

# Martha C Giraldo Zapata

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8908599/publications.pdf>

Version: 2024-02-01

13  
papers

2,011  
citations

759233

12  
h-index

1125743

13  
g-index

13  
all docs

13  
docs citations

13  
times ranked

2210  
citing authors

#	ARTICLE	IF	CITATIONS
1	Growth and colonization of organic matter in soil by <i>Fusarium proliferatum</i> . <i>Canadian Journal of Plant Pathology</i> , 2019, 41, 242-250.	1.4	3
2	The Small GTPase MoSec4 Is Involved in Vegetative Development and Pathogenicity by Regulating the Extracellular Protein Secretion in <i>Magnaporthe oryzae</i> . <i>Frontiers in Plant Science</i> , 2016, 7, 1458.	3.6	24
3	Characterization and regulation of expression of an antifungal peptide from hemolymph of an insect, <i>Manduca sexta</i> . <i>Developmental and Comparative Immunology</i> , 2016, 61, 258-268.	2.3	30
4	pFPL Vectors for High-Throughput Protein Localization in Fungi: Detecting Cytoplasmic Accumulation of Putative Effector Proteins. <i>Molecular Plant-Microbe Interactions</i> , 2015, 28, 107-121.	2.6	26
5	Filamentous plant pathogen effectors in action. <i>Nature Reviews Microbiology</i> , 2013, 11, 800-814.	28.6	417
6	Two distinct secretion systems facilitate tissue invasion by the rice blast fungus <i>Magnaporthe oryzae</i> . <i>Nature Communications</i> , 2013, 4, 1996.	12.8	321
7	Gene-based SSR markers for common bean ( <i>Phaseolus vulgaris</i> L.) derived from root and leaf tissue ESTs: an integration of the BMC series. <i>BMC Plant Biology</i> , 2011, 11, 50.	3.6	79
8	Translocation of <i>Magnaporthe oryzae</i> Effectors into Rice Cells and Their Subsequent Cell-to-Cell Movement. <i>Plant Cell</i> , 2010, 22, 1388-1403.	6.6	426
9	Interaction Transcriptome Analysis Identifies <i>Magnaporthe oryzae</i> BAS1-4 as Biotrophy-Associated Secreted Proteins in Rice Blast Disease. <i>Plant Cell</i> , 2009, 21, 1273-1290.	6.6	346
10	Development and diversity of Andean-derived, gene-based microsatellites for common bean ( <i>Phaseolus</i> ) Tj ETQq0 0,0 rgBT /Overlock 10	3.6	62
11	Development of microsatellite markers for common bean ( <i>Phaseolus vulgaris</i> L.) based on screening of non-enriched, small-insert genomic libraries. <i>Genome</i> , 2009, 52, 772-782.	2.0	37
12	Characterization of AT-rich microsatellites in common bean ( <i>Phaseolus vulgaris</i> L.). <i>Theoretical and Applied Genetics</i> , 2008, 118, 91-103.	3.6	39
13	Microsatellite marker diversity in common bean ( <i>Phaseolus vulgaris</i> L.). <i>Theoretical and Applied Genetics</i> , 2006, 113, 100-109.	3.6	201