

## Frisinga - TUM 34/70<sup>®</sup> & Proles - TUM 34/78<sup>®</sup>

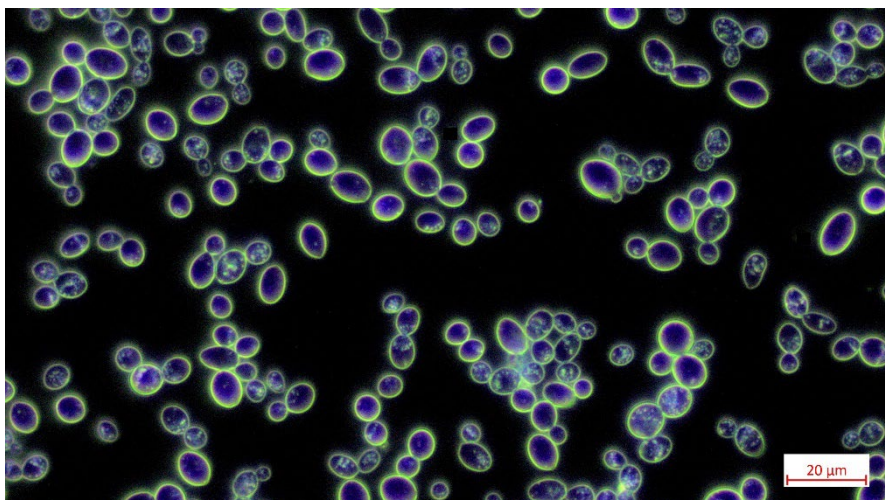
*Saccharomyces pastorianus ssp. carlsbergensis*  
bottom fermenting brewing yeast

### Short description

These yeast strains are excellently suited for the production of bottom-fermented beers of all types. The resulting beer possesses an extremely pure flavor, a fine, subtle aroma and a mild overall impression. Fermentation is rapid with optimal yeast flocculation. The degree to which the color lightens during fermentation is sufficient.

The principle difference between Frisinga - TUM 34/70<sup>®</sup> and Proles - TUM 34/78<sup>®</sup> is their flocculation behavior during primary fermentation. Proles - TUM 34/78<sup>®</sup> can remain somewhat longer in suspension. The length of time that the yeast remains in suspension is dependent on conditions present in the brewery.

<b>Examination parameters</b>	<b>Result</b>
Original extract (%)	11.6
Apparent degree of attenuation (%)	73
Cells in suspension (Mio/ml)	12.5
pH value	4.6
Diacetyl (mg/l)	0.7
Acetaldehyde (mg/l)	6
Higher aliphatic alcohols (mg/l)	58
Esters (mg/l)	18.5
Foam according to Ross & Clark (sec)	132
<b>Short characteristics</b>	
Fermentation rate	high
pH drop	normal
Flocculation	optimal
Diacetyl reduction	very good
Foam	very good
Acetaldehyde	normal
Higher alcohols	very low
Esters	pronounced



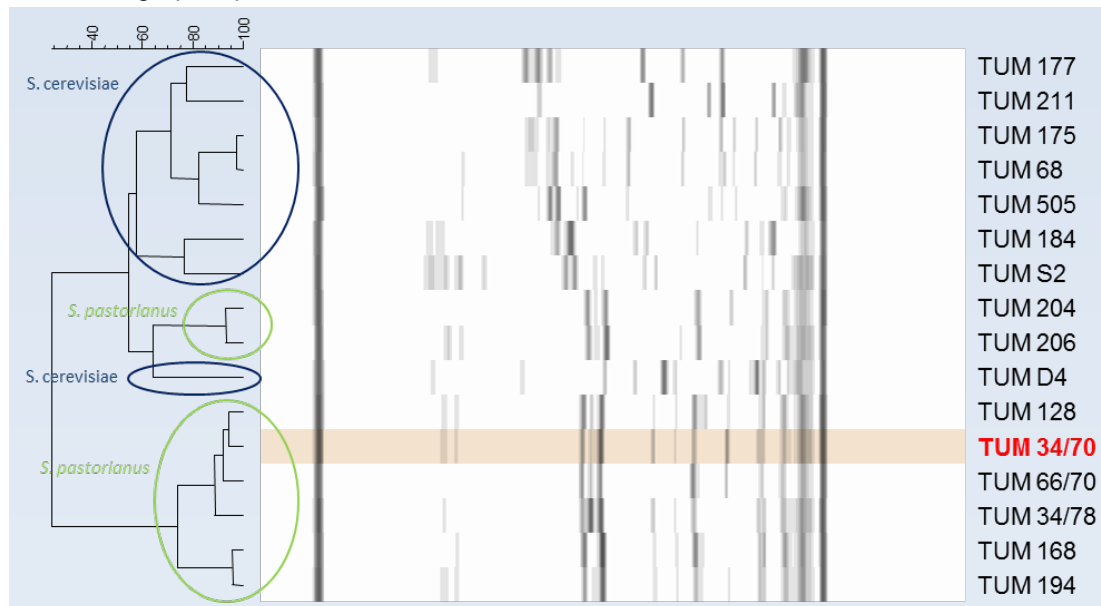
Microscopic view of yeast strain Frisinga – TUM 34/70<sup>®</sup>  
(Picture Frisinga – TUM 34/70<sup>®</sup> © FZW BLQ)

Real-time PCR Screening Profile:

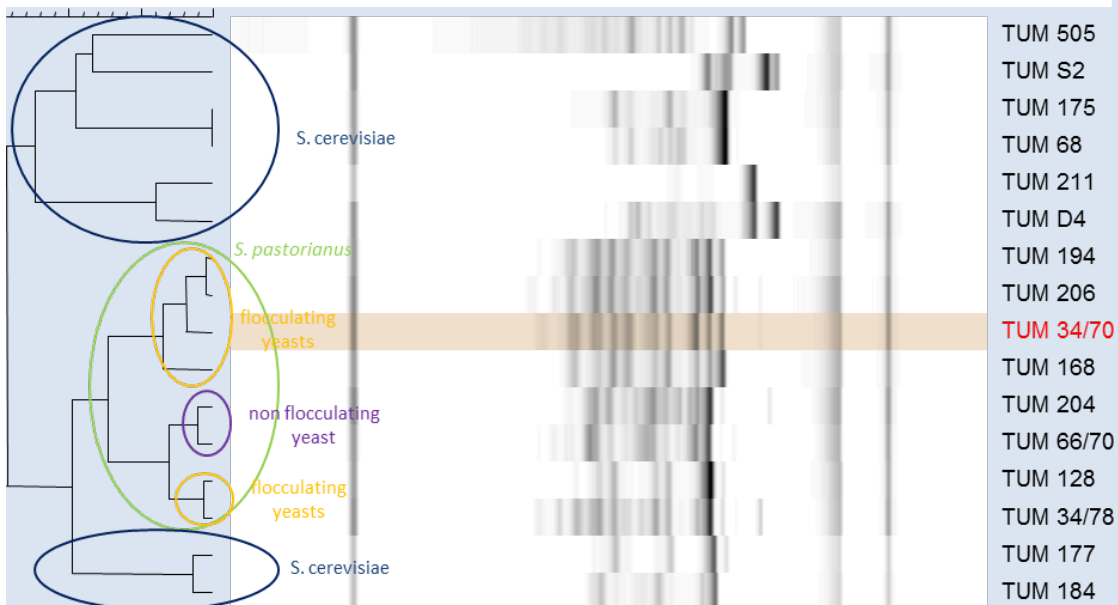
Comparison of individual qualitative results from real-time PCR systems for the differentiation of industrial *Saccharomyces* strains (with a focus on *S. cerevisiae* and *S. pastorianus* (bottom-fermenting strains)) (according to Hutzler, M. 2009, Hutzler, M. 2010)

Yeast	Strain	PCR System					
		Sc-GRC3	Sc	OG-COXII	Sbp	UG-LRE1	UG-300
<i>S. bayanus</i>	DSM 70412T, 70547, BTII K 1-C-3	-	-	-	+	-	-
	70411, 70508	-	-	-	+	+	+
<i>S. bayanus/pastorianus</i>	CBS 2440, 6017	-	-	-	+	+	+
<i>S. pastorianus</i>	CBS 1503, 1513, 1538, DSM 6580NT, 6581	-	-	-	+	+	+
<i>S. pastorianus</i> (bottom-fermenting)	TUM 26, 44, 34/70, 34/78, 44, 54, 59, 69, 84, 105, 109, 120, 128, 168, 172, 180, 194, 199, 206 (flocculent yeasts)	+	+	-	+	+	+
	TUM 71, 144 (low or non-flocculent yeasts)						
	CBS 1484, 5832, CBS 6903, NBRC 2003, BTII K B-I-4, B-J-4, B-J-5						
	TUM 120 (flocculent yeast) TUM 66, 66/70, 204 (low or non-flocculent yeasts) CBS 5832, CBS 6903	+	+	-	+/-	+	+
<i>S. cerevisiae</i>	DSM 70424, 70449T, 70451, CBS 1464, 8803, BT II K 3-A-1, 3-C-3, 3-G-1, 5-A-7, 6-I-1, 6-F-4	+	+	+	-	-	-
<i>S. cerevisiae</i> (top-fermenting)	TUM 68, 127, 149, 175, 205, BTII K 5-A-8 (Southern German style wheat beer yeast)	+	+	+	-	-	-
	TUM 148, 184, 208 (alt yeast)	+	+	+	-	-	-
	TUM 165, 177 (kölsch yeast)	+	+	+	-	-	-
	TUM 210, 211, 213 (ale yeast)	+	+	+	-	-	-
	TUM Bingen, Bordeaux, Eperney, Laureiro, Stein, Wädensvill (wine yeast)	+	+	+	-	-	-
	TUM B4 (distillers' yeast)	+	+	+	-	-	-
	TUM S2 (sparkling wine yeast)	+	+	+	-	-	-
<i>S. cerevisiae var. diastaticus</i>	CBS 1782, DSM 70487, BTII K 1-B-8, 1-H-7, 2-A-7, K 2-F-1, 3-D-2, 3-H-2, 3-H-4	+	+	+	-	-	-
<i>S. cariocanus</i>	CBS 7995, 8841	-	+	-	-	-	-
	CBS 5313	+	+	+	-	-	-
<i>S. kudriavzevii</i>	CBS 8840	-	-	-	-	-	-
<i>S. mikatae</i>	CBS 8839	-	-	-	-	-	-
<i>S. paradoxus</i>	CBS 406, 432, 2908, 5829, 7400, 8436	-	+	-	-	-	-

Genetic fingerprint profile:



The virtual electrophoresis gel depicted above was created using capillary electrophoresis separated by interdelta PCR ( $\theta_{12}/\theta_{21}$ ). Pictured are results for strain TUM 34/70 compared with a selection of yeast strains most frequently supplied by the Weihenstephan Research Center (FZW BLQ).



A virtual image of the rDNA IGS2\_314 genetic fingerprint of yeast strains separated using capillary gel electrophoresis is depicted above. Pictured are the results for strain Frisinga - TUM 34/70<sup>®</sup> compared with a selection of yeast strains most frequently supplied by the Weihenstephan Research Center (FZW BLQ)

- References:
- |                                |  |
|--------------------------------|--|
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| Technische Universität München | · Research Center Weihenstephan for Brewing and Food Quality   |