

Table S8. List of primers used in this study.

Gene model name	Application	Primer pair	Sequence 5' – 3'
<i>Vna4.761</i>	RT-qPCR	CHR4.761-F CHR4.761-R	CAACGGCGATGCACACATTA CGCAGCCTCCGGTGATC
<i>Vna5.694</i>	RT-qPCR	CHR5.694-F CHR5.694-R	CCTGGCGACTACACTCCTCCTA CAGACCTGCCAGCCTTGCT
<i>Vna7.443</i>	RT-qPCR	CHR7.443-F CHR7.443-R	TGGCAGCTGGTATCCTGG GTTCCCTTGCCCTCGTACTT
<i>Vna8.691</i>	RT-qPCR	CHR8.691-F CHR8.691-R	CCAAGGGAGGATACAGCACCTA CCTCGCTGAGATCCTTCTGTAC
<i>VnaUn.279</i>	RT-qPCR	UN.279-F UN.279-R	TTCCCATTACAGCCCTTTTCAC CGGCCCTCCATGGACAAGA
<i>Vna8.801</i> (splicing factor 3a2)	RT-qPCR	SF-F SF-R	GCGCTCGAGACGATCAACCT AGGTACGAGCCGTCGTTCTG
<i>VnaUn.148</i> (DNA topoisomerase)	RT-qPCR	Top-F Top-R	GCATGGTTTTTCGGATCCAT TACCTCGCGAAATCCTCAA
<i>Vna5.694</i>	KO ^b	5.694-O1-F ^a 5.694-O2-R ^a	GGTCTTAAUACACATCTGCGAGCACTGAG GGCATTAAUTGTTGGCGTGTGTGAAGAT
	KO ^c	5.694-A3-F ^a 5.694-A4-R ^a	GGA CTTAAUATAAGTGGCAAGCGGTTTT GGGTTAAUGTTGTCTCGAGCCGTTTGT
	KO ^d	5.694-T1-F ^a 5.694-T2-R ^a	ATGCAGTTCCTCACTCTCCT CTAAATGCAGTAGGGGCTCT
	KO ^e	5.694-T4-F ^a RF-2	GGGAAAAGCGGCAGATATTA TCTCCTTGCATGCACCATTCCCTTG
	KO ^e	5.694-T3-F ^a RF-1	AGGTGGTTGTGGAAGTCTGG AAATTTTGTGCTCACCGCCTGGAC
<i>Vna7.443</i>	KO ^b	7.443-O1-F ^a 7.443-O2-R ^a	GGTCTTAAUCACGAGCCTCTCTGTGTG GGCATTAAUTGGTGTGGGTTGGGTAGAT
	KO ^c	7.443-A3-F ^a 7.443-A4-R ^a	GGA CTTAAUGGTTCTCCTCCATCCTCTT GGGTTAAUCCCTACATTTGCCTTGCATT
	KO ^d	7.443-T1-F ^a 7.443-T2-R ^a	ATGCGCTACTCCCTTACTCTC TCACTGCTCAGGGTGCCACTT
	KO ^e	7.443-T4-F ^a RF-2	CATTGTCGAGGCTGGCTATT TCTCCTTGCATGCACCATTCCCTTG
	KO ^e	7.443-T3-F ^a RF-1	CACTCAGCATCCGTGACAGT AAATTTTGTGCTCACCGCCTGGAC
<i>Vna8.691</i>	KO ^b	8.691-O1-F ^a 8.691-O2-R ^a	GGTCTTAAUAGATTGAGGGGAAACCCTGT GGCATTAAUCATCGCCTCAAGAAGTCCTC
	KO ^c	8.691-A3-F ^a 8.691-A4-R ^a	GGA CTTAAUGCGTTGATCAATGGTCATGT GGGTTAAUCTGAGACTTCTCCGCCTCAC
	KO ^d	8.691-T1-F ^a 8.691-T2-R ^a	ATGGGAGACCAACGAACA CTATTCTCGGCATCCTG
	KO ^e	8.691-T4-F ^a RF-2	GGACCATGCCGAGACATAAC TCTCCTTGCATGCACCATTCCCTTG
	KO ^e	8.691-T3-F ^a RF-1	CACTCGATCCTGCAACAAAG AAATTTTGTGCTCACCGCCTGGAC
<i>VnaUn.279</i>	KO ^b	UN.279-O1-F ^a UN.279-O2-R ^a	GGTCTTAAUAGTTGTGTAACGGCCCTAGC GGCATTAAUTCACGCCTTTAGGAATTATCC
	KO ^c	UN.279-A3-F ^a UN.279-A4-R ^a	GGA CTTAAUTGGATTGAGGTTTCCGACAT GGGTTAAUTACGCCATCATCCAAGACAG
	KO ^d	UN.279-T1-F ^a UN.279-T2-R ^a	GCAGAGCAGGAGGATACGTC GAATGCTGCCAGAAGTGCA
	KO ^e	UN.279-T4-F ^a RF-2	GCAGAGCAGGAGGATACGTC TCTCCTTGCATGCACCATTCCCTTG
	KO ^e	RF-1 UN.279-T3-R ^a	AAATTTTGTGCTCACCGCCTGGAC GAATGCTGCCAGAAGTGCA
AFLP-9-1 ^f	fungal biomass quantification	9-1gs-F 9-1gs-R	GGTAACGTCATCGAACGACATC CACACGCTACATATCAAACAGCATAT