

S9 Table: Primers used in this study.

Primer	Sequence	Usage
Bar-up	CGCCTGGACGACTAAACC	Confirmation of the gene disruptions
Bar-down	TCAGCCTGCCGGTACCGC	
ΔMAA-Pks2-5-1	GGGGACAGCTTCTTGTACAAAGTGAAGTTGTGACTTCGGGTAG	Disruption of <i>Pks2</i> in <i>M. robertsii</i>
ΔMAA-Pks2-5-2	GGGGACTGCTTTTTGTACAAACTTGTGTGGACAGTATGCTATC	
ΔMAA-Pks2-3-1	GGGGACAACCTTTGTATAGAAAAGTTGTTAAGACGTGACATGAAC	
ΔMAA-Pks2-3-2	GGGGACAACCTTTGTATAATAAAGTTGTTCCGTGCTGTATCCATC	
ΔMAA-Pks2-CF-1	AATCACACGTCCGCAC	Confirmation of <i>Pks2</i> disruption in <i>M. robertsii</i>
ΔMAA-Pks2-CF-2	GTGCTCTTGTGCGTGAAC	
ΔMAN-Pks1-5-1	GGGGACAGCTTCTTGTACAAAGTGAAGAATGAGGACCGTCTTG	Disruption of <i>Pks1</i> in <i>M. anisopliae</i>
ΔMAN-Pks1-5-2	GGGGACTGCTTTTTGTACAAACTTGTGCTTACCAAGTGGTCCAC	
ΔMAN-Pks1-3-1	GGGGACAACCTTTGTATAGAAAAGTTGTTAGGTTCTGCTCATGATG	
ΔMAN-Pks1-3-2	GGGGACAACCTTTGTATAATAAAGTTGTGCACGTCCAAGAAGTTG	
ΔMAN-Pks1-CF-1	CGCACGAAAGTCTTCAC	Confirmation of the <i>Pks1</i> disruption in <i>M. anisopliae</i>
ΔMAN-Pks1-CF-2	ATGTTGGCCTTGACAAG	
ΔMBR-Pks1-5-1	GGGGACAGCTTCTTGTACAAAGTGAAGTTTCCTCTGCCATGTG	Disruption of <i>Pks1</i> in <i>M. brunneum</i>
ΔMBR-Pks1-5-2	GGGGACTGCTTTTTGTACAAACTTGTCTTGAGCACAGACATG	
ΔMBR-Pks1-3-1	GGGGACAACCTTTGTATAGAAAAGTTGTTACGAACGCAACGTTCCAC	
ΔMBR-Pks1-3-2	GGGGACAACCTTTGTATAATAAAGTTGTGTGCGCATCAAAGATG	
ΔMBR-Pks1-CF-1	TGATCAAGTTCTGCTC	Confirmation of <i>Pks1</i> disruption in <i>M. brunneum</i>
ΔMBR-Pks1-CF-2	CATGCTTCAAAGTCGTG	
ΔMGU-Pks1-5-1	GGGGACAGCTTCTTGTACAAAGTGAAGTTATCGTCGCGCAACAC	Disruption of <i>Pks1</i> in <i>M. guizhouense</i>
ΔMGU-Pks1-5-2	GGGGACTGCTTTTTGTACAAACTTGTGTACATTGAGCCATGAG	
ΔMGU-Pks1-3-1	GGGGACAACCTTTGTATAGAAAAGTTGTTAGGTTCTGCTCATGATG	
ΔMGU-Pks1-3-2	GGGGACAACCTTTGTATAATAAAGTTGTGCAAGCCTTCGCTCTTG	
ΔMGU-Pks1-CF-1	TGCTTCGCTCCATCAAC	Confirmation of <i>Pks1</i> disruption in <i>M. guizhouense</i>
ΔMGU-Pks1-CF-2	CATCTTGTGTCGCATC	
MAJ-Pks1-RNAi-5	gggaattcCGTGCTACAGCGGATGCCGGTCCAAAGTCCgatacgtgcctgccc	Cloning the genomic clone for knocking down <i>Pks1</i> in <i>M. majus</i>
MAJ-Pks1-RNAi-3	ggtctagaCGTGCTACAGCGGATGCCGGTCCAAAGTCCtgggaaagcgegttaca	
MAJ-Pks1-Promoter-F	ccgaattcAGAGCACTCAAATCATAG	Cloning the promoter for knocking down <i>Pks1</i> in <i>M. majus</i>
MAJ-Pks1-Promoter-R	ccgaattcGTTGATCCGAAGGTTGC	
ΔMAC-Pks1-5-1	GGGGACAGCTTCTTGTACAAAGTGAAGGGCATACTCTAATTTG	Disruption of <i>Pks1</i> in <i>M. acridum</i>
ΔMAC-Pks1-5-2	GGGGACTGCTTTTTGTACAAACTTGTCTGCTTCTGTTATG	
ΔMAC-Pks1-3-1	GGGGACAACCTTTGTATAGAAAAGTTGTTATCATTAGACCTGCGAG	
ΔMAC-Pks1-3-2	GGGGACAACCTTTGTATAATAAAGTTGTATGATAGCTGACTGGTG	
ΔMAC-Pks1-CF-1	ATTTCCAGCTCTCTAG	Confirmation of <i>Pks1</i> disruption in <i>M. acridum</i>
ΔMAC-Pks1-CF-2	GTGAAGACTTTCGTGAG	
MAM-Pks1-RNAi-5	gggaattcGCTTTGTTTCAGTCACCAAGCCTCGGCCAATgatacgtgcctgeccc	Cloning the promoter for knocking down <i>Pks1</i> in <i>M. album</i>
MAM-Pks1-RNAi-3	ggtctagaGCTTTGTTTCAGTCACCAAGCCTCGGCCAATtgggaaagcgegttaca	
MAM-Pks1-Promoter-F	ccgaattcGTAAGCAATCTGTCTTATC	Cloning the promoter for knocking down <i>Pks1</i> in <i>M. album</i>
MAM-Pks1-Promoter-R	ccgaattcGTTGATGCAGGCAGGCGTGC	
MAA-Gpd-RT-5	GTCGTCATCTCTGCTCCCTC	<i>gpd</i> gene in <i>M. robertsii</i>
MAA-Gpd-RT-3	CAATGGTGAACCTGTCGTGG	
MAA-Tef-RT-5	AGGCTGACTGCGCTATTCTC	<i>tef</i> gene in <i>M. robertsii</i>
MAA-Tef-RT-3	ACTTGGTGGTGCCATCTTG	
MAA-Pks1-RT-5	AACTACTTTGGAGACGGCCA	qRT-PCR for <i>Pks1</i> in <i>M. robertsii</i>

MAA-Pks1-RT-3	ACCAAGAGGCCAACTTTGAG	
MAA-Pks2-RT-5	GCACTGTCGAGAACCTGTCA	qRT-PCR for <i>Pks2</i> in <i>M. robertsii</i>
MAA-Pks2-RT-3	CACCTCGTCCCAGTTGATTT	
MAN-Pks1-RT-5	ACGATCAAGCAGTCTGATAC	qRT-PCR for <i>Pks1</i> in <i>M. anisopliae</i>
MAN-Pks1-RT-3	ACTAGAAAAGGCAGGGAATG	
MAN-Pks2-RT-5	TGGCCTGTTTCATCTACAAA	qRT-PCR for <i>Pks2</i> in <i>M. anisopliae</i>
MAA-Pks2-RT-3	CCAGCCACTATTACATCTCC	
MBR-Tef-RT-5	TATTCTCATTATCGCTGCCG	<i>tef</i> in <i>M. brunneum</i>
MBR-Tef-RT-3	CTCCTTGATGATTTCTCTGGT	
MBR-Pks1-RT-5	ACGATCAAGCAGTCTGATAC	qRT-PCR for <i>Pks1</i> in <i>M. brunneum</i>
MBR-Pks1-RT-3	TCGAGCAACCAGATTTTGTA	
MBR-Pks2-RT-5	AAAGTTTGACAAGACATGG	qRT-PCR for <i>Pks2</i> in <i>M. brunneum</i>
MBR-Pks2-RT-3	TTGCTGTGATAGTTTTCGGAA	
MGU-Gpd-RT-5	AAGAAGGTCATCATCTCTGC	<i>gpd</i> in <i>M. guizhouense</i>
MGU-Gpd-RT-3	CTCGACAATGGTGAAGTTGT	
MGU-Pks1-RT-5	ACGATCAAGCAGTCTGATAC	qRT-PCR for <i>Pks1</i> in <i>M. guizhouense</i>
MGU-Pks1-RT-3	ACTAGAAAAGGCAGGGAATG	
MGU-Pks2-RT-5	AATTTGCACAAGACATGGAC	qRT-PCR for <i>Pks2</i> in <i>M. guizhouense</i>
MGU-Pks2-RT-3	TGGGAAGGTGATTTTCTCG	
MAJ-Gpd-RT-5	CAGCTTTACTCTTGCGAATG	<i>gpd</i> in <i>M. majus</i>
MAJ-Gpd-RT-3	AGAACTTGACCTTCTTGCC	
MAJ-Pks1-RT-5	ACGATCAAGCAGTCTGATAC	qRT-PCR for <i>Pks1</i> in <i>M. majus</i>
MAJ-Pks1-RT-3	TCGAGCAACCAGATTTTGTA	
MAJ-Pks2-RT-5	TTTTGGCGAGAAAATACAGC	qRT-PCR for <i>Pks2</i> in <i>M. majus</i>
MAJ-Pks2-RT-3	GTGACAGTTTTGGAATTGGG	
MAC-Gpd-RT-5	CCAGCTTTTAGTTTTGCGAA	<i>gpd</i> in <i>M. acridum</i>
MAC-Gpd-RT-3	TTGATGTCGCCCTTGAAAAAT	<i>tef</i> in <i>M. acridum</i>
MAC-Tef-RT-5	GGGTAAGGAAGACAAGACTC	
MAC-Tef-RT-3	TGTAGGCTCCAAAACATACC	
MAC-Pks1-RT-5	CAAGAGGCATATCCAAC	qRT-PCR for <i>Pks1</i> and the fragment PS2 in Fig. S1 in <i>M. acridum</i>
MAC-Pks1-RT-3	GTCCGAGCAACCAAGAG	
MAC-Pks2-RT-5	GAAATAATGGAAGCTGTCCG	qRT-PCR for <i>Pks2</i> in <i>M. acridum</i>
MAC-Pks2-RT-3	ACACCACCCAATACTTTAC	
MAM-Gpd-RT-5	AAGAAGGTCATCATCTCTGC	<i>gpd</i> in <i>M. album</i>
MAM-Gpd-RT-3	CCTCAACAATGGTGAAGTTG	
MAM-Pks1-RT-5	CCAAATACCCACGTCTC	qRT-PCR for <i>Pks1</i> and the fragment PS2 in Fig. S1 in <i>M. album</i>
MAM-Pks1-RT-3	AGAGGACGAGCAAGAAG	
MAC-Pks1-RT-6-F	CCAGCTGTACGATCCTG	RT-PCR analysis of the fragment PS3 in <i>M. acridum</i> in Fig. S1
MAC-Pks1-RT-6-R	CACCTTCTCCAAGAATC	
MAM-Pks1-RT-6-F	CTGATTATGACTGCATC	RT-PCR analysis of the fragment PS3 in <i>M. album</i> in Fig. S1
MAM-Pks1-RT-6-R	GCAAACCTCTGGTGGTC	
PS1-5	MAC-Pks1-RT-6-F/ MAM-Pks1-RT-6-F	RT-PCR analysis of the fragment PS1 in Fig. S1.
PS1-3	MAC-Pks1-RT-3/ MAM-Pks1-RT-3	
MAC-Pks1-Re-F	ATGGAACACGTGACGATC	RT-PCR analysis of the fragment PS4 in <i>M. acridum</i> in Fig. S1.
MAC-Pks1-Re-R	GCAACCAAGAGGCCAAC	
MAM-Pks1-Re-F	ATGGAGCACGTGATAATC	RT-PCR analysis of the fragment PS4 in <i>M. album</i> in Fig. S1.
MAM-Pks1-Re-R	TTGAGTGACACAAGGAC	
MAA_Arp1-5	GGATCCATGAACCAATCTTTAACCC	Cloning the genomic clone of <i>M. robertsii</i> <i>Arp1</i> for constructing expression plasmid
MAA_Arp1-3	GATATCCTACGAATGTTTCTCCGAC	

MAA-Pks2-OE-5	GGATTTAAATATGGCAACGCAGGATGCGCCTAT	Cloning the genomic clone of <i>M. robertsii</i>
MAA-Pks2-OE-3	GGATTTAAATCTAGCTGGGACCAAGCAAAAATTC	<i>Pks2</i> for constructing overexpressing vector
MAA-Pks2-SEQ-F	AGTCCAGCAACGGACAG	Confirming the insertion of <i>M. robertsii Pks2</i>
Trpc-seq-5	TTCTTGTGACATGGAG	into the genomes of <i>M. album</i> and $\Delta Pks2$
MAA_Pks1_F1	CTTCAGTATATTCATCTTCCCATCCAAGAACCTTTAATCGATGAACCACGTGACGA TCAAGCAG	Amplification of <i>M. robertsii's Pks1</i> for heterologous expression in <i>A. nidulans</i>
MAA_Pks1_R1	CTCGACTTGGCAAGCTATGGC	
MAA_Pks1_F2	GATGCTGGCCATCCGAAGCAG	
MAA_Pks1_R2	CACAACATATTTTCGTCAGACACAGAATAACTCTCGCTAGCGGTTGAGAGACCAT AAACAATG	
MAA_Pks1_RT_F	CTGGTCAGCGATTGGCTCTAG	Confirmation of the insertion of <i>Pks1</i> into the genome of <i>A. nidulans</i>
MAA_Pks1_RT_R	GTCGTGATCTGCAATGGCATC	
MAA_Pks2_F1	CTTCAGTATATTCATCTTCCCATCCAAGAACCTTTAATCGCCGTCAAGAGAAGAGC TGTAATG	Amplification of <i>M. robertsii's Pks2</i> for heterologous expression in <i>A. nidulans</i>
MAA_Pks2_R1	GCACCACAATGGTAACCAAGG	
MAA_Pks2_F2	CTTGCTGCGCTGTATCAGGC	
MAA_Pks2_R2	CACAACATATTTTCGTCAGACACAGAATAACTCTCGCTAGGACTGGCTGCTCGATT GGACC	
MAA_Pks2_RT_F	GCATATCGCCCAAGGAGGCAG	Confirmation of the insertion of <i>Pks2</i> into the genome of <i>A. nidulans</i>
MAA_Pks2_RT_R	CGAAGGTGCGCAGGACAGAAC	
The loop DNA	GATACGCTGGCCTGCCAACCTTTTCGGTATAAAGACTTCGCGCTGATACCAGACG TTGCCCGCATAATTACGAATATCTGCATCGGCGAACTGATCGTTAAAACCTGCCTG GCACAGCAATTGCCCGGCTTTCTTGTAAACGCGTTTCCCA	The DNA fragment for the loop in a dsRNA for RNAi
