

Comparative Genome Analysis of Filamentous Fungi Re Associated with Fungal Pathogenesis

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Citation Report

#	ARTICLE	IF	CITATIONS
1	The <i>Laccaria</i> genome: a symbiont blueprint decoded. <i>New Phytologist</i> , 2008, 180, 296-310.	3.5	92
2	Funabase: a Fungal phylogenomic database. <i>BMC Bioinformatics</i> , 2008, 9, 456.	1.2	60
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5	Constructing Physical and Genomic Maps for <i>Puccinia striiformis</i> sp. <i>tritici</i> , the Wheat Stripe Rust Pathogen, by Comparing Its EST Sequences to the Genomic Sequence of <i>P. graminis</i> sp. <i>tritici</i> , the Wheat Stem Rust Pathogen. <i>Comparative and Functional Genomics</i> , 2009, 2009, 1-13.	2.0	2
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9	Molecular cloning and characterization of the <i>AVR-Pia</i> locus from a Japanese field isolate of <i>Magnaporthe oryzae</i> . <i>Molecular Plant Pathology</i> , 2009, 10, 361-374.	2.0	68
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17	FUNGIpath: a tool to assess fungal metabolic pathways predicted by orthology. <i>BMC Genomics</i> , 2010, 11, 81.	1.2	39
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38	Conservation of ciliary proteins in plants with no cilia. <i>BMC Plant Biology</i> , 2011, 11, 185.	1.6	26

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40	The biology of blast: Understanding how <i>Magnaporthe oryzae</i> invades rice plants. <i>Fungal Biology Reviews</i> , 2011, 25, 61-67.	1.9	26
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164	Virulence and pathogenicity determinants in whole genome sequence of <i>Fusarium udum</i> causing wilt of pigeon pea. <i>Frontiers in Microbiology</i> , 0, 14, .	1.5	0
165	Role of Cytochrome P450 Enzyme in Plant Microorganismsâ€™ Communication: A Focus on Grapevine. <i>International Journal of Molecular Sciences</i> , 2023, 24, 4695.	1.8	8
166	Ancient and pervasive expansion of adaptin-related vesicle coat machinery across Parabasalia. <i>International Journal for Parasitology</i> , 2023, 53, 233-245.	1.3	0