

**Table S1.** Identification, based on blast comparison in GenBank of ribosomal region 26S rRNA of 10 yeast strains selected on the basis of RAPD-PCR cluster analysis.

Cluster	Strain	Size	Closest Relative	% Identity	Sequence ID *
A	YFi12	573	<i>Saccharomyces cerevisiae</i> isolate 20170721	100%	MF521980
B	YG7	573	<i>Torulaspora delbrueckii</i> YMTT5-1	100%	LC387296
C	YC15	590	<i>Pichia kudriavzevii</i> strain LY49	100%	KY705008
D	YG8	576	<i>Hanseniaspora uvarum</i> isolate 20170728	98%	MF521987
E	YA17	578	<i>Hanseniaspora uvarum</i> isolate SFM28	100%	MG017569
F	YC18	588	<i>Hanseniaspora uvarum</i> isolate 11-1288 la	100%	MH468682
G	YA27	610	<i>Saccharomyces cerevisiae</i> strain LY183	100%	KY705009
H	YFa31	833	<i>Metschnikowia pulcherrima</i> culture CBS:2243	98%	KY108490
I	YA28	596	<i>Hanseniaspora uvarum</i> isolate 11-1288 la	100%	MH468682
L	YS20	921	<i>Zygotorulaspora florentina</i> CBS 746	99%	NG_058449

\* Accession number of the sequence of the closest blast search.