

**Supplementary Table S2.** List of primers used in qRT-PCR analysis.

ID	Gene Product Name	Primers (5' - 3')	Efficiency (%)	Concentration (nM)
TERG_00216	endochitinase ( <i>T. equinum</i> )	FW: TCGACAACGGTCTAGGGAGGT REV: CCTCGACGAAGGTAGCAATC	102.30	700
TERG_00759	conidiophore development protein <i>hym1</i> ( <i>T. violaceum</i> )	FW: GCGTACGTGGACAGTGGTGAT REV: GCGGTCGTCTCGAAGCA	105.94	400
TERG_01127	1,3-beta-glucan synthase component FKS1	FW: TTGACTGAGCGTGGTTTCTG REV: GGCGTAGATCTGACACACGA	98.35	400
TERG_01721	STE/STE20/YSK protein kinase <i>kic1</i>	FW: AGCGGGCAGGTCAAGCT REV: TGGTCATTGTAGCGGAAAGTTG	102.20	400
TERG_01788	cell morphogenesis protein <i>tao3</i> ( <i>T. equinum</i> )	FW: CCGGGTGGTAGACGAACCT REV: CGTACTAACTGGGTGATTGGTTCTG	96.35	500
TERG_02705	class III chitinase ( <i>T. tonsurans</i> )	FW: CAAGGAGATGCTGGAAGGAG REV: AGCCGAGGACGAAGTTGTTA	101.48	400
TERG_02719	glycosyl hydrolase ( <i>T. equinum</i> )	FW: AGCCCTTCTTCCTCATCCTC REV: GGTCCGGTAGAACTTGTCCTCA	103.95	300
TERG_02863	protein kinase activator <i>mob2</i> ( <i>T. equinum</i> )	FW: CAGCCCCTCATCGATATCTACAC REV: ACCGGCAAGGGCTTCAG	102.88	400
TERG_03379	AGC/NDR/NDR protein kinase <i>cbk1</i>	FW: AAATTGTCAAGAAGGCCACCTGCG REV: TGGAAGCTCGAAATGAGCGGTACA	95.93	200
TERG_03963	mannosyl phosphorylinositol ceramide synthase SUR1 ( <i>T. equinum</i> )	FW: AGCTGTACGACACAACGGAAT REV: ATACAGCCATGAAAGGACTGC	105.40	200
TERG_04234	hydrophobin, putative ( <i>T. verrucosum</i> )	FW: GGCATACATCTTGGTGGTTTC REV: CAGACAGTGGAGGTGGATGTT	95.09	400
TERG_04564	mixed-linked glucanase ( <i>T. equinum</i> )	FW: TCACTACGGAGTGCAAGACG REV: TTGGCTCCTGGTAGACGTTT	96.76	300
TERG_05576	cell wall glucanase (Scw11), putative ( <i>A. benhamiae</i> )	FW: CCCGACTCCAGGAACTTACA REV: GAAGGACAGACGGTTGCACT	98.00	500
TERG_07599	cell morphogenesis protein Sog2 ( <i>T. tonsurans</i> )	FW: TCCTGGAAATCCTCGACCTAAG REV: GCGGACTTCCTCAGGCAAT	101.34	400
TERG_08191	glucooligosaccharide oxidase ( <i>T. equinum</i> )	FW: CAGCTCAATGCTTTTGGAGAG REV: TCTCCGTCTACCATGGACTGT	91.90	300
TERG_12319	chitin synthase 2 <i>chs</i>	F: AGCCAACCTGCCTTGTACCAT R: GTAATCCGACCCATCCCTTT	90.29	600
TERG_04402	glyceraldehyde 3-phosphate dehydrogenase	FW: GCGTGACCCAGCCAACA REV: CGGTGGACTCGACGATGTAGT	99.90	200
TERG_05742	DNA-dependent RNA polymerase II	FW: TGCAGGAGCTGGTGGAGA REV: GCTGGGAGGTACTGTTTGATCAA	94.99	300