b>2015</b> , 33, 1876-1892	1.7	45
36	The use of SNP markers for linkage mapping in diploid and tetraploid peanuts. <i>G3: Genes, Genomes, Genetics</i> , <b>2014</b> , 4, 89-96	3.2	44
35	Root Transcriptome Analysis of Wild Peanut Reveals Candidate Genes for Nematode Resistance. <i>PLoS ONE</i> , <b>2015</b> , 10, e0140937	3.7	42
34	Matita, a new retroelement from peanut: characterization and evolutionary context in the light of the Arachis A-B genome divergence. <i>Molecular Genetics and Genomics</i> , <b>2012</b> , 287, 21-38	3.1	40
33	The effect of tetraploidization of wild Arachis on leaf morphology and other drought-related traits. <i>Environmental and Experimental Botany</i> , <b>2012</b> , 84, 17-24	5.9	40
32	Identification of QTLs for Rust Resistance in the Peanut Wild Species Arachis magna and the Development of KASP Markers for Marker-Assisted Selection. <i>G3: Genes, Genomes, Genetics</i> , <b>2015</b> , 5, 1403-13	3.2	38
31	Arachis batizocoi: a study of its relationship to cultivated peanut (A. hypogaea) and its potential for introgression of wild genes into the peanut crop using induced allotetraploids. <i>Annals of Botany</i> , <b>2015</b> , 115, 237-49	4.1	34
30	Genetic Mapping of Resistance to Meloidogyne arenaria in Arachis stenosperma: A New Source of Nematode Resistance for Peanut. <i>G3: Genes, Genomes, Genetics</i> , <b>2015</b> , 6, 377-90	3.2	32
29	A Study of Gene Expression in the Nematode Resistant Wild Peanut Relative, Arachis stenosperma, in Response to Challenge with Meloidogyne arenaria. <i>Tropical Plant Biology</i> , <b>2010</b> , 3, 183-192	1.6	28
28	Genome-wide analysis of expansin superfamily in wild Arachis discloses a stress-responsive expansin-like B gene. <i>Plant Molecular Biology</i> , <b>2017</b> , 94, 79-96	4.6	27

## (2017-2008)

27	BAC libraries construction from the ancestral diploid genomes of the allotetraploid cultivated peanut. <i>BMC Plant Biology</i> , <b>2008</b> , 8, 14	5.3	26	
26	The repetitive component of the A genome of peanut (Arachis hypogaea) and its role in remodelling intergenic sequence space since its evolutionary divergence from the B genome. <i>Annals of Botany</i> , <b>2013</b> , 112, 545-59	4.1	25	
25	A survey of genes involved in Arachis stenosperma resistance to Meloidogyne arenaria race 1. <i>Functional Plant Biology</i> , <b>2013</b> , 40, 1298-1309	2.7	24	
24	Phenotypic effects of allotetraploidization of wild and their implications for peanut domestication. <i>American Journal of Botany</i> , <b>2017</b> , 104, 379-388	2.7	23	
23	Ultrastructure of the Initial Interaction of Puccinia arachidis and Cercosporidium personatum with Leaves of Arachis hypogaea and Arachis stenosperma. <i>Journal of Phytopathology</i> , <b>2010</b> , 158, 792-796	1.8	20	
22	Development of a PCR test for the detection of Curtobacterium flaccumfaciens pv. flaccumfaciens. <i>Antonie Van Leeuwenhoek</i> , <b>2001</b> , 80, 1-10	2.1	20	
21	Ex vitro hairy root induction in detached peanut leaves for plant-nematode interaction studies. <i>Plant Methods</i> , <b>2017</b> , 13, 25	5.8	19	
20	Comparative root transcriptome of wild Arachis reveals NBS-LRR genes related to nematode resistance. <i>BMC Plant Biology</i> , <b>2018</b> , 18, 159	5.3	17	
19	Comparative proteomics and gene expression analysis in Arachis duranensis reveal stress response proteins associated to drought tolerance. <i>Journal of Proteomics</i> , <b>2019</b> , 192, 299-310	3.9	17	
18	Segmental allopolyploidy in action: Increasing diversity through polyploid hybridization and homoeologous recombination. <i>American Journal of Botany</i> , <b>2018</b> , 105, 1053-1066	2.7	16	
17	Early responses to dehydration in contrasting wild Arachis species. PLoS ONE, 2018, 13, e0198191	3.7	13	
16	Contrasting Effects of Wild Dehydrin Under Abiotic and Biotic Stresses. <i>Frontiers in Plant Science</i> , <b>2019</b> , 10, 497	6.2	12	
15	Characterization of raffinose metabolism genes uncovers a wild Arachis galactinol synthase conferring tolerance to abiotic stresses. <i>Scientific Reports</i> , <b>2020</b> , 10, 15258	4.9	9	
14	Comparative Genomics Reveals Novel Target Genes towards Specific Control of Plant-Parasitic Nematodes. <i>Genes</i> , <b>2020</b> , 11,	4.2	9	
13	The genome structure of (Linnaeus, 1753) and an induced allotetraploid revealed by molecular cytogenetics. <i>Comparative Cytogenetics</i> , <b>2018</b> , 12, 111-140	1	8	
12	Proteomics unravels new candidate genes for Meloidogyne resistance in wild Arachis. <i>Journal of Proteomics</i> , <b>2020</b> , 217, 103690	3.9	6	
11	Evolutionarily conserved plant genes responsive to root-knot nematodes identified by comparative genomics. <i>Molecular Genetics and Genomics</i> , <b>2020</b> , 295, 1063-1078	3.1	4	
10	Functional Genomics in Peanut Wild Relatives. Compendium of Plant Genomes, 2017, 149-164	0.8	4	

9	Overexpression of Wild Arachis Lipocalin Enhances Root-Knot Nematode Resistance in Peanut Hairy Roots. <i>Plant Molecular Biology Reporter</i> , <b>2019</b> , 37, 74-86	1.7	3
8	Transgenic tobacco plants expressing Tarin 1 inhibit the growth of Pseudomonas syringae pv. tomato and the development of Spodoptera frugiperda. <i>Annals of Applied Biology</i> , <b>2003</b> , 143, 349-357	2.6	3
7	Defining the combined stress response in wild Arachis. <i>Scientific Reports</i> , <b>2021</b> , 11, 11097	4.9	3
6	Presence of resveratrol in wild Arachis species adds new value to this overlooked genetic resource. <i>Scientific Reports</i> , <b>2020</b> , 10, 12787	4.9	1
5	Overexpression of DUF538 from Wild Arachis Enhances Plant Resistance to Meloidogyne spp <i>Agronomy</i> , <b>2021</b> , 11, 559	3.6	1
4	Ectopic expression of an expansin-like B gene from wild Arachis enhances tolerance to both abiotic and biotic stresses. <i>Plant Journal</i> , <b>2021</b> , 107, 1681-1696	6.9	1
3	Brazilian Kayabi Indian accessions of peanut, Arachis hypogaea (Fabales, Fabaceae): origin, diversity and evolution. <i>Genetics and Molecular Biology</i> , <b>2020</b> , 43, e20190418	2	О
2	Curtobacterium <b>2017</b> , 1-16		

1 Curtobacterium **2015**, 1-14