

Table S1. Distribution of yeast species

Participant	No. of spp.	<i>C. albicans</i>	<i>C. glabrata</i>	<i>C. parapsilosis</i>	<i>C. tropicalis</i>	<i>C. pararugosa</i>	Others
TCH-029-27	5	1	0	0	0	1	<i>C. krusei</i> , <i>C. orthopsilosis</i> , <i>Lodderomyces elongisporus</i>
GLS-011-144	5	1	0	1	0	0	<i>C. intermedia</i> , <i>C. lusitaniae</i> , <i>Pichia aff. fermentans</i>
GLS-019-167	4	0	0	1	0	1	<i>Exophiala dermatitidis</i> , <i>Trichosporon mucoides</i>
TCH-011-118	4	1	0	0	1	1	<i>Saccharomyces cerevisiae</i>
FSC-021-90	4	1	0	1	0	0	<i>C. guilliermondii</i> , <i>C. orthopsilosis</i>
HWA-012-21	4	1	0	1	0	1	<i>C. intermedia</i>
HWA-017-63	4	1	1	0	0	0	<i>C. metapsilosis</i> , <i>Magnusiomyces capitatus</i>
FSC-049-93	4	1	1	0	1	0	<i>Saccharomyces cerevisiae</i>
GFC-014-154	4	1	1	0	1	0	<i>C. orthopsilosis</i>
FSC-001-33	4	1	1	1	0	0	<i>Candida guilliermondii</i>
HWA-015-127	3	0	0	0	0	0	<i>C. intermedia</i> ,
GLS-017-136	3	0	0	1	0	1	<i>Trichosporon asahii</i>
YEOo-002-89	3	1	0	0	0	0	<i>C. guilliermondii</i> , <i>Magnusiomyces capitatus</i>
FAM-007-168	3	1	0	1	0	0	<i>C. norvegensis</i>
FSC-041-126	3	1	0	1	0	0	<i>Saccharomyces cerevisiae</i>
FSC-008-46	3	1	0	1	0	0	<i>C. fermentati</i>
FSC-004-42	3	1	0	1	0	0	<i>C. guilliermondii</i>
TCH-013-104	3	1	0	1	0	0	<i>Trichosporon asahii</i>
FAM-001-73	3	1	0	1	0	1	

WES-001-2	3	1	0	1	1	0	
FSC5-039-114	3	1	1	0	0	0	<i>Trichosporon asahii</i>
FSC-029-68	3	1	1	0	0	0	<i>Rhodotorula minuta</i>
GLS5-010-66	3	1	1	0	0	0	<i>Candida ethanolica</i>
FAM-011-34	3	1	1	0	1	0	
FSC5-025-11	3	1	1	0	1	0	
FSC5-016-50	3	1	1	0	1	0	
GLS-008-15	3	1	1	0	1	0	
GLS-007-67	3	1	1	0	1	0	
HWA-002-13	3	1	1	0	1	0	
TCH-010-78	3	1	1	0	1	0	
WES-013-96	3	1	1	0	1	0	
WES-018-40	3	1	1	0	1	0	
WES-020-75	3	1	1	0	1	0	
YEO5-006-108	3	1	1	0	1	0	
TCH-014-117	3	1	1	1	0	0	
HWA-007-56	2	0	1	0	0	0	<i>Candida dubliniensis</i>
TCH5-016-115	2	0	1	0	0	0	<i>Candida metapsilosis</i>
FSC-042-155	2	1	0	0	0	0	<i>Saccharomyces cerevisiae</i>
HOP-008-148	2	1	0	0	0	0	<i>Trichosporon asahii</i>
HWA-014-138	2	1	0	0	0	0	<i>Exophiala dermatitidis</i>
HWA-001-157	2	1	0	0	0	0	<i>Candida guilliermondii</i>
YEO-008-35	2	1	0	0	0	0	<i>Candida metapsilosis</i>
GFC-015-163	2	1	0	0	1	0	
HOP-001-101	2	1	0	0	1	0	
TCH5-043-113	2	1	0	0	1	0	
GLS-014-110	2	1	0	1	0	0	

TCH-021-6	2	1	0	1	0	0	
FSC-026-76	2	1	1	0	0	0	
FSC5-023-38	2	1	1	0	0	0	
FSC-033-137	2	1	1	0	0	0	
FSC-018-57	2	1	1	0	0	0	
GLS-003-102	2	1	1	0	0	0	
GLS-015-24	2	1	1	0	0	0	
GLS-018-16	2	1	1	0	0	0	
GLS-003-128	2	1	1	0	0	0	
HOP-003-7	2	1	1	0	0	0	
HOP-005-111	2	1	1	0	0	0	
HOP-006-29	2	1	1	0	0	0	
HOP-010-121	2	1	1	0	0	0	
HWA-020-39	2	1	1	0	0	0	
HWA-009-53	2	1	1	0	0	0	
HWA-006-109	2	1	1	0	0	0	
TCH-037-150	2	1	1	0	0	0	
TCH5-009-25	2	1	1	0	0	0	
TCH-024-62	2	1	1	0	0	0	
WES-016-92	2	1	1	0	0	0	
WES-019-20	2	1	1	0	0	0	
TCH-012-149	1	0	0	0	0	0	<i>C. utilis</i>
WES-040-44	1	0	0	0	0	0	<i>Candida dubliniensis</i>
YEO-009-141	1	0	0	0	0	0	<i>Candida dubliniensis</i>
YEO-003-52	1	0	0	0	0	0	<i>Candida dubliniensis</i>
HWA-037-123	1	0	0	0	0	1	
GFC-022-98	1	0	0	1	0	0	

GFC-004-94	1	0	0	1	0	0
HOP-007-145	1	0	0	1	0	0
TCH-006-4	1	0	0	1	0	0
FAM-006-146	1	1	0	0	0	0
FAM-004-69	1	1	0	0	0	0
FAM-002-125	1	1	0	0	0	0
FAM-008-70	1	1	0	0	0	0
FSC-031-14	1	1	0	0	0	0
FSC-030-91	1	1	0	0	0	0
FSC-027-51	1	1	0	0	0	0
FSC-019-12	1	1	0	0	0	0
FSC-002-3	1	1	0	0	0	0
FSC-012-41	1	1	0	0	0	0
FSC-032-105	1	1	0	0	0	0
FSC-007-122	1	1	0	0	0	0
FSC-044-82	1	1	0	0	0	0
FSC-014-18	1	1	0	0	0	0
FSC-013-71	1	1	0	0	0	0
GFC5-002-95	1	1	0	0	0	0
GFC-017-119	1	1	0	0	0	0
GLS5-004-134	1	1	0	0	0	0
GLS-006-81	1	1	0	0	0	0
GLS5-012-165	1	1	0	0	0	0
GLS-007-22	1	1	0	0	0	0
GLS-001-97	1	1	0	0	0	0
GLS-002-99	1	1	0	0	0	0
HWA-013-156	1	1	0	0	0	0

Table S2. Distribution of species causing multiple species colonization

Total spp No	No. residents	<i>C. albicans</i>	<i>C. glabrata</i>	<i>C. parapsilosis</i>	<i>C. tropicalis</i>	<i>C. pararugosa</i>	Others
5	1	1	0	1	0	0	<i>C. intermedia</i> , <i>C. lusitaniae</i> , <i>Pichia aff. fermentans</i>
5	1	1	0	0	0	1	<i>C. krusei</i> , <i>C. orthopsilosis</i> , <i>Lodderomyces elongisporus</i>
4	1	1	1	1	0	0	<i>C. guilliermondii</i>
4	1	1	1	0	1	0	<i>Saccharomyces cerevisiae</i>
4	1	1	1	0	1	0	<i>C. orthopsilosis</i>
4	1	1	1	0	0	0	<i>C. metapsilosis</i> , <i>Magnusiomyces capitatus</i>
4	1	1	0	1	0	1	<i>C. intermedia</i>
4	1	1	0	1	0	0	<i>C. guilliermondii</i> , <i>C. orthopsilosis</i>
4	1	1	0	0	1	1	<i>Saccharomyces cerevisiae</i>
4	1	0	0	1	0	1	<i>Exophiala dermatitidis</i> , <i>Trichosporon mucoides</i>
3	11	1	1	0	1	0	
3	1	1	1	1	0	0	
3	1	1	1	0	0	0	<i>Trichosporon asahii</i>
3	1	1	1	0	0	0	<i>Rhodotorula minuta</i>
3	1	1	1	0	0	0	<i>C. ethanolica</i>
3	1	1	0	1	1	0	
3	1	1	0	1	0	1	

3	1	1	0	1	0	0	<i>C. norvegensis</i>
3	1	1	0	1	0	0	<i>Saccharomyces cerevisiae</i>
3	1	1	0	1	0	0	<i>C. fermentati</i>
3	1	1	0	1	0	0	<i>C. guilliermondii</i>
3	1	1	0	1	0	0	<i>Trichosporon asahii</i>
3	1	1	0	0	0	0	<i>C. guilliermondii</i> , <i>Magnusiomyces capitatus</i>
3	1	0	0	1	0	1	<i>Trichosporon asahii</i>
3	1	0	0	0	0	0	<i>C. intermedia</i> ,
2	20	1	1	0	0	0	
2	3	1	0	0	1	0	
2	2	1	0	1	0	0	
2	1	1	0	0	0	0	<i>Saccharomyces cerevisiae</i>
2	1	1	0	0	0	0	<i>Trichosporon asahii</i>
2	1	1	0	0	0	0	<i>Exophiala dermatitidis</i>
2	1	1	0	0	0	0	<i>C. guilliermondii</i>
2	1	1	0	0	0	0	<i>C. metapsilosis</i>
2	1	0	1	0	0	0	<i>C. dubliniensis</i>
2	1	0	1	0	0	0	<i>C. metapsilosis</i>

No, number.