

Supplementary Figure S1. Alignment of *BcmfsG* coding sequence and homologous gene sequences from different *Botrytis* species: *B. cinerea* B05.10, *B. pseudocinerea* BP362, *B. medusae* B555, *B. fragariae* BCB16 and *B. byssoidea* MUCL94. Gene sequences were selected after running similarity search by Blastn using *BcmfsG* as query sequence. Multiple alignments of protein sequences were carried out with Clustal-Omega and were visualized using ESPrpt 3.0. Blue and green box represent conserved regions between *Botrytis* species within the region corresponding to the domain MFS-1 (pfam07690; interval 136-1140), targets for the design of primers for the identification of *mfsG* gene in other *Botrytis* species by PCR. **The arrows indicate the orientation of the primers used for the *mfsG* gene detection.**



Supplementary Figure S1 (part 2). Alignment of *BcmfsG* coding sequence and homologous gene sequences from different *Botrytis* species: *B. cinerea* B05.10, *B. pseudocinerea* BP362, *B. medusae* B555, *B. fragariae* BCB16 and *B. byssoidea* MUCL94. Gene sequences were selected after running similarity search by Blastn using *BcmfsG* as query sequence. Multiple alignments of protein sequences were carried out with Clustal-Omega and were visualized using ESPrpt 3.0. Blue and green box represent conserved regions between *Botrytis* species within the region corresponding to the domain MFS-1 (pfam07690; interval 136-1140), targets for the design of primers for the identification of *mfsG* gene in other *Botrytis* species by PCR. The arrows indicate the orientation of the primers used for the *mfsG* gene detection.



Supplementary Table S1. List of oligonucleotides used for the expression analysis study.

Name of the Oligo	Sequence	Description of the primer
CKmfsG1	GTTATGGGAATAGCG	Identification of the <i>mfsG</i> gene in other <i>Botrytis</i> species
CKmfsG2	AGAAGAATGGGAAAAAGA	Identification of the <i>mfsG</i> gene in other <i>Botrytis</i> species
CKmfsG3	GGAATGTCATATGAAGA	Identification of the <i>mfsG</i> gene in other <i>Botrytis</i> species
CKmfsG4	GAATACATTCAAAGTTC	Identification of the <i>mfsG</i> gene in other <i>Botrytis</i> species
FwMfsGcinerea	GTATACTGGAGCACAAAGCTT	mRNA level quantification of the gene <i>mfsG</i> in <i>B. cinerea</i>
RvMfsGcinerea	GGTGAATTTCTCGTCTTCTAG	mRNA level quantification of the gene <i>mfsG</i> in <i>B. cinerea</i>
FwMfsGpseudocinerea	ATACTGGAGCACAAAGCTTT	mRNA level quantification of the gene <i>mfsG</i> in <i>B. pseudocinerea</i>
RvMfsGpseudocinerea	CTGAATTTCTCGTCTTCTAGC	mRNA level quantification of the gene <i>mfsG</i> in <i>B. pseudocinerea</i>
FwMfsGbysssoidea	TTAGTCTTGGGAGGCTCAAT	mRNA level quantification of the gene <i>mfsG</i> in <i>B. bysssoidea</i>
RvMfsGbysssoidea	TTATATCCTCGCCCAGATG	mRNA level quantification of the gene <i>mfsG</i> in <i>B. bysssoidea</i>
FwActAcinerea	TTTGAGACCTTCAACGCCCC	mRNA level quantification of the gene <i>BcactA</i> in <i>B. cinerea</i>
RvActAcinerea	ACGTGAGTAACTCCGTCACC	mRNA level quantification of the gene <i>BcactA</i> in <i>B. cinerea</i>
FwActApseudocinerea	GGTGGTATCCACGTCATA	mRNA level quantification of the gene <i>BcactA</i> in <i>B. pseudocinerea</i>
RvActApseudocinerea	CATAACAATGTTACCATACAAATC	mRNA level quantification of the gene <i>BcactA</i> in <i>B. pseudocinerea</i>
FwActAbysssoidea	TGACGATGCTCCAAGAGC	mRNA level quantification of the gene <i>BcactA</i> in <i>B. bysssoidea</i>
RvActAbysssoidea	ATTGTCAATGACGAGGGCT	mRNA level quantification of the gene <i>BcactA</i> in <i>B. bysssoidea</i>
Fwtubcinerea	TCCTTTTCGGTCAACTCTTCCG	mRNA level quantification of the gene β -tubuline in <i>B. cinerea</i>
Rvtubcinerea	CACCCTCAGTGTTAATGACCC	mRNA level quantification of the gene β -tubuline in <i>B. cinerea</i>
Fwtubpseudocinerea	GATCTCCAACCTTGAGCGTA	mRNA level quantification of the gene β -tubuline in <i>B. pseudocinerea</i>
Rvtubpseudocinerea	TGCTCGCCAGAGATAGTTT	mRNA level quantification of the gene β -tubuline in <i>B. pseudocinerea</i>
Fwtubbysssoidea	GGTGATCAATTCAGTCTAT	mRNA level quantification of the gene β -tubuline in <i>B. bysssoidea</i>
Rvtubbysssoidea	TTCATGTTGGACTCAGCCT	mRNA level quantification of the gene β -tubuline in <i>B. bysssoidea</i>