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Supplementary Fig. S3. ITS tree with *Chrysothrix fagicola*. GTRG model with gamma distribution was used as a model of sequence evolution. Trees were constructed using Bayesian inference run for 597,000 generations.

Coniocarpon cinnabarinum MN734117

0.6 — Andreiomyces morozianus KF740836
— Andreiomyces morozianus KF740837
0.64 — Andreiomyces morozianus KF740838
— Andreiomyces morozianus KF740839
— Andreiomyces morozianus KF740840
0.67 — Andreiomyces obtusaticus AF517896
— Andreiomyces obtusaticus PRA-Vondrák26191 OQ717300

0.63 — Chrysothrix fagicola Malíček13300 OQ717368, Malíček13310 OQ717369
— Chrysothrix fagicola Malíček14013 OQ717370
— Chrysothrix fagicola Malíček13296 OQ717367
— Chrysothrix fagicola Malíček14081 OQ511312

Chrysothrix candelaris PRA-Vondrák24317 OQ717814

0.88 — Chrysothrix sp. (sub Lepraria aff. obtusatica) JQ673478
— Chrysothrix xanthina MH714516

1 — Arthonia mediella TYPE1 PRA-Vondrák24180 OQ717698, PRA-Vondrák24183 OQ717699, PRA-Vondrák25623 OQ717700
— Arthonia mediella TYPE2 PRA-Vondrák23828 OQ717701
— Arthonia mediella TYPE3 PRA-Palice32406 OQ717702

0.64 — Felipes leucopellaeus PRA-Vondrák24395 OQ717838

Zwackhia viridis PRA-Vondrák24383 OQ718154

S3