

Figure S1. Phylogenetic relationship based on the sequence of the D1/D2 domain of the LSU rRNA gene of yeast species isolated from corals and zoanthids in sampling site S1 (●) and sampling site S2 (■) in the phylum Ascomycota, and their closely related yeast species retrieved from the GenBank databases. The tree was constructed with the maximum-likelihood method and the general time reversible (GTR) evolutionary model in the MEGA software version 7.0. Numbers at nodes indicate the bootstrap percentage (>50%) derived from 1000 datasets. *Schizosaccharomyces pombe* NRRL Y-12796^T (JQ689077) was used as the outgroup. The scale bar indicates an evolutionary distance of 0.02 K_{nuc} .

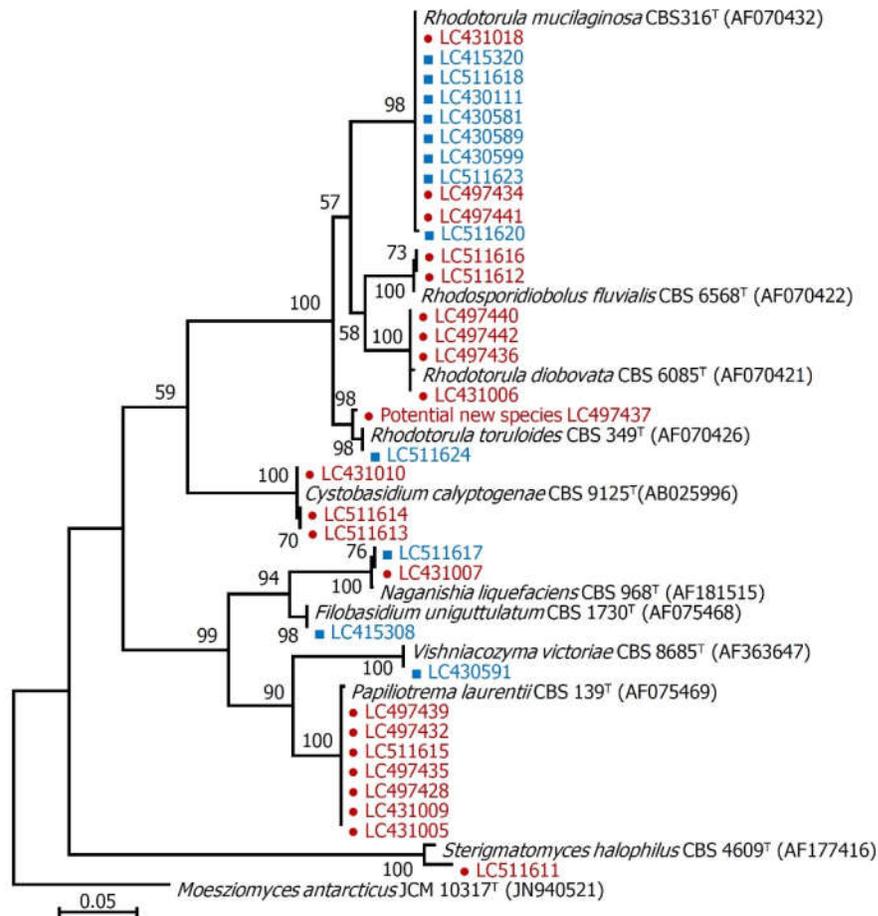


Figure S2. Phylogenetic relationship based on the sequence of the D1/D2 domain of the LSU rRNA gene of yeast species isolated from corals and zoanthids in sampling site S1 (●) and sampling site S2 (■) in the phylum Basidiomycota, and their closely related yeast species retrieved from the GenBank databases. The tree was constructed with the maximum-likelihood method and the GTR evolutionary model in the MEGA software version 7.0. Numbers at nodes indicate the bootstrap percentage (>50%) derived from 1000 datasets. *Moesziomyces antarcticus* JCM 10317^T (JN940521) was used as the outgroup. The scale bar indicates an evolutionary distance of 0.05 K_{mc}.

Table S1. Identification of yeast isolated from corals and zoanthids by analysis of the D1/D2 domain of the LSU rRNA gene and the ITS region.

Strain DMKU	Closest species (Accession no. of D1/D2; ITS)	D1/D2 domain			ITS region				Result of identification	
		Accession no.	No. of identity	No. of gap	No. of substitution	Accession no.	No. of identity	No. of gap		No. of substitution
Coral samples										
J7-1	<i>Papiliotrema laurentii</i> CBS139 ^T (AF075469)	LC431005	573/574	0	1	-	-	-	-	<i>P. laurentii</i>
JC7-2	<i>Rhodotorula diobovata</i> CBS6085 ^T (AF070421)	LC431006	532/533	0	1	-	-	-	-	<i>R. diobovata</i>
JC7-4	<i>Naganishia liquefaciens</i> CBS968 ^T (AF181515)	LC431007	555/556	0	1	-	-	-	-	<i>N. liquefaciens</i>
JC7-5	<i>Candida parapsilosis</i> CBS604 ^T (U45754)	LC431008	528/528	0	0	-	-	-	-	<i>C. parapsilosis</i>
JC8-1	<i>Papiliotrema laurentii</i> CBS139 ^T (AF075469)	LC431009	571/572	0	1	-	-	-	-	<i>P. laurentii</i>
JC9-1	<i>Cystobasidium calyptogenae</i> CBS9125 ^T (AB025996)	LC431010	565/566	0	1	-	-	-	-	<i>Cys. calyptogenae</i>
JC14-1	<i>Candida parapsilosis</i> CBS604 ^T (U45754)	LC431017	519/519	0	0	-	-	-	-	<i>C. parapsilosis</i>
JC15-1	<i>Rhodotorula mucilaginosa</i> CBS316 ^T (AF070432)	LC431018	543/544	0	1	-	-	-	-	<i>R. mucilaginosa</i>
J18-1	<i>Candida spencermartinsiae</i> CBS10894 ^T (FJ008044)	LC431020	505/505	0	0	-	-	-	-	<i>C. spencermartinsiae</i>
JC22-1	<i>Candida parapsilosis</i> CBS604 ^T (U45754)	LC497427	535/535	0	0	-	-	-	-	<i>C. parapsilosis</i>
JC24-1	<i>Sterigmatomyces halophilus</i> CBS4609 ^T (AF177416)	LC511611	585/592	6	1	-	-	-	-	<i>S. halophilus</i>
JC25-1	<i>Papiliotrema laurentii</i> CBS139 ^T (AF075469)	LC497428	606/607	0	1	-	-	-	-	<i>P. laurentii</i>
JC25-2	<i>Candida tropicalis</i> CBS94 ^T (U45749)	LC497429	529/529	0	0	-	-	-	-	<i>C. tropicalis</i>
J33-2	<i>Rhodotorula mucilaginosa</i> CBS316 ^T (AF070432)	LC497434	532/533	0	1	-	-	-	-	<i>R. mucilaginosa</i>
J33-3	<i>Papiliotrema laurentii</i> CBS139 ^T (AF075469)	LC497435	579/580	0	1	-	-	-	-	<i>P. laurentii</i>
JC33-1	<i>Rhodotorula diobovata</i> CBS6085 ^T (AF070421)	LC497436	549/550	0	1	-	-	-	-	<i>R. diobovata</i>
JC33-2	<i>Rhodospiridiobolus fluvialis</i> CBS6568 ^T (AF070422; NR077089)	LC511612	535/538	0	3	LC522501	593/598	0	5	<i>Rh. fluvialis</i>
JC33-4	<i>Cystobasidium calyptogenae</i> CBS9125 ^T (AB025996; AB025996)	LC511613	607/609	0	2	LC522502	557/562	1	4	<i>Cys. calyptogenae</i>
JC35-1	<i>Rhodotorula diobovata</i> CBS6085 ^T (AF070421)	LC497442	532/533	0	1	-	-	-	-	<i>R. diobovata</i>

Strain DMKU	Closest species (Accession no. of D1/D2; ITS)	D1/D2 domain				ITS region				Result of identification
		Accession no.	No. of identity	No. of gap	No. of substitution	Accession no.	No. of identity	No. of gap	No. of substitution	
JC35-2	<i>Cystobasidium calyptogenae</i> CBS9125 ^T (AB025996; AB025996)	LC511614	607/609	0	2	LC522503	559/564	1	4	<i>Cys. calyptogenae</i>
JC46-1	<i>Papiliotrema laurentii</i> CBS139 ^T (AF075469)	LC511615	562/563	0	1	-	-	-	-	<i>P. laurentii</i>
C322-1	<i>Filobasidium uniguttulatum</i> CBS1730 ^T (AF075468)	LC415308	564/564	0	0	-	-	-	-	<i>F. uniguttulatum</i>
3222-1	<i>Rhodotorula mucilaginosa</i> CBS316 ^T (AF070432)	LC415320	552/555	2	1	-	-	-	-	<i>R. mucilaginosa</i>
3222-2	<i>Naganishia liquefaciens</i> CBS968 ^T (AF181515)	LC511617	596/597	0	1	-	-	-	-	<i>N. liquefaciens</i>
3236-1	<i>Rhodotorula mucilaginosa</i> CBS316 ^T (AF070432)	LC511618	554/555	0	1	-	-	-	-	<i>R. mucilaginosa</i>
729-1	<i>Rhodotorula mucilaginosa</i> CBS316 ^T (AF070432)	LC430111	554/555	1	0	-	-	-	-	<i>R. mucilaginosa</i>
7231-1	<i>Rhodotorula mucilaginosa</i> CBS316 ^T (AF070432)	LC430581	550/550	0	0	-	-	-	-	<i>R. mucilaginosa</i>
7235-1	<i>Rhodotorula mucilaginosa</i> CBS316 ^T (AF070432)	LC430589	549/550	0	1	-	-	-	-	<i>R. mucilaginosa</i>
C7235-3	<i>Candida zeylanoides</i> CBS619 ^T (U45832)	LC430590	515/515	0	0	-	-	-	-	<i>C. zeylanoides</i>
C7235-4	<i>Vishniacozyma victoriae</i> CBS8685 ^T (AF363647)	LC430591	550/550	0	0	-	-	-	-	<i>V. victoriae</i>
7241-1	<i>Rhodotorula mucilaginosa</i> CBS316 ^T (AF070432)	LC430599	555/556	1	0	-	-	-	-	<i>R. mucilaginosa</i>
FC12-1	<i>Meyerozyma caribbica</i> CBS9966 ^T (AY187283)	LC511619	519/519	0	0	-	-	-	-	<i>M. caribbica</i>
FC12-2	<i>Rhodotorula mucilaginosa</i> CBS316 ^T (AF070432)	LC511620	533/534	0	1	-	-	-	-	<i>R. mucilaginosa</i>
FC12-3	<i>Candida parapsilosis</i> CBS604 ^T (U45754)	LC511621	524/524	0	0	-	-	-	-	<i>C. parapsilosis</i>
FC12-4	<i>Candida tropicalis</i> CBS94 ^T (U45749)	LC511622	523/523	0	0	-	-	-	-	<i>C. tropicalis</i>
J37-1	<i>Rhodotorula toruloides</i> CBS349 ^T (AF070426; AF444489)	LC497437	583/588	0	5	LC522504	579/588	3	6	Potential new species closest to <i>R. toruloides</i>
Zoanthid samples										
319-1	<i>Meyerozyma guilliermondii</i> CBS2030 ^T (U45709)	LC415303	542/545	1	2	-	-	-	-	<i>M. guilliermondii</i>
319-2	<i>Candida tropicalis</i> CBS94 ^T (U45749)	LC415304	544/545	1	0	-	-	-	-	<i>C. tropicalis</i>

Strain DMKU	Closest species (Accession no. of D1/D2; ITS)	D1/D2 domain				ITS region				Result of identification
		Accession no.	No. of identity	No. of gap	No. of substitution	Accession no.	No. of identity	No. of gap	No. of substitution	
C319-1	<i>Wickerhamomyces anomalus</i> NRRL_Y-366 ^T (EF550341)	LC415305	553/556	2	1	-	-	-	-	<i>W. anomalus</i>
C319-2	<i>Candida metapsilosis</i> CBS10907 ^T (KY106577)	LC415306	552/554	1	1	-	-	-	-	<i>C. metapsilosis</i>
J29-5	<i>Meyerozyma guilliermondii</i> CBS2030 ^T (U45709)	LC497430	532/534	0	2	-	-	-	-	<i>M. guilliermondii</i>
JC29-2	<i>Kodamaea ohmeri</i> CBS5367 ^T (U45702)	LC497431	442/442	0	0	-	-	-	-	<i>K. ohmeri</i>
JC29-6	<i>Papiliotrema laurentii</i> CBS139 ^T (AF075469)	LC497432	573/574	0	1	-	-	-	-	<i>P. laurentii</i>
JC29-7	<i>Meyerozyma caribbica</i> CBS9966 ^T (AY187283)	LC497433	519/519	0	0	-	-	-	-	<i>M. caribbica</i>
JC34-1	<i>Papiliotrema laurentii</i> CBS139 ^T (AF075469)	LC497439	576/577	0	1	-	-	-	-	<i>P. laurentii</i>
JC34-2	<i>Rhodotorula diobovata</i> CBS6085 ^T (AF070421)	LC497440	532/533	0	1	-	-	-	-	<i>R. diobovata</i>
JC34-5	<i>Rhodospiridiobolus fluvialis</i> CBS6568 ^T (AF070422; NR077089)	LC511616	546/549	0	3	LC522505	588/593	0	5	<i>Rh. fluvialis</i>
JC34-7	<i>Rhodotorula mucilaginosa</i> CBS316 ^T (AF070432)	LC497441	533/534	0	1	-	-	-	-	<i>R. mucilaginosa</i>
F9-1	<i>Rhodotorula mucilaginosa</i> CBS316 ^T (AF070432)	LC511623	533/534	0	1	-	-	-	-	<i>R. mucilaginosa</i>
FC9-1	<i>Rhodotorula toruloides</i> CBS349 ^T (AF070426)	LC511624	532/532	0	0	-	-	-	-	<i>R. toruloides</i>