Table S1. Morphological characteristics of the yeast strains isolated on WL agar and their designation, sequence similarity with species type strain and GenBank (*29*) accesion numbers of ITS and D1/D2 sequences

Species	No.	Yeast strain	Colony colour	Colony morphology	*N*(compared nucleotide)	Sequence similarity/%	GenBank acc. no.
S. cerevisiae	1.	R2161	Cream to green	Grown in centre, smooth, opaque, butyrous		100	
	2.	K2161				100	
	3.	Mer382				100	
	4.	K3161			1312	99.3	MT734685
	5.	K3162				100	
	6.	R3165				100	
	7.	CH1213				100	
	8.	R3161				100	
	9.	R3163				100	
	10.	MAL382			1310	100	MT734684
	11.	ZW2271				100	
	12.	Mer384				100	
	13.	Mer182				100	
	14.	K1161			1313	99.8	MT734686
	15.	R1161				100	
	16.	CH1212				100	
	17.	Mer281				100	
	18.	R1163				100	
	19.	MAL282				100	
	20.	CH1211				100	
	21.	K3163			1312	99.8	MT734687
	22.	Mer282				100	
	23.	MAL381				100	
	24.	R2162				100	
	25.	ZW2273				100	
	26.	Mer283				100	
	27.	ZW2272				100	
	28.	R1162				100	
	29.	MAL384				100	
	30.	K2163				100	
	31.	R3164				100	
	32.	Mer181				100	
	33.	K1162				100	
	34.	MAL281				100	
	35.	MF1161				100	
	36.	ZW3271				100	
	37.	CH1214				100	
	38.	MAL283				100	
	39.	MAL383				100	
	40.	ZW3272				100	
	41.	MF1162				100	
	42.	Mer183				100	
	43.	R2163				100	
	44.	ZW3273				100	
	45.	ZW3274				100	
	46.	K2162				100	
Hanseniaspora uvarum	47.	Mer381	Intense green	Flat surface, smooth, opaque, butyrous		100	
	48.	Š3272				100	
	49.	Mer383			1251	100	MT734682
	50.	Š3271				100	
	51.	Š3273				100	
Pichia kudriavzevii	52.	Š2271	White to cream	Low convex with a flattened centre, lobed, fringed margin, butyrous	998	100	MT734683
	53.	Š2273				100	
	54.	Š2272				100	
Saturnispora diversa	55.	MF1163	White to cream	Smooth, butyrous, with entire margin	550	100	MT730589*
	56.	M1282				100	
Starmerella bacillaris	57.	M2281	Light to intense green	Low colonies with smooth to finely lobed margins, butyrous		100	
	58.	MF3163				100	
	59.	K2164				100	
	60.	MF3162			890	100	MT734681
	61.	M1281				100	
	62.	MF3161				100	
	63.	M2282				100	
	64.	MF3164				100	

*only D1/D2 region