

**S2 Table. Dominant fungal species significantly different between the two nematode species regardless of the soils (average ± SEM).** Only species with a relative abundance > 0.1% are displayed. Significance was inferred using a negative binomial regression and generalized linear model fit (nbGLM). A Likelihood ratio test was used, complemented with a post-hoc false discovery rate multiple correction test (LHR FDR,  $P < 0.05$ ). Analysis was done using the EdgeR package.

Phylum	Family	Species	Pratylench	Meloidogyne	Adj. FDR	
Ascomycota	Ascomycota (unid.)	SH212862.07FU	-	0.49±0.49b	3.91E-05	
	Ascomycota (unid.)	SH210351.07FU	-	0.49±0.47b	1.77E-04	
	Chaetomiaceae	<i>Chaetomium jatropheae</i>	-	1.37±1.25b	9.27E-05	
	Chaetomiaceae	<i>Hemicola grisea</i>	-	0.13±0.11b	2.04E-03	
	Chaetomiaceae	<i>Chaetomium aterrimum</i>	-	0.3±0.3b	4.17E-03	
	Cordycipitaceae	<i>Cordyceps bassiana</i>	-	0.3±0.3b	6.64E-03	
	Davidiellaceae	<i>Cladosporium exasperatum</i>	0.03±0.01a	9.16±3.01b	1.77E-04	
	Dothideomycetes	unidentified	-	1.15±1.15b	7.14E-05	
	Dothioraceae	<i>Aureobasidium pullulans</i>	-	1.91±1.25b	5.59E-05	
	Helotiales (i.sedis)	<i>Tetracladium maxilliforme</i>	0.01±0.01a	5.2±4.57b	7.14E-05	
	Helotiales (i.sedis)	unidentified	-	0.92±0.91b	3.41E-03	
	Hyaloscyphaceae	<i>Mycoarthris corallina</i>	-	0.48±0.47b	5.05E-05	
	Hypocreales (i.sedis)	<i>Myrothecium roridum</i>	-	0.11±0.11b	0.013	
	Mycosphaerellaceae	<i>Pseudocercosporaella</i>	-	0.41±0.41b	5.05E-05	
	Nectriaceae	SH219102.07FU	0.01±0.002	1.9±1.89b	9.27E-05	
	Nectriaceae	SH220702.07FU	0.02±0.01a	6.15±3.52b	2.12E-04	
	Oxygenaceae	<i>Chrysosporium lobatum</i>	-	0.31±0.31b	5.55E-03	
	Pezizales (i.sedis)	<i>Cephaliophora tropica</i>	-	0.63±0.63b	3.17E-03	
	Plectosphaerellaceae	<i>Lectera longa</i>	-	1.54±1b	3.82E-05	
	Plectosphaerellaceae	<i>Plectosphaerella alismatis</i>	-	0.18±0.11b	3.61E-03	
	Pleosporaceae	<i>Stemphylium herbarum</i>	-	5.08±2.6b	3.91E-05	
	Pleosporaceae	<i>Alternaria alternata</i>	0.02±0.01a	10.95±7.36	6.20E-05	
	Pleosporaceae	<i>Chalastospora ellipsoidea</i>	-	0.38±0.33b	1.77E-04	
	Pleosporales (unid.)	unidentified	-	0.9±0.9b	3.61E-03	
	Pyronemataceae	unidentified	-	0.49±0.49b	3.61E-03	
	Pyronemataceae	<i>Pseudaleuria</i> sp.	-	0.39±0.39b	5.55E-03	
	Saccharomycetales	<i>Candida tropicalis</i>	0.01±0.01a	3.65±3.64b	7.14E-05	
	Sordariales (unid.)	unidentified.	-	1.3±0.83b	1.00E-04	
	Sordariomycetes	unidentified	0.01±0.003	0.75±0.52b	4.17E-03	
	Trichocomaceae	<i>Penicillium chrysogenum</i>	-	3.51±2.31b	4.57E-05	
	Trichocomaceae	<i>Penicillium aurantiogriseum</i>	-	2.14±2.09b	7.34E-05	
	Sordariomycetes	unidentified	0.01±0.003	4.59±2.63b	2.12E-04	
	Tremellales (i.sedis)	<i>Cryptococcus heimaeyensis</i>	-	2.75±2.74b	1.54E-05	
	Tremellales (i.sedis)	<i>Hannaella oryzae</i>	-	3.47±3.47b	5.05E-05	
	Chytridiomycota	Rhizophlyctidaceae	<i>Rhizophlyctis rosea</i>	-	0.12±0.12b	0.012
	Fungi (unid.)	Rhizophydiaceae	<i>Rhizophydium</i> sp.	-	0.18±0.18b	6.64E-03
	Fungi (unid.)	unidentified	-	0.3±0.3b	5.62E-05	
Ascomycota	Davidiellaceae	<i>Cladosporium haerospermum</i>	1.04±1.04b	-	3.82E-05	
	Davidiellaceae	<i>Cladosporium halotolerans</i>	1.54±1.53b	-	1.51E-03	
	Hypocreaceae	<i>Trichoderma asperellum</i>	0.96±0.96b	-	5.50E-05	
	Hypocreales (i.sedis)	unidentified	0.47±0.47b	-	3.91E-05	
	Saccharomycetales	<i>Debaryomyces nepalensis</i>	0.9±0.9b	-	3.91E-05	
	Taphrinaceae	<i>Taphrina padi</i>	0.65±0.49b	-	3.82E-05	
	Trichocomaceae	<i>Penicillium brevicompactum</i>	0.21±0.21b	-	5.44E-03	
	Trichocomaceae	<i>Aspergillus niger</i>	63.35±4.76	1.77±1.72a	8.49E-03	
	Trichocomaceae	<i>Penicillium novae-zeelandiae</i>	0.12±0.12b	0.01±0.01a	0.037	
	unidentified	<i>Pleosporales</i> sp.	2.12±2.11b	0.01±0a	2.96E-04	
Basidiomycota	Xylariaceae	<i>Hypoxyylon howeanum</i>	0.37±0.37b	-	5.62E-05	
	Malasseziaceae	<i>Malassezia SH176396.07FU</i>	2.77±2.61b	-	5.47E-05	
	Malasseziaceae	<i>Malassezia SH176395.07FU</i>	0.31±0.31b	-	4.17E-03	
	Marasmiaceae	<i>Baeospora myosura</i>	1.7±1.69b	-	1.07E-03	
Zygomycota	Mortierellaceae	<i>Mortierella alpina</i>	1.47±1.46b	-	4.57E-05	