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320	Pheromone-regulated target genes respond differentially to MAPK phosphorylation of transcription factor Prf1. <b>2008</b> , 69, 1041-53		34
319	The information highways of a biotechnological workhorsesignal transduction in Hypocrea jecorina. <b>2008</b> , 9, 430		61
318	MADS-box transcription factor mig1 is required for infectious growth in Magnaporthe grisea. <i>Eukaryotic Cell</i> , <b>2008</b> , 7, 791-9		72
317	Mitogen-activated protein kinase cascade required for regulation of development and secondary metabolism in Neurospora crassa. <i>Eukaryotic Cell</i> , <b>2008</b> , 7, 2113-22		75
316	Interaction transcriptome analysis identifies Magnaporthe oryzae BAS1-4 as Biotrophy-associated secreted proteins in rice blast disease. <b>2009</b> , 21, 1273-90		248
315	Advances in Genetics, Genomics and Control of Rice Blast Disease. 2009,		24
314	The PMK1 MAP Kinase Pathway and Infection-Related Morphogenesis. <b>2009</b> , 13-21		3
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312	Under pressure: investigating the biology of plant infection by Magnaporthe oryzae. <b>2009</b> , 7, 185-95		596
311	Study of mRNA Expression by Real Time PCR of Cpkk1, Cpkk2 and Cpkk3, three MEKs of Cryphonectria parasitica, in Virus-free and Virus-infected Isogenic Isolates. <b>2009</b> , 158, 409-416		9
310	MoFLP1, encoding a novel fungal fasciclin-like protein, is involved in conidiation and pathogenicity in Magnaporthe oryzae. <b>2009</b> , 10, 434-44		40
309	MAP kinase signalling pathway components and targets conserved between the distantly related plant pathogenic fungi Mycosphaerella graminicola and Magnaporthe grisea. <b>2009</b> , 46, 667-81		39
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307	The cAMP Signaling and MAP Kinase Pathways in Plant Pathogenic Fungi. 2009, 157-172		7
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305	Fusarium oxysporum Ste12 controls invasive growth and virulence downstream of the Fmk1 MAPK cascade. <b>2009</b> , 22, 830-9		67
304	A yeast STE11 homologue CoMEKK1 is essential for pathogenesis-related morphogenesis in Colletotrichum orbiculare. <b>2010</b> , 23, 1563-72		26

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303	The cAMP signaling pathway in Fusarium verticillioides is important for conidiation, plant infection, and stress responses but not fumonisin production. <b>2010</b> , 23, 522-33	63
302	Cpkk1, MAPKK of Cryphonectria parasitica, is necessary for virulence on chestnut. <b>2010</b> , 100, 1100-10	7
301	Role of the Slt2 mitogen-activated protein kinase pathway in cell wall integrity and virulence in Candida glabrata. <b>2010</b> , 10, 343-52	53
300	The role of mitogen-activated protein (MAP) kinase signalling components and the Ste12 transcription factor in germination and pathogenicity of Botrytis cinerea. <b>2010</b> , 11, 105-19	93
299	An in vitro method for the analysis of infection-related morphogenesis in Fusarium graminearum. <b>2010</b> , 11, 361-9	38
298	A Candida albicans cell wall-linked protein promotes invasive filamentation into semi-solid medium. <b>2010</b> , 76, 733-48	30
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292	The homeodomain transcription factor Ste12: Connecting fungal MAPK signalling to plant pathogenicity. <b>2010</b> , 3, 327-32	29
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290	Microarray analysis of differential gene expression elicited in Trametes versicolor during interspecific mycelial interactions. <b>2010</b> , 114, 646-60	27
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287	The FUS3 MAPK signaling pathway of the citrus pathogen Alternaria alternata functions independently or cooperatively with the fungal redox-responsive AP1 regulator for diverse developmental, physiological and pathogenic processes. <b>2010</b> , 47, 381-91	42
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280	A MADS-box transcription factor MoMcm1 is required for male fertility, microconidium production and virulence in Magnaporthe oryzae. <b>2011</b> , 80, 33-53	74
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278	A Pmk1-interacting gene is involved in appressorium differentiation and plant infection in Magnaporthe oryzae. <i>Eukaryotic Cell</i> , <b>2011</b> , 10, 1062-70	25
277	Regulatory circuitry governing fungal development, drug resistance, and disease. <b>2011</b> , 75, 213-67	365
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150	Candidate gene based association mapping in Fusarium culmorum for field quantitative pathogenicity and mycotoxin production in wheat. <b>2017</b> , 18, 49	11
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148	Genome sequence of the ectophytic fungus Ramichloridium luteum reveals unique evolutionary adaptations to plant surface niche. <b>2017</b> , 18, 729	8
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