

Table S1. Primers used in this study.

Primers	Sequence (5'-3')	Remarks
<i>MaSom1</i> -F	ATTTCCCCTAAGTACTTCTAGAAATGGCCATGAACCCGAATAT	Used to clone the DNA fragment encompassing the <i>MaSom1</i> open reading frame
<i>MaSom1</i> -R	TTGCTCACCATGGATCCGTCGGCACCAATCGGGTCGC	
<i>MaSom1</i> -QF	TGGTAATGGCACTGTAA	Used for detecting the expression level of <i>MaSom1</i> gene
<i>MaSom1</i> -QR	TCTGAGGATTATTCTGGTTAT	
<i>Gapdh</i> -QF	AGATGGAGGAGTTGGTGTG	Used for detecting the expression level of internal control gene
<i>Gapdh</i> -QR	GACTGCCCGCATTGAGAAG	
ITS-F	TGGCATCTTCTGAGTGGTG	Used for measuring the concentration of fungal genomic DNA of hyphal bodies
ITS-R	CCCGTTGCGAGTGAGTTA	
<i>MaSom1</i> -RF	GCTGGCCGCCCATGGGATACAATGGCAACAGGGTCCTA	Used for screening the <i>MaSom1</i> -overexpression transformants
EGFP-VR	CCATCACCAGAATCCAACAC	
PgpdA-VF	TCTCCGAAGTAGGTAGAGC	Used for screening the empty transformant
<i>sur</i> -PF	CCTGAAATGTTGTGCTGTC	Used for Southern blot probe1 amplification
<i>sur</i> -PR	CGAGATTATGCCCAAGAAC	

<i>MaUve1</i> -QF	TAACAACGCACGAGATAT	Used for detecting the expression level of <i>MaUve1</i> gene
<i>MaUve1</i> -QR	AGGCTATAACCATACTCTG	
<i>MaUbi1</i> -QF	CAAGAAGAAGGTCTACACTACTC	Used for detecting the expression level of <i>MaUbi1</i> gene
<i>MaUbi1</i> -QR	AATGTTGCCGTCCTTGTC	
<i>MaSsb1</i> -QF	CTCTCGGTGTCGCTATGG	Used for detecting the expression level of <i>MaSsb1</i> gene
<i>MaSsb1</i> -QR	TCGGCAACAGTGGTGAAG	
<i>MaNTH1</i> -QF	TGTTGGCTGGCACTGAGAA	Used for detecting the expression level of <i>MaNTH1</i> gene
<i>MaNTH1</i> -QR	ACCAGTCCAAGCAAGCATCT	
<i>MaMpl1</i> -QF	GCTGTCTCTACCGCCTTTGT	Used for detecting the expression level of <i>MaMpl1</i> gene
<i>MaMpl1</i> -QR	CTACTGCTTGACCTTCTCCTTGAC	
<i>MaMad2</i> -QF	GCACTATGTCCATCCTTG	Used for detecting the expression level of <i>MaMad2</i> gene
<i>MaMad2</i> -QR	TGATGAGGGTCTTGATTTG	
