

## Supplementary material

# News from the News from the Sea: a new genus and seven new species in the pleosporalean families Roussoellaceae and Thyridariaceae

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**Table S1.** The eight variable sites detected in nrITS region among *P. dematiacea* and the neighbor species

Position								
Species	62	86	87	98	107	112	257	365
<i>P. dematiacea</i>	C	C	A	A	T	T	T	C
<i>C. goaensis</i>	T	T	G	G	C	G	C	T
<i>C. leucanae</i>	T	T	G	G	C	G	C	T
<i>L. muriformis</i>	T	-	G	G	C	G	C	T
<i>T. mahakoshae</i>	T	T	G	G	C	G	C	T
<i>T. mangrovei</i>	T	T	G	G	C	C	C	T

**Table S2.** The single variable site detected in nrLSU region among *P. dematiacea* and the neighbor species

Position	
Species	662
<i>P. dematiacea</i>	A
<i>C. goaensis</i>	G
<i>C. leucanae</i>	G
<i>L. muriformis</i>	G
<i>T. mahakoshae</i>	G
<i>T. mangrovei</i>	G

**Table S3.** The five variable sites detected in nrITS region among *P. dematiacea* and the neighbor species

Position					
Species	245	326	341	342	615
<i>P. dematiacea</i>	T	G	C	T	T
<i>C. goaensis</i>	C	A	T	C	C
<i>C. leucanae</i>	C	A	T	C	C
<i>L. muriformis</i>	C	A	T	C	C
<i>T. mahakoshae</i>	C	A	T	C	C
<i>T. mangrovei</i>	C	A	T	C	C

**Table S4.** The six variable sites detected in the TEF1 $\alpha$  partial gene among *P. dematiacea* and the neighbor species

Position						
Species	301	316	323	529	571	616
<i>P. dematiacea</i>	T	C	C	C	T	G
<i>C. goaensis</i>	C	G	T	T	C	A
<i>C. leucanae</i>	C	G	T	T	C	A
<i>L. muriformis</i>	C	T	A	T	C	A
<i>T. mahakoshae</i>	C	G	A	T	C	A
<i>T. mangrovei</i>	C	T	A	T	C	A

**Table S5.** The six variable sites detected in nrITS region among *P. tyrrhenica*, *P. flabelliae* and their neighbor species

Position						
Species	60	163	164	419	458	473
<i>P. tyrrhenica</i>	C	T	C	T	A	A
<i>P. flabelliae</i>	T	T	A	C	A	-
<i>P. percutanea</i>	A	A	A	C	C	G
<i>P. ramulicola</i>	A	A	A	C	C	G
<i>P. robiniae</i>	A	A	A	C	C	G

**Table S6.** The eight variable sites detected in the nrLSU region among *P. tyrrhenica*, *P. flabelliae* and their neighbor species

Position								
Species	784	786	116	1118	1137	1190	1392	1478
<i>P. tyrrhenica</i>	C	C	C	C	C	G	T	G
<i>P. flabelliae</i>	T	T	T	T	T	G	C	A
<i>P. percutanea</i>	T	C	C	C	C	A	C	A
<i>P. ramulicola</i>	T	C	C	C	C	A	C	A
<i>P. robiniae</i>	T	C	C	C	C	A	C	A

**Table S7.** The six sites detected in the in the TEF1 $\alpha$  partial gene among *P. tyrrhenica*, *P. flabelliae* and their neighbor species

Position						
Species	781	883	1009	1012	1126	1135
<i>P. tyrrhenica</i>	T	C	T	C	T	C
<i>P. flabelliae</i>	T	T	C	T	C	T
<i>P. percutanea</i>	C	C	C	C	C	T
<i>P. ramulicola</i>	C	C	C	C	C	T
<i>P. robiniae</i>	C	C	C	C	C	T

**Table S8.** The 33 variable sites detected in the in the RPB2 partial gene among *P. tyrrhenica*, *P. flabelliae* and their neighbor species

	Position																																
Species	248	278	305	353	359	384	407	413	494	496	509	518	557	581	593	612	623	689	716	728	729	749	752	779	809	839	869	875	881	899	908	914	929
<i>P. tyrrhenica</i>	T	T	G	C	A	T	C	C	C	G	C	T	G	C	A	T	T	G	G	A	T	G	A	T	C	T	G	G	A	G	C	C	G
<i>P. flabelliae</i>	C	T	G	T	G	T	C	T	T	C	C	C	G	A	A	T	C	G	G	A	T	G	G	C	C	C	G	G	A	G	C	T	G
<i>P. percutanea</i>	C	C	A	A	G	C	T	T	T	C	A	C	A	A	T	C	C	T	C	T	C	A	G	C	G	C	A	A	G	A	T	T	C
<i>P. ramulicola</i>	C	A	C	T	G	C	T	T	T	C	A	C	A	A	C	C	C	C	A	T	C	A	G	A	T	C	A	A	G	C	T	T	C
<i>P. robiniae</i>	C	A	A	T	G	C	T	T	T	C	T	C	A	A	G	C	C	C	A	T	C	A	G	A	A	C	A	A	G	A	A	T	C

**Table S9.** The two variable sites detected in nrITS region among *R. mediterranea*, *R. padinae*, and the neighbor species

Position		
Species	54	155
<i>R. padinae</i>	A	T
<i>R. mediterranea</i>	A	T
<i>R. neopustulans</i>	T	A
<i>R. kunmingensis</i>	C	A
<i>R. chiangraina</i>	T	G

**Table S10.** The single variable site detected in nrLSU region among *R. mediterranea*, *R. padinae*, and the neighbor species

Position	
Species	58
<i>R. padinae</i>	T
<i>R. mediterranea</i>	T
<i>R. neopustulans</i>	C
<i>R. kunmingensis</i>	C
<i>R. chiangraina</i>	C

**Table S11.** The six sites detected in the in the TEF1 $\alpha$  partial gene among *R. mediterranea*, *R. padinae*, and the neighbor species

Position						
Species	311	360	366	408	597	695
<i>R. padinae</i>	A	A	T	A	A	T
<i>R. mediterranea</i>	A	A	T	A	A	T
<i>R. neopustulans</i>	C	G	G	G	T	C
<i>R. kunmingensis</i>	C	G	G	G	G	C
<i>R. chiangraina</i>	C	G	G	G	T	C

**Table S12.** The six sites detected in the in the RPB2 partial gene among among *R. mediterranea*, *R. padinae*, and the neighbor species

Position						
Species	180	207	244	282	321	705
<i>R. padinae</i>	A	T	T	T	T	G
<i>R. mediterranea</i>	T	G	C	G	G	A
<i>R. neopustulans</i>	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.
<i>R. kunmingensis</i>	G	G	C	C	C	A
<i>R. chiangraina</i>	G	G	C	C	C	A

**Table S13.** The eight variable sites detected in the nrITS region among *N. lignicola* and the neighbor species

Position									
Species	180	192	233	402	424	484	519	528	531
<i>N. lignicola</i>	T	C	C	C	A	C	A	T	G
<i>N. hevae</i>	C	A	G	A	T	T	G	C	T
<i>N. entadae</i>	C	A	A	A	T	A	G	C	C
<i>N. leucanae</i>	C	A	A	A	T	A	G	C	C
<i>N. solani</i>	C	A	A	A	T	A	G	C	C
<i>N. alishanense</i>	C	A	T	A	T	T	G	C	A
<i>N. bambusae</i>	C	A	A	A	T	T	G	C	A

**Table S14.** The three variable sites detected in the nrLSU region among *N. lignicola* and the neighbor species

Species	562	563	693
<i>N. lignicola</i>	T	C	C
<i>N. hevae</i>	C	A	T
<i>N. entadae</i>	C	A	T
<i>N. leucanae</i>	C	A	T
<i>N. solani</i>	C	A	T
<i>N. alishanense</i>	C	A	T
<i>N. bambusae</i>	C	A	T

**Table S15.** The eight variable sites detected in the nrSSU region among *N. lignicola* and the neighbor species

Position								
Species	520	522	526	527	528	530	531	537
<i>N. lignicola</i>	T	A	C	A	G	G	A	G
<i>N. hevae</i>	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.
<i>N. entadae</i>	A	C	G	G	T	A	T	A
<i>N. leucanae</i>	A	C	G	G	T	A	T	A
<i>N. solani</i>	A	C	G	G	T	A	T	A
<i>N. alishanense</i>	A	C	G	G	T	A	T	A
<i>N. bambusae</i>	A	C	G	G	T	A	T	A

**Table S16.** The ten sites detected in the in the TEF1 $\alpha$  partial gene among *N. lignicola* and the neighbor species

Position										
Species	423	487	492	496	552	684	705	835	836	837
<i>N. lignicola</i>	C	T	C	A	T	T	C	G	C	A
<i>N. hevae</i>	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.
<i>N. entadae</i>	T	G	T	G	C	C	T	A	G	C
<i>N. leucanae</i>	T	G	T	G	C	C	T	A	G	C
<i>N. solani</i>	T	G	T	G	C	C	T	A	G	C
<i>N. alishanense</i>	T	G	T	G	C	C	T	A	G	C
<i>N. bambusae</i>	T	G	T	G	C	C	T	A	G	C

**Table 17.** The three variable sites detected in nrITS region among *R. margidoriensis* and the neighbor species

Position			
Species	401	406	481
<i>R. margidorensis</i>	T	T	T
<i>R. tuberculata</i>	C	A	G
<i>R. thailandica</i>	C	A	A
<i>R. pseudohysterioides</i>	A	G	G
<i>R. nitidula</i>	A	G	A

**Table S18.** The 29 variable sites detected in the in the TEF1 $\alpha$  partial gene among *P. tyrrhenica*, *P. flabelliae* and their neighbor species

Position																													
Species	124	139	145	155	163	211	271	275	295	305	316	317	3892	412	427	506	508	511	613	643	685	733	745	754	760	766	767	769	781
<i>R. margidorensis</i>	T	T	T	C	T	T	G	A	G	A	C	G	T	T	C	T	T	C	T	G	T	T	T	C	A	T	T	A	A
<i>R. tuberculata</i>	C	C	C	A	C	C	C	T	T	C	A	T	C	C	T	G	C	T	C	A	C	C	C	T	T	C	C	C	G
<i>R. thailandica</i>	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.
<i>R. pseudohysterioides</i>	C	C	C	A	C	G	C	T	T	C	A	T	C	C	T	G	C	T	C	A	C	C	C	T	T	C	C	C	T
<i>R. nitidula</i>	C	C	C	A	C	C	C	T	T	C	A	T	C	C	T	G	C	T	C	A	C	C	C	T	T	C	C	C	T