



Figure S1. *Trichoderma* species phylogenetic tree. The tree was inferred from alignments of coding nucleotide sequences of *act1*, *call*, *fas1*, *lcb2*, *rpb2*, and *tef1* housekeeping genes. Trees for these six genes were inferred by two methods, and the results were combined: 1) nucleotide sequences from each gene were aligned separately, the alignments were concatenated, and the resulting concatenated alignment was subjected to maximum likelihood analysis using IQ-tree software 1.6.7[1]. Branch support was determined by bootstrap analysis after 1000 pseudoreplicates. 2) nucleotide sequences from each housekeeping gene were aligned separately and subjected to maximum likelihood analysis separately. The resulting six trees were used to generate a consensus tree using RAxML software [2]Branch support was determined by internode certainly (IC) analysis as implemented in RAxML[3]. The tree in the figure was inferred using method 1 but shows branch support values derived from both methods: bootstrap values are shown before the forward slash (i.e. /), and the IC values are shown in red color after the forward slash. At the right size of the tree, there are indicated in red color the names of the *Trichoderma* Clades or Sections where the different species included in the present study are located. The strains isolated and analyzed in the present work is squared with a blue line.

1. Nguyen, L.T.; Schmidt, H.A.; Von Haeseler, A.; Minh, B.Q. IQ-TREE: A fast and effective stochastic algorithm for estimating maximum-likelihood phylogenies. *Mol. Biol. Evol.* **2014**, *32*, 268–274, doi:10.1093/molbev/msu300.
2. Stamatakis, A. RAxML version 8: A tool for phylogenetic analysis and post-analysis of large phylogenies. *Bioinformatics* **2014**, *30*, 1312–1313, doi:10.1093/bioinformatics/btu033.
3. Salichos, L.; Stamatakis, A.; Rokas, A. Novel information theory-based measures for quantifying incongruence among phylogenetic trees. *Mol. Biol. Evol.* **2014**, *31*, 1261–1271, doi:10.1093/molbev/msu061.

Supplementary Data. Appendix. Sequences retrieved from the genomes analyzed in this work.

Appendix S1. *act1* resulting coding sequences retrieved from the genomic sequences

>*Trichoderma* _sp. T214

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ATGGCCAGCAGCTTGGCCACGTCGGCGAGCCAACCGACATTACGGCGGAGATGAGGT
CTCTGCCCTCGTCTCGACCCGGCTACTGTAACACCGCGAGCAGGTTTGCCGGCGAAGATGT
TCCCAATCAATCCTGCCGTCTTCTACGGCACATCACCAGCGATCCCCCGAGAACCTGTT
TGGCGATGAGTGCATCATCCCGCGCGACTTCGAGGGTCCGAACATACATGAACAAGGACA
GCGTTGGAGGATTGGGATGTCGCCGCCAAGATGTGGGAGTTATGCTGGTAAGCGCCTGC
AGCCCGAGAGACAGACGCCCTCGTCCAAGAATGGATTGAATGACGACGTGAAAGACCAAGGA
TGGCGAGGGCGATGTCGATGGAGGGAGATCGAGGGTCAGGAAAAGCCTCTGGAGGGAGTT
CCATTGCTGATGACAGAACGCTCCCTGGAACTCGCCAACGGCGTGAAAAGGCTATTGAGCT
TAGCATGGAGAGCTGGGGACCCGGCTTTGGTTGAGCCGGACACCTGTTATCTGCCTT
CGCTGCTGGCAAAGCGACGGCTTGTCAATCGATATCGGTGGCGAAACACCTCCGTACAG
CTATGCACGACGGCATGGTCTCAAGAGATCTACAGCGATCACCTGCAGGTGGCTGTGGC
TGTCTTCGAGATCCGCAACATGTGGAAACCTCTGAGGCCAACGCGACTTGTACCCGACAT
TCATGGTTGAGAACAAAGACTCCTGTCGATGCCCTCGCCCCGCCAGGCACGACTACGCGAA
TTCCCATACAAGATTGACTCCTCCGAGCATTGAAAGAGGGAGCGACTGCTCACCGAGTT
AAAGAGTCCGTCGAGGTTGGGTGGCCCCGGACGGTATGGTGGCTGGGAACGAGG
AGTACATCAAGTCACAGCCTGGCCCGTATTGAGATGCCGATGGCTACAACCAGATGTGGC
GCGAGCAGCGATTCAAGGTGGCCGAGGGAATGTGGGATGAGAACGCCGATACCCGTTCCC
GAGGCGGAGCGTCTCACCAGGCTCAGACTATCCCTGAGCTTATCCGCTCTGCTCTCACGCG
ATTGATGTTGATCTCGGGCAACCTCTCGCTAATGTTGTCGTTACTGGTAGACAAGCTTGA
TCAACGGCTTCAACGACCGACTCAACAATGAGCTGACAGCCATGTATCCGGGATTGAAGATC
AAGATTCACGCGCCGGCTTGTGAGCGAGCGTAGATTGGTGGATCACGGCAGCAT
TCTGGCAGTCTGGTACTTCCATCAGATGTGGATATCCAAGAAGGAGTATGAGGAAAACGG
ACCGGGCATTGTCGAGAACGCGATGCAAATAG
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>*Trichoderma* _sp. T065

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ATGGCCAGCAGCTTGGCCACGTCGGCTAGCCAACCGACATCTACGGCGGAG
ATGAGGTCTCTGCCCTCGTCTGACCCGGCTACTGCAATACCGCGAGCAGGCTCGCTGGCG
AAGATGCCCCAAGTCAAATCTACCATCGTTCTACGGCATGTCACCAGCGACCCCTCCCCGAG
ATTATTGGCGACGAGTGCATCATCCACGAAGCGACTTGGAGGTCCGCAACTACATGAACA
AGGACAGCGTCTGAGAGGACTGGGATGTGGCGTAAGATGTGGGATTATGCTGGTTAAG
CGCCTACAGCCCGAGCGACAGACGCCCATCAAAGAATAAGCTAACGACGACGTGAAGG
AGCAGGATGGCGAAGGAGATGTCGCGATGAAAGAAGTCGAGACGATGGAGAAGGCTCTAGA
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ATCGAGCTCAGCATGGAGAGCTGGGGACTCCGCCTCTGGCTAAGCCGGACTCCGTTCT
GTCGTCTTCGCGCTGGAAAAGCTACTGCCCTGTCATTGACGTTGGTGGCGCAAATACCTC
CGTTACAGCTATCCATGATGGCATGGTTGAAGCGATCTACCGCGATACCCGCGGGCGGT
CTATGGCTGTCTCGAGATCGCAACATGTGGAAACTTCCGAGGCCAAAGTGGACTTGGTT
CCGACGTTATGGTGGAGAACAGTCTCCGTTGATGCTCTTCCTGCCAGGCCGACTA
CGCGAGTCCCTACAAGATCCACGACTCTTCGAACATTGAAGAACAGGCCGCTGGAAA
GAGTTCAAGGAGTCCGTTGAGGTTGGCGTGGACCTGGAAGATAACGGCGCCGCTGGAAA
CGAAGAGTACATCAAATCTAGCCTGGCGTGTGTTGAGATGCCGTGATGGCTACAACCAGAT
GTGGCGCGAGCAGCGATTCAAGGTCGCTGAGGGCATGTGGGAGAAAACGCCGGTACCC
GTCCCCGAGTCGAGCGCCTCACCAAGGCCAGACCATCCCCGAACTTATCCGTTCCGCCCTC
AATGCTATTGACGTTGACCTGCGGGCAACCTCCTGCCAACGCGATGTACGGAAAGCAC
AAGTTGATCAACGGCTCAACGACCGTCTAAATAACGAGCTGACAGCGATGTACCCCTGGGCT
GAAGATCAAGATTACGCTGCAAGGTCTGTCAGCGAGCGCAGATTGGTGGCTGGATTGGT
GAAGCATTCTGCTAGCTGGAAACTTCCACCAAATGTGGATATCCAAAAAAGGAGTATGAGG
AAAACGGAGTGGCATTGTCGAGAACGCGATGCAAATAG
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Appendix S2. *call* resulting coding sequences retrieved from the genomic sequences

>*Trichoderma* _sp. T214

```
ATGGCTGATTCTCTTACCGAAGAGCAGGTCTCTGAGTTCAAGGAGGCCCTCCCTTTGAC
AAGGACGGTGATGCCAGATCACCACCAAGGAGCTGGCACCGTGATGCCCTCCCGCCA
GAACCCCTCCGAGTCAGAGCTCCAGGACATGATCAACGAGGTGATGCCGACAACACGGAT
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CTATCGATTCCCCGAATTCTGACCATGATGGCACGAAAGATGAAGGACACTGATTCCGAGG
AGGAGATCCGAGAGGCATTCAAGGTCTTGACCGCAGACAACAACGGCTCATCTCAGCCGCT
GAGCTGCGCCATGTCATGACATCCATCGCGAGAAGCTCACCACGACGAGGTCGATGAGAT
GATTGCGAGGCCGATCAGGATGGTATGCCATTGATTACAATGAGTTGTCCAGCTTAT
GATGCAAAATAA

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ATGGCTGATTCACTTACCGAAGAGCAGGTCTTGAGTTCAAGGAGGCCTCTCTCTTGTAC
AAGGACGGCGATGCCAGATCACCACCAAGGAGCTCGCACTGTCATGCGCTCTTGGGACA
GAACCCCTCGAGTCAGAGCTCAGGACATGATTAACGAGGTGATGCCACAACAACGGAT
CCATCGATTCCCTGAGTTCTTACCATGATGGCACGAAAAATGAAGGACACCGATTCCGAGG
AGGAGATCCGAGAGGCTTCAAGGTCTCGACCGCGATAACAAACGGCTCATCTCAGCTGCC
GAGCTGCGCCATGTCATGACATCTATTGGCGAGAAGCTCACCACGACGAGGTCGATGAGAT
GATTGCGAGGCCGATCAGGATGGTATGCCATTGATTACAACGAGTTGTCCAGCTCAT
GATGCAAAATAA

Appendix S3. *fas1* resulting coding sequences retrieved from the genomic sequences

>*Trichoderma* sp. T214

ATGCGTCCCAGTCGAGCAAGAGCTCGCCACACGCTCCTCGAGCTTCTGGCATACCA
GTTCGCCTCTCCCGTGAGGTGGATTGAGACCCAGGATGTCTTCTGGCAGAGAGAACAGCCG
AGCGCATCGTCGAAGTCGGCCCCCGAGATAACCTTGGAGTCATGGCAAAGCGCACACTGGCA
TCCAAGTACGAGGCCATCGACGCCAACGTCGGTTAGCGACAGATCCTCTGCTACAACAA
AGACGCCAAGGAGATCTACTATGATGTAGACCCGTTGAGGAAGAGCCCGAGCCTGTTGCCA
GCTCATCCGCTGCTCCCACAAGTCACCCGCTGCTGCCAGCGCTCTGCCGGCTGCTGCCA
GTTGCTGCTGCTCCGCTGCCGGCCTGGACCTGCAAGCTCAGGTTCCGATGTCCTGCTGCCA
GCTGTCGAGATTGTCACACCTTGGTTGACAGAAGCTCAAGAAGTCTCTAGCTGATGTGCC
CTTAGCAAGGCCATCAAGGATCTGGTTGGAGGCAAATCCACTCTCAAAATGAGATTCTGGT
GACCTTGGAAAGGAATTGGCTCAACACCTGAGAAGCCCGAAGATAACACCTCTGACGAGCT
CGCCGCCTCATGCAAGCTACCTTGGATGGCAACCTTGGCAAGCACACAGTATCCCTCATCGC
TAGGCTCATCTCTCAAGATGCTGGTGGCTCAACATCACAGTGGCCAGGAATATCTGGA
AACAAAGATGGGGTCTCGGCCCCGGCAGGCAAGACGGAGCATTGCTGCTGGCTTGACCATGG
AGCCTACTGCCCGTCTAGGAAACGACGGTGAGGCAAGGGATTCTCGACGGAATTGTCCAG
AAATACGCCGAGCGCCGGTATCAGCTGACATCTGCAAGCTGCTCTGGTGGCTGAAAGGA
GGTGGGGAGGCATGATGATGGATCCGCCATCGATGCTCTCACCAAGGACCGAGCGCG
CCTGTTCAAGCAGCAGTTGGAGCTTGTCCAGATACTCAAGATGGACCTCCGGCCGGCG
ACAAGGCACACATTGACTCCCAGAAATCAGAAAAGTTCTCAAGCTCAGCTGGATGTGG
ACCGCCGAGCACGGGATTCTACGCTTCTGGTATCGAGGCTGCTTACGGCTCAAGGCT
AGATCCTACGACTCCTCTGGAACTGGCTCGCAGGATGCGCTCACCATGTTCTACGACATT
ATCTTGGTAGACTCAAGACCGTCGACCGTGAAATTGTCAGCCAGTGCATTGCACTATGAAC
AACTAAATCCAAGCTCCTCGAGTTATGCACTGACAACTGCCCTACTGAGCG
GGTAGAGACCTACAGCTGGCAAGGAGCTGGTACAGAGCTCATGAGAAACTGCAAGGACG
TTCTGAACCTTCCCCGTCTACAAGGACGTTGCTGCTGCCCCACTGGCTCGCACCACTGTGG
ACGCTGTTGCAACCTCAACTACGAAGAGGTTCCCCGCGCAGCTGCAGAAAGCTCGAGCA
CTACGTTCAAGCAGATGGCTGAGGGTGGAAAGATTCCAGTATGGCAACCGCACCAAGGTGC
AAAACGACCTGAGCGTATCTACAAGTTGATCAAGCAGCAGCACAAAGATTCCAAGACTTCT
CAGCTCGAGATCAAGAGCCTGTACGGCGATGTTCTGCGCTCTGCCATGAACGAGAGCCA
GATCCTCCCCAAGGAGAATGGCAAGGGCGAAAGCCAGGTCTCAAGGGCACAAACCCCAAC
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TCACCTCCAGGACAAGTATGTCCTGATGACTGGTGGCGGTGCCATTGGTGGCGAGG
TCCTGCAAGGCCTCATCGCGAGGTGCCAAGGTTGGTACCAACTAGCCGATTCTCTGAC
AGGTGACCGAGTACTACCAAGTCCATGTACAGCCGCTTGGCTCGCGGATCGCAAATCGTC
TCGTTCCCTCAACCAGGGAAAGCAAGCAGGATGTTGGAGGCTTGTCAACTACATACGATG
CCAAGACCGGTCTGGCTGGATCTGACTTCATTGTTCCCTCGCTGCCATTCTGAGAACG
GCCGACAGATTGATAGCATCGATTCAAGTGGAGCTGGCTCACCGTATTATGCTCACCAACT
TGATCCGATGCTGGCTGCGTCAAGACTCAAAGGCTGAGCGTGGCTCGAGACCCGTCCC
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GAGTCCAAGCTGGCTTTGAGACCCCTTCAACCGATGGCACTCCGAGGACTGGGGCCACTA
CCTCACCATGCGGTGCTGTCATGGACTCGTGGTACTGGTCTCATGTCAGGCAACAA

CGTTGTTGCTGAGGGTGTGGAGGCCTGGCGTGCACCTCTCCAGCAAGAGATGGCCT
TCAACCTGTTGGGCCTGATGTCAGCTACCATGTTGATCTGCCAGTCAGAGCCGCTTTG
CCGACTTGAACGGTGGCCTGCAATTCACTCCCCAACCTGAACGAGTCCATGACCAAGCTCCGC
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CACCATGTTAACGGAGCCGACTCCGAGGTTCTTACAAGAAGAAGACTATCGAGCCCCGTG
CCAACATCAAGTGTGACTCCCTCACCTGCCTGACTGGAAGACGGAGGTGGCCCTCAAC
GACCAGCTGAAGGGCATGGITGACTTGGAGAAGGTCATTGTTGACCGGTTTGCAGGAAGT
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TGCCCTGCAACCTGGGCTTACCATGATGATGTCAAGGTGGCCTCGTCCACGGCACATC
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GAAGGACATTGAGATTCTAAGGACGACACCCGGCGCCCTGTTGTTACTCTCCATGGTGAAG
CCGCCGCCGCTGCCAACGAGGCTGGTCAAGGAGGCTCGGTCTCCATTACACGCCGAC
AACGAGGCCGTTGCCAGTCAGTTGCCATTCTAA

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ATGCGTCCCTGAAGTTGAGCAAGAGCTGCCACAGCTCCTCGTTGAGCTTAGCATACCAG
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GACGCCAAAGAGATTACTACGATGTCGACCCATTGAAGAACAGAGGCCAGGCCGCTGCC
CAGCTCATCAGATGCCCGAGCCAGCCTGCTGCCGGCGCCCGCTGCTGCTCCGGCTG
CTGCTGCCCTAGCTCCCTCAGGGCCTGCAGCTCAGGTTCTGATCAGCCAGTGCAGGCT
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CAAGATGGGCCTTGGTTCTGGTAGACAAGACGGAGCACTGCTGTTGGCTTGACCATGGAG
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GAGCTTGCCACCGTATTGAGTCGACATAGGTGTGGAGACCATTGGAAAGGCAAG
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 CCATCAACATTGAGAATGACACTTTATCGAGCGCAACTCACTAGCCAGGAGATTGCTTACT
 GCAAGAGCTCTGCTAGCCCCCAGAGCTCATTGCTGGCCATGGAGCGCAAGGAGGCCAGTC
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 AAGGATGACTCTGGCGCCCTACTGTTACTCTTCATGGCGATGCGGCTGCTGCTAAACAG
 GCTGGCGTGAAGGAGGTTCCGCTCCATCTCGCATGCTGATAAGCAGGCTGTTGCCGTTGCT
 GTTGCTCACTTTAA

Appendix S4. *lcb2* coding sequences

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ATGCCCGACGGCTGCAAATCCCTCTCCAGCCAGTCTGCCATGTCAGCGTCCGAAAAG
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 CTCAATTGGCCCGCTGTTCTGAGCACTGTACACCAGCCAGGCCGGGGAAAGCCCT
 TCGAGCGCCGATTGAAGACGAGCCTCCGTACTACTATCTCTGACCACCTACCTGAGCTACT
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 GCCGCTTCATCACCCCTCAAGGACCGCAAGTCGGACGACTACAACCTACCGCTACACT
 GGCACCTACACTGAGACGCTAACATGAGTTCTACAACACTATCGGTTTGCTCAGTCTGAG
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 CTCATGCAAGATTCTGCCCTCAAGCTGATCAACGGAGATATTGCCCGGCCAGGGCGAG
 GAGCGCCTGAAGCGCATTGCCCTCAACTCTCGCTATCTCCGTCTCGGACTCAAGCGTCTCGGC
 ATGATTGTCAGGGTACGACGACTCACCCATCATTCCGTCCTGCTTACAACCCGGAAAG
 ATGCGCTGCCAGATGCTGCCAGATGCTGCCGAAAATCTCGTCGTCGTCGTTATCCTG
 CCACTCCGCTCATCAGCTCGGTGCTCGTTCTCGCTCTGCTGCCACAACAAGGATGACC
 TCGACCGAATGCTTATTGCCGTGACGAGGTTGCCGACCTTCTCCAGCTCAAGTACTCCACTG
 GAATTGCCGGCGGTTGGACCCCTCTGCCGAGGGCGTCGACCCAGAAGACGAGGCTGAGTG
 GAGAAAAGAACACTGGCACTCCATCAAGGCCCGATGGAAGGTTGAGGACGTCATTGAC
 GAGGTGCGTTGACTGCAAGCTCCCTCCGGTGA

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ATGCCCGACGGCTGCAAACCCCTCTCCCTACCCAGTCTGCTATGTCAGCATCCGAGAAG
 ACTGGTGAAAGACGTCTCGTCCAGACCAACCGACTGAGCCAGCTTCTCCAG

CCCCAAAGTCCAAGGACCAGTCACCACTCGCCGCCATGCCGAGGCCAGATGAACACCATCTCGCGTCGACGGTGTGCTGCCACCAATTGCGCTGACGGCAGTGGCGAGGCCAACAGCATCGAGCCAACCAAGATGCTGTTCAAGCCGAATCGGCCAGGAGCAAGAGCAGCGACGACTCGCCAGTCTCAATTGGCCCCCTGCTTGATCCGGCCCACCGCTATGTCAGCCAGTCCAATGCGGAGGAGTTCAAGGAGCCGATTGAGGATATGCCCTCGTACTTCTATCTCTGACCACCTACCTGAGCTATCTCTTGTGATCATGATGGGCCATTGTCGCGATTACTTGGCAAGCGCTCGGAGATAAAAAAGCGCTACGACCCCCCTCAAAGTGCAGAACGGGTTGCCCTCTCACTGATGATTTCGACAGCTTCTACACTCGTCGGCTGAAGGGTCGTTGGATGACTGCTTGCTGCCCTACTTCGGCTCCCGGTGCGTACATTACTCTAAGGAACGTACGGCAGACAGGCTTAACCGCAACTACCAACTACACTGGAAACCACGTCGAGACACTCAATGTGAGCTCTACAACACTACCTCGGTTTGCTCAATCGCAGGGCCCCCTGTGCCGATGCTGTCGATGAATGTGCAAGAAGTACGGTGTACCGCTGCAAGGCCCGCGTGGCGATAGCGGCACCTCCGACCTGGCCCTCGAAGGTTGAACCGCGAAGGTTGCAACCTTGTGCGAAAGCCAGAGGCCATGGTCTTCTCTATGGGTTACGTTACCAACTCCAGTACCTTCCCCGCTCTCGTGTCAAAGGGCTGCCGTGATTGTCGACGAACTGAACCATGCCCTCATTCTGTCGGTCTCGTCTCAGTGGCCTGTTATCCAGTCTTCAAGCACAAACGACGTGGCCGCTGGAGCGCGTCTCGTGAAGCCATCTCCAGGGCCAGCCAAGGACTCACCGCCCCCTGGAAAGAAGATTCTCGTCGTTGTAAGGGTCTCTATTCTATGGAAGGCACTATGGTTAATCTGCCGGCAATTGTGGCTCTCAAGCGCAAGTACAAGTTCTACCTGTACGTTGACGAGGCTCACTCCATTGGTGCCCTGGGACCCCGTGGCGTGGTGTGATTACTTGGTATTGATCCCTCCGAGGTTGACATCCTCATGGGACCCCTGACCAAGTCCTTGGCGCAACGGAGGCTACATTGCGGAGAGAACACATCATTGACAAGCTCCGTGCCACCAACGTCGCTCAATCTATGGCGAAGCCCCCTCTCTCTTGTGCTCATGCAAGATTCTTACCTCGATCAAGCTGATTAACGGCGATATTCCCCCTGGCCAGGGCGAGGAGCGTCTCCAGCAATTGCTTCAACTCTCGCTATCTCGCCTTGGACTTAAGCGCTTGGCCTCATCGTTGCTGGCTCTGATGATTCTCCATTATCCCCGTTCTGCTGTACAACCCCTGGAAAGATGCCTGCCTTAGCCGCAAATGCTCAAGCGCAACATCTCTGTTGTCGTCGTTGGTACCCGGCTACCCCGCTCATCAGCTCGCGTGGCCGTTCTGCATCTCTGCCGCTCACAAACAAGGATGACCTTGACCGCATGATCAGAGCTTGCAGCAGGGTCGGCGAATTGCTCAGCTCAAGTCTCTTCAGGCAATTGCTGCTGAGCACCAGAAAACGAGGCTGAGTGGAGGAGGGCCAACAATGTCGCCATCAAGCCTCTCGATGGGATGTTGAAGAGGTCATTGACGAGGTGCCATGCAAGCTTCCCTGCGATAA

Appendix S5. *tef1* resulting coding sequences retrieved from the genome

>*Trichoderma*_sp._T214

ATGGGTAAGGAGGACAAGACTCACATCAACGTGGTCGTTATCACCCTGACTTGATCTACCAG
TGCAGGTGGTATCGACCGTCGTACCATCGAGAAGTTCGAGAAGGAAGGCCGCGAACTCGGCAA
GGGTTCTTCAAGTACGCTTGGTTCTTGACAAGCTCAAGGCCGAGCGTGAGCGTGGTATCA
CCATCGACATTGCTCTGTGGAAGTTCGAGACTCCCAGTACTATGTCACCGTCATTGACGCTC
CCGGCCACCGTGATTTCATCAAGAACATGATCACTGGTACTTCCCAGGCCGATTGCCCTATCCT
CATCATGCCGCCGTACTGGTGAGTTGAGGCTGGTATCTCCAAGGATGCCAGACCCGTGA
GCACGCTCTGCTGCCAACCCCTGGGTGTCAGCAGCTCATTGTTGCCATCAACAAGATGGA
CACTGCCAACTGGGCCAGGGCTGTTACCAGGAAATCATCAAGGAGACTTCAACTTCA
AGAACAGTCGGCTTCAACCCCAAGGCTGTTGTTCTGCCCCATCTCCGGCTCAACGGTGAC
AACATGCTCCAGCCCTCCACCAACTGCCCTGGTACAAGGGCTGGGAGAAGGAGACCAAGG
CTGGCAAGTTCACCGCAAGACCCCTCTGAGGCTATCGACTCCATCGAGCCCCCAAGCGT
CCCACGGACAAGCCCTCCGTTCCCTCCAGGATGTCATAAGATCGTGGTATTGGAACA
GTTCCCGTGGCCGTATCGAGACTGGTATCCTCAAGCCGGTATGGTCGTACCTTCGCTCCCT
CCAACGTCACCAACTGAAGTCAAGTCCGTCGAGATGCAACCACGAGCAGCTCGTGAGGGTGT
CCCAGGTACAACGTTGGTTCAACGTCAAGAACGTTCCGTTAGGAAATTGCCGTGGTAA
CGTTGCCGGTGAUTCAAGAACGACCCCCCATGGTGCCGCTTCTTCAACGCTCAGGTCA
CGTCATGAACCACCCCTGGCCAGGTGGTGGCTACGCCCGTCTTGAUTGCCACACTG
CCCACATTGCCCTGCAAGTTCGCCAGCTCCAGGAGAAGATCGACCGCCGTACCGTAAGGCT
ACCGAGACTGCCCAAGTTCATCAAGTCCGGTGAUTGCCATCGTCAAGATGATTCCCTCC
AAGCCCATGTGCGTTGAGGCTTCAACGACTACCCCTCCCTGGGTGTTGCGCCGTCGTGAC
ATGCGACAGACCGTCGCTGCGTGTCAAGGCTGTCGACAAGTCCGCCACCGCTGG
CAAGGTACCAAGTCCGTCGCCAAGGCCACCAAGAAATAA

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ATGGGTAAGGAGGAGAAGACTCACATCAACGTCGTCATCAGTGGTCACTTGATCTACCAG
TGCGGTGGTATTGACAAGCGTACCATCGAGAAGTTGAGAAGGCCCGAACCTGGCAA

GGGTTCTTCAAGTATCGTGGTTCTTGACAAGCTAAGGCCAGCGTGAGCGTGGTATCAC
 CATCGACATTGCCCTCTGGAAGTCGAGACTCCCAAGTACTATGTCACCCTCATTGACGCTCC
 CGGCCACCGTATTTCATCAAGAACATGATCACTGGTACCTCCCAGGCTGACTGCCTATCCT
 GATTATCGCTGCCGGTACTGGTGAGTCGAGGCTGGTATCTCCAAGGATGCCAGACCCGTGA
 GCACGCTCTCGCCTACACCCCTGGGTGCAAGCAGCTCATCGTGCCTCAACAAAGATGGA
 CACTGCCAAGTGGCTGAGGCTCGTTACCTTGAGATCATCAAGGAGACCTCAACTCATCAA
 GAAGGTCGGCTCAACCCCAAGACCGTTGCCCTCGTCCCCATCTCCGGCTCAACGGTGACA
 ACATGTTGGCCGCTCCACCAACTGCCCTGGTACAAGGGCTGGGAGAAGGAGACCAAGGC
 TGGCAAGTCCACCGCAAGACCCCTCTCGAGGCCATTGACGCCATTGAGCCCCCAAGCGTC
 CCACAGACAAGCCCCCTCGTCTGCCCTCAGGATGTTACAAGATCGGTGGTATCGGAACAG
 TCCCTGCGCCGTATCGAGACTGGTGTCTCAAGCCGGTATGGTGTACCTTCGCTCCCTC
 CAACGTCAACCCTGAAGTCAAGTCCGTTGAGATGCACCACGAGCAGCTCGTGAGGGTGTCC
 CCGGTGACAACGTTGGTTCAACGTCAAGAACGTCTCCGTCAAGGATATCGCCGTGGTAAC
 GTTGGCGGTGACTCCAAGAACGACCCCCCATGGTGCCGCTTCAACGCCAGGTACATC
 GTCATGAACCACCCCTGGCCAGGTGGTCCGGAACGCTCCGCTCGATTGCCACACCGCC
 CACATTGCCCTGCAAGTTCTGAGCTCCTCGAGAAGATCGACCCGGTACCGTAAGGCTACT
 GAGGCCTCCCCAAGTTCATCAAGTCTGGTACTCCGCCATGTCAAGATGGTCCCTCCAAG
 CCCATGTGCGTTGAGGCCTCACCGACTACCCCTCCCGTGGTGTCCGTGACATG
 CGCCAGACCGTCGCCGCGTGTCAAGTCCGTCAGAACGACTGGTGTACCCGCAA
 GGTACCCAAGTCCGCCAAGGCCAAGAAATAA

Appendix S6. *rpb2* resulting coding sequences retrieved from the genome

>*Trichoderma* sp. T214

ATGGCTGACTACGAGGACGAATACGACTACGAGAACTACGGAGACGAGGATGAAGGCATCAC
 TCCCGAGGATTGCTGGACTGTCATCTCCTCCTCTCGAAACCAAGGGTCTCGTATCGCAGCA
 GACCGACTCTTGACGAATTCACCCAGACGACGATCAGGACCTCGTAAACGAATACTCCAC
 CATCACACTCGACCAGCCAAACCCCTCCTCGCCACCCGGCCGAACAATAGCCCTCGTCGATA
 TGAGATCAAGTTGGAAGCGTCATGGTGTACGTCCCCTACTATCAGTGAGACGGACGGCACTG
 TGACTTCTCTGCTCCCTACGAGTGGCAGACCGTAACCTGACCTACGCCAGTCCCTCTATAT
 CAAGATCACTAAAAAGGTGCGGCTGCTGTTGAGAGGGAGGTTCCCTGCACGAAATGGACG
 ATGCTCAGCAGGAGGAATACGCAAGAACCGCGAACACCCCTACAAAGCTCGAGTGGGAAGA
 GGAGGAGAACGGCGAACGATGACAACATCGCAAGTCTGATGACTGGAAGGACATGGTCTTC
 GTTGGCAAGCTGCCCATGGTCAAATCCAAGATTGTCATCTGAGCCGTGAACAGGATGAT
 AGCCTGTTCTGTCAACGAATGTCCTACGATCAGGGTGGCTACTTGTATCAACGGCAGT
 GAAAAGGTCTCATGCCAACAGAGAGATCCGCCAACATTGTCAGTGGCAAGTCTCAAGAAGGC
 CCAGCCCAGTGCCTATACCTATACGGCTGAAATCCGAAGTGCCTGGAAAAGGGCTACGGC
 TCATCTCTAGCATGATGCTCAAGCTGTATGGCAAGGGAGACTCTGCTCGTGGTGGCTTGGCC
 AGACTATCCACACCACCCCTGCCCTTGTCAAGTCAGATCTTCCCGCCATTGTCTCCGTGC
 CCTGGGTGCTTCTGATGAAGATATCCTCAACCACATTGCTACGCCAACAGACAGCCA
 GATGCTGGAGATGCTCGACCTTGTATTGAGGGAGGCCTCTGTGTCCAGGACCGAGAGGTTGC
 TCTGGATTTCATGGAAACGCGGAAACCGAGACCAAGCTGGTCTCGACGCGAGAACGCT
 GTCCCGTGGCTAAGGATATCCTCAGAAGGAGACTCTTCCCGCCATTGTCTCCGTGC
 AAGTCAAACCCAGAAAGGATTTCTGGGATACATGGTGCACAAGCTGGCAATGTGCGCT
 CGGAAGAACGAGAGCCGACGATCGTGCACACTTGGAAAGAACGCTCTGGATCTGGCGGGT
 CCCCTGCTGGCAAAGCTGTTCCGTGGTATCATGCGAAGGATGAAACACTGAGTTGGCAACTAT
 CTGAGACGATCGTCGAGGGCAACCGACACTCAACCTGCTGTGGGTATCAAGCCGGCAC
 GCTTCAAACGGATTGAAGTATTGCTTGCCACAGGAAACTGGGGTATCAGAACAGGCCA
 TGAGCTCAACTGCAGGTGTGCTCCAGGTGCTAACCGTTACACGTTGCTCGACCCATCAC
 ATTGCGTCGTAACAAACTCCTATCGGAAGAGATGGTAAGCTCGAAAGCCTCGACAGCTTC
 ACAACACGCACGGGGTTGGTCTGCCGGCGAGACACCCGAGGGACAGGCTGTGGTCT
 GGTCAAGAACTTGTCTTGATGTGTTACGTCACTGTCGGTTCTCCCTCGAACCTCTGATTGA
 GTTCATGATCAACAGAGGTATGGAAGTCGTTGGAAGAGTACGAGCCGCTCGGTATCCTCATGC
 TACAAAGATTGGTGAACGGTGTCTGGGTTGGAGTCACCAAGACCCCTAACGACTTGGTGA
 ACCAGGTCTGGACACTCGTCGCAAGTCCTATCTGCAATACGAAGTCTCTCGTGAGAGAA
 ATTGAGACGAGGAATTCAAATCTTCCGACGCGAGGCCGTGTAATGCGGCCAGTCTTAC
 GTTCAGCAGGAAGATGATCCGAAACGGCATCAACAAAGGGCACCTGGTATTGACCAAGGA
 GCTCGTCAATAGATTGGCAAGGAGCAGGCTGAACCTCCGGAAAGACCCAGCATGAAGATTG
 GATGGGAGGGATTGATTAGGGCTGGTGCCTGAATATCTGACGCCAGGAAGAGGGAGACG
 TCCATGATCTGCGATGACGCCAGAGGATCTGAGCTGTATCGTCTCAGAACGGCTGGTATTAAC

ACTGAGGAAGACATGGGAGATGACCGAACAAAGCGACTAAAGACCAAGACAAACCCGACTA
CTCACATGTACACCCATTGCGAGATTCAACCAAGTATGATCTTAGGCATCTGTGCTAGTATCATT
CCTTCCCCGATACAACCAGTCCCCCGTAACACTTACCAATCTGCCATGGTAAGCAAGCT
ATGGGTTTCTTCCTCACAAACTATTCCCGCGCATGGACACCATGCCAACATTCTACTACC
CCCAGAAGCCGCTGGTACCACTCGATCCATGGAGTTTGAGATTCCGAGAGTTGCCTGCTG
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GAACCAAGAGTAGTATTGACAGAGGTCTTCCGAAGTCTCTTCCGATCATACTCGGATCA
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GTGTCCGAACGACCAAGATTCCCTCAGATTGGAGACAAGTTCGCCCTCGTCACGGTCAAAG
GGCACTATTGGTGTACTTACCGACAAGAGGACATGCCCTCACAAGAGAAGGTGTCACTCC
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CTCCTAAGTAAGGTCTCACGCTGGAGGTATGGAGGGTATGCCACCCCTCACGGATGTC
ACCGTCGACTCGGTCTCAGAGCTGTGCAAGACGGCTACCAAGTCACTCGTGTGCCC
TATGTACAATGGCCACACAGGCCGAAGCTGAGAGGCCAGGTGTTGGACCAACATACTA
CCAGCGACTCCGTCACATGGTGGACGACAAGATCCACGCTCGTGTGCCC
TTATGACAAGACAACCGTGGAGGGTGTGCCAGAGATGGTGGTCTCCGATTGGAGAAATG
GAACGTGATTGTATGATTGCTCACGGTGCCTCGCTTCAAGGAGCGATTGTTGAGGTG
TCAGACGCCCTCCGAGTTCACATTGCGAGATTGTGGACTCATGACGCCATTGCCAACCTC
TCTAAACAATCGTCAGTGTGACCTTGTAAAGAACAGACCAAGATTGACAGATTACATC
CCTTATGCCCAAGCTCCTGTTCCAGGAACCTCAGTCGATGAACATTGCCATGTC
ACAAACCGGTCTGGCGCCATCAGGTAA

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ATGGCTGATTACGAAGACGATTACGACTATGAGAACTATGGGATGAAGATGAGGGCATCAGC
CCCGAGGATTGCTGGACTGTGATTCCCTCTTCGAAACCAAGGGTCTGTATCGCAGCAG
ACCGACTCCTTGACGAATTACCGCAGACGACAATCCAGGATCTGTCACGAATATTCCACC
ATCACACTCGATGCCAATCCTCCTCGCACCTGGTCAACGATAGCCCTCGCGATATG
AAATCAAATTGGAAGTGTATGGTACCGTCCACTATCAGTGGAGACGGATGGAACGTG
CATCTGCTCCCTACGAATGCCAGACCGTAACCTGACTTACGCCAGTCCGCTTACATCAA
GATCACAAGAAGGTGTCTCGGCCGTGAGAGGGAGGTTCCGCTGACGAGATGGACGAT
GCCCAACAAGAACAGTATGCAAGGACGGAGAAAACCCCACAAAGCTGGAATGGGAGGAG
GAAGAAAATGGCAAGACGACAATCTGGCAAGTCTGATGACTGGAAGGACATGGTTTCTG
TGGCAAGCTGCCCATCGTAAAGATTGTCATCTGAGCCGTGAACAGGATGACAG
CCTGTTCTCGTCAACGAGTGCCCTACGACCAAGGAGGCTACTTGTATCAACGGTAGTGA
AAAGGTCTCATGCCAAGAGCCTCCGCCAAACATTGTCAGGTCTCAAGAACAGGCC
AGCCCAAGTGCCTACACTGCTGAAATCCGAAGTGCCTGGAAAAGGGATCTGACTC
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GCGAGACCAAGAACGGATTCTCCTGGATACATGGTCACAAGCTGTCATGCGCACTC
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TCGGTCTGGTGCATCTGTTGGTAG