SCREENING OF KILLER-SENSITIVE PATTERN (KSP) FOR BIOTYPING YEAST STRAINS ISOLATED FROM DAIRY PRODUCTS

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Abstract

Killer-Sensitive Pattern (KSP) was screened by cross reactions in 50 yeast species belonging to 20 genera which were previously isolated from different dairy products. The Killer-Sensitive Pattern (KSP) appeared as strain character rather than species level. Among all yeasts, strain designated as YF19-*Lipomyces starkeyi* appeared as the most killer *i.e.* showed 46.93% killing activity and strains appeared as most sensitive were YF45-*Bullera pyricola* (77.55%), YF42-*Pichia heimii* (77.50%), YF87-*Bullera pseudoalba* (51.02%) and Y90-*Williopsis californica* (42.86%).

Introduction

Certain yeasts produce killer toxins (mycocins), which are lethal to closely related strains but the killer yeast itself has a killer resistant phenotype (Bevan & Makower, 1963; Woods & Bevan, 1968; Bussey, 1972; Pfeiffer & Radler, 1982; Spencer & Spencer, 1997). The killer phenomenon provides an excellent model system to study host-virus interactions in eukaryotic cells (Wickner, 1979, 1989) and to investigate the mechanisms of protein processing and secretion (Douglas et al., 1988). Possible uses of killer phenomenon, which aroused great interest, include the differentiation of pathogenic strains (Morace et al., 1984) and their possible role in ecosystems mainly in natural fermentation processes (Starmer et al., 1987; Vagnoli et al., 1993; Hidalgo & Flores, 1994). Killer activity is one of the mechanisms of antagonism among yeasts during spontaneous fermentations and because of this mechanism killer strains could dominate at the end of the wine fermentation (Bussey et al., 1988; Jacobs et al., 1988; Longo et al., 1990).

Killer toxins are protein in nature and active at low pH (Young & Yagiu, 1978; Pfeiffer & Radler, 1982; Radler et al., 1985). They are secreted in an inactive glycosylated form that once secreted to the cell plasma membrane, becomes cleaved (Zhu et al., 1993). A portion of the toxin with the glycosylated site remains associated with the membrane and conveys immunity to the cell. The cleaved mature toxin is available to bind at the sites located on the cell wall and the plasma membrane of sensitive yeasts. However the phenomenon of insensitivity towards killer toxins generally occurs at the cell wall level. Resistant yeasts lack receptors necessary for the formation of the link and thus for the action of the killer toxin (Marquina et al., 2002; Golubev, 2006). As a result, if different cell wall chemical compositions are taxon-associated; resistance, causing insensitivity could be a taxon-related property as well (Golubev, 1998, 2006). Based on evidence that the chemical composition of yeast cell walls is a taxon related characteristic, Golubev (2006) hypothesized that KSP profiles may have taxonomic relevance. The theoretical rationale supporting this conclusion is related to the resistance mechanism. In a previous study we screened Killer-Sensitive-Pattern (KSP) among yeast species previously isolated from slime fluxes of different trees and flowers' nectar (Mushtaq et al., 2010). In the present study Killer-Sensitive-Pattern (KSP) has been screened by cross reactions in yeast species isolated from

different dairy products (Mushtaq et al., 2007; Mushtaq et al., 2006).

Materials and Methods

A modified method of Abranches et al., (1997) was used to screen Killer-Sensitive Pattern (killer, sensitive and neutral phenotypes) in 23 yeasts species belonging to 13 genera previously isolated from slime fluxes of trees and 57 yeast species belonging to 23 genera from flowers' nectar, on yeast extract-malt extract agar supplemented with 0.003% methylene blue (YM-MB Agar). Twentyfour h old yeast culture grown on YM agar (Kreger-van Rij, 1984) was diluted in double distilled sterile water to obtain a suspension of $4x10^5$ cells/ml and spread with a sterile cotton swab as seeded (lawn) cultures on the surface of YM-MB agar in Petri plates and dried. Fresh cultures of the yeasts to be tested were grown on YM agar for 24 h and each inoculated in a single streak on plates seeded with the yeast culture and incubated at 25±1°C for 10 days and observed daily. The seeded yeast was considered as killer if a blue colored killing zone appeared on streak and sensitive if killing zone appeared around the streak on lawn. Intensity of the killer activity was recorded as K^{+1} (very light blue killing zone), K^{+2} (blue killing zone), K^{+3} (dark blue killing zone) and K^{+4} (intense dark blue killing zone) reaction. The sensitivity of the yeast was also recorded in the same manner as S^{+1} . S^{+2} , S^{+3} and S^{+4} . A negative reaction indicated by (-) when yeasts did not show any reaction. Percentages of killing activity and sensitivity of yeast species were calculated. Strains that showed >40% killing activity or sensitivity were considered as super killers and super sensitive.

Results

Killer-Sensitive Pattern (KSP) (killer, sensitive & neutral phenotypes) was screened by cross reactions in 50 yeast species belonging to 20 genera previously isolated from different dairy products. Spectrum of killing activity and sensitivity (species vise) is presented respectively in table 1 and their percentages in table 2. One of the strain of *Lipomyces starkeyi* designated as YF19 showed 46.93% killing activity and appeared as the most killer strain, on the other hand, yeast strains designated as YF45-*Bullera pyricola* (77.55%), YF42-*Pichia heimii* (77.50%), YF87-*B. pseudoalba* (51.02%) and Y90-*Williopsis californica* (42.86%) appeared as the most sensitive yeast strains.

		Streak strains																
		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17
No.	Seeded strains	Arxula adeninovorans	Bensingtonia intermedia	B. naganoensis	Bullera pseudoalba	B. pyricola	C. diddensiae	C. etchellsii	C. friedrichii	C. haemutonii	C. membranifaciens	C. pseudointermedia	C. shehatae	C. succiphila	C. valdiviana	C. xestobii	Clavispora lusitaniae	Cryptococcus albidus
1.	Arxula adeninovorans	x	-	S ⁺¹	-	K ⁺¹	-	-	-	S ⁺¹			-	-	-	-	-	
2.	Bensingtonia intermedia	-	x		K*2	K ⁺²	S*3	-	-	-				-		-		
3.	B. naganoensis	K^{+1}	-	х	-	K^{+1}	-	-	-	-	-	S^{+2}		-	-	-	-	
4.	Bullera pseudoalba		S^{+2}		х	K+1S+2	S^{+2}	S^{+1}	-	S^{+2}	S^{+1}		S ⁺³		K^{+2}		S^{+4}	S^{+1}
5.	B. pyricola	S ⁺¹	S^{+2}	S ⁺¹	K+2S+1	х	S^{+1}	-	S^{+1}	-	$K^{+2}S^{+1}$	-	S ⁺¹	S ⁺²	S ⁺¹	K^{+2}	S ⁺¹	$K^{+2}S^{+1}$
6.	Candida diddensiae	-	K^{+3}		K^{+2}	K^{+2}	х	-	-	-	-			-	-	K^{+2}	K^{+1}	K^{+2}
7.	C. etchellsii	K^{+1}	-	-	K^{+1}	-	-	х	K^{+1}	-	-	-	-	-	K^{+1}	-	-	-
8.	C. friedrichii	-	K^{+3}	-	K^{+1}	K^{+2}	-	-	х	-	K^{+1}	-	-	-	K^{+2}	-	-	-
9.	C. haemulonii	-	-	-	K^{+2}	-	-	-	-	х	-	-	-	-	-	-	-	K^{+1}
10.	C. membranifaciens	-	-	-	K^{+1}	$\mathrm{K}^{+1}\mathrm{S}^{+2}$	-	-	-	-	х	-	S^{+3}	-	-	-	S^{+1}	-
11.	C. pseudointermedia	-	-	K^{+2}	-	-	-	-	-	-	-	х	-	-	-	-	-	-
12.	C. shehatae	-	-	-	K^{+3}	K^{+2}	-	-	-	-	K^{+3}	-	х	-	K^{+3}	S^{+1}	K^{+3}	K^{+3}
13.	C. succiphila	-	-	-	-	K^{+2}	-	-	-	-	-	-	-	х	-	-	-	-
14.	C. valdiviana	-	-	-	S ⁺²	K^{+1}	-	S^{+1}	-	-	-	-	S^{+3}	-	х	K^{+1}	-	-
15.	C. xestobii	-	-	-	-	S*2	K^{+2}	-	-	-	-	-	S ⁺¹	-	S*1	х	-	-
16.	Clavispora lusitaniae	-	-	-	K^{*2}	K^{+1}	S^{+2}	-	-	-	K^{+1}	-	S^{+3}	-	-	-	х	K^{*1}
17.	Cryptococcus albidus	•	•	•	K^{+1}	$K^{+1}S^{+2}$	S^{+2}	-	-	S^{+1}	-	•	S^{+3}	•	-	-	S^{+1}	х
18.	C. gastricus	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
19.	Debaryomyces castellii	-	S^{+1}	•	K*2	K^{+1}	-	-	-	-	•	-	•	-	-	K^{+1}	•	•
20.	D. hansenii	S ⁺¹	K^{+1}	-	S ⁺¹	K^{+1}	-	-	-	-	-	S ⁺¹	-	-	-	S ⁺¹	-	-
21.	D. nepalensis	-	-	-	-	K^{+1}	-	-	-	-	-	-	-	-	-	-	-	-
22.	D. vanrijii	-	-	K^{+2}	-	-	-	-	-	-	-	-	-	-	-	-	-	-
23.	D. yamadae	-	-	-	-	K^{+2}	-	-	-	-	-	-	-	-	-	-	-	-
24.	Fibulobasidium inconspicuum	-	-	K*2S*2	K*1	-	S^{*1}	-	-	-	-	S*1	-	-	K ⁺¹	-	-	-
25.	Filobasidiella neoformans	-	-	-	K*2	-	-	-	-	-	-	-	-	-	-	-	-	-
26.	Filobasidium uniguttulatum	-	-	-	K ⁺¹	K+2	-	-	-	-	-	-	-	-	K ⁺¹	-	-	-
27.	Kluyveromyces polysporus	S**	-	S	-	K**	S-1	-	-	-	-	S ⁻¹	S	-	-	-	-	-
28.	Lipomyces lipofer	S	- 17 ⁺¹	-	-	K **	-	-	-	-	K **	-	-	-	K 1	-	K **	K** 17*2
29.	L. starkeyi	-	K.	K	K	K ·	К -	-	K -	K -	K -	-	K -	-	K 1	-	K -	K -
30.	Piahia avousta	•	•	•	•	K	•	-	K	•	•	•	•	•	ĸ	•	•	•
32	P anomala	s+1	s+1			- K ⁺²	- S ⁺³	- K ⁺¹	-	г. К ⁺¹	-	s+1	s+3	-	-	г. К ⁺¹	-	
32.	P. anomaia P. amharhiinhila				K ⁺¹	K ⁺¹		ĸ		ĸ			3			к -		
34	P. guilliermondii				K ⁺²	K ⁺¹		-	-									
35.	P. heimii	S ⁺²	S ⁺⁴	S ⁺³	K ⁺⁴ S ⁺²	K ⁺³ S ⁺²	S^{+3}	-	-	S^{+1}	S ⁺³	S ⁺²	K ⁺² S ⁺²	-	K+3S+2	K*3	K ⁺² S ⁺²	K+3S+2
36.	P. iadinii		-			K ⁺¹	-	-	-	-	-	-	S ⁺¹	-	-	-	-	
37.	P. lvnferdii	K^{+2}	-	K^{+2}	K^{+2}	K ⁺³	-	-	-	-	K ⁺²	K*2		-	-	-	-	K^{*2}
38.	P. methanolica	-	-	-	K*2S*2	K^{+2}	-	-	-	-	-			-	-	-	-	-
39.	P. mexicana	S ⁺¹	S^{+2}			K^{+1}	-	-	-	-	-	-	S ⁺³	-	-	-	-	-
40.	P. ofunaensis	-	-		K+3	-	-	-	-	-	$K^{+1}S^{+1}$	S^{+2}	-	-	-	-	-	$K^{+2}S^{+1}$
41.	P. ohmeri	-	-			K ⁺²	-	-	-	-	-			-	-	-	-	
42.	P. strasburgensis	-	-	-	K^{+2}	-	S^{+1}	-	-	-	-		-	-	-	-	-	-
43.	P. sydowiorum	-	-	K^{+1}	-	-	-	-	-	-	-	S*3		-	-	-	-	
44.	Saccharomycodes ludwigii	-	-	-	K^{*2}	K^{+1}	-	K^{+3}	-	-	-	-	-	S^{+1}	-	-	-	-
45.	Sporidiobolus ruineniae			K^{+2}	K^{+2}	-	-		-	-	-	S^{+2}	-		-	-	K^{+2}	-
46.	S. salmonicolor	-	-	K^{+2}	K^{+4}	K^{+2}	-	-	K^{+3}	S^{+1}	-	S ⁺¹	S ⁺¹	S ⁺¹	K^{+1}	-	-	S ⁺²
47.	Sporobolomyces tsugae	K^{+1}	S^{+3}	K^{+2}	K^{+2}	K^{+2}	-	S^{+2}	-	S^{+2}	-	$\mathrm{K}^{+1}\mathrm{S}^{+1}$	-		K^{+2}	K^{+3}	-	-
48.	Stephanoascus ciferrii	-	-	-	-	K^{+2}	-	-	-	-	-	-	-	-	-	-	-	-
49.	Tremella encephala	-	-	-	-	K^{+1}	S^{+1}	-	-	-	-	-	K^{+1}	-	-	-	S^{+1}	-
50.	Williopsis californica	-	S*2	K ⁺¹ S ⁺²	S ⁺¹	K ⁺¹	S*2	-	S*3	-	-	S*2	-	-	K ⁺¹	K^{+2}	-	-

Table 1. Cross reaction screening of killer, sensitive and neutral phenotypes in yeast species isolated from dairy products.

								<u> </u>	S	treak str	ains							
		18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34
No.	Seeded strains	C. gastricus	Debaryonyces castellii	D. hansenii	D. nepalensis	D. vanrijii	D. yamadae	Fibulobasidium inconspicuum	Filobasidiella neoformans	Filobasidium uniguttulatum	Kluyveromyces polysporus	Lipomyces lipofer	L. starkeyî	Phaffia rhodozyma	Pichia angusta	P. anomala	P. euphorbiiphila	P. guilliermondii
1.	Arxula adeninovorans		-	K ⁺¹			-			· .	K ⁺¹		-	- <u>-</u>				· -
2	Rensingtonia intermedia		K ⁺¹	S ⁺¹				-		-		-	S ⁺¹	-		K ⁺¹		
3	R. naganoensis										K ⁺¹	-						
4	Bullera pseudoalba	-	S^{+2}	K ⁺¹	-	-	-	S ⁺¹	S^{+2}	S^{+1}		-	S ⁺³				S^{+1}	S^{+2}
5.	B. pyricola	-	S ⁺¹	S ⁺¹	S ⁺¹	S*2	K^{+2}	K ⁺² S ⁺²		S*2	S ⁺¹	S^{+1}	S ⁺³	S ⁺²		S ⁺¹	S ⁺¹	S*1
6	Candida diddensiae			-				K ⁺¹		-	K ⁺¹		S ⁺³	-		K ⁺⁴		
7	C etchellsii												·.			S ⁺¹		
8	C. friedrichii	-		K ⁺³	-	K ⁺²	-			-		-	K ⁺¹	S ⁺¹			-	
9	C haemulanii											_	S*2			K ⁺¹		
10	C. membranifaciens											K ⁺³	S ⁺²					
11.	C. pseudointermedia	-	-	K ⁺¹		-	-	K ⁺¹		-	K ⁺¹					K ⁺¹		
12	C. shehatae						_				K ⁺³		S ⁺¹		_	K ⁺⁴		
13	C. succiphila																	
14.	C. valdiviana		-		-	-	-	S ⁺¹		S ⁺¹	-	-	S ⁺¹	S ⁺¹				-
15.	C. xestobii	-	S ⁺¹	K ⁺¹	-	-	-		-	-		-	-			S ⁺¹		
16.	Clavispora lusitaniae											S ⁺²						
17.	Cryptococcus albidus	-	-		-	-	-	-	-	-	-	S^{+1}	S^{+2}		-		-	
18.	C. gastricus	х	-		-	-	-		-	-	-	-	S ⁺¹			K+4	-	
19.	Debarvomvces castellii	-	х		-	-	-		S*1	-		-	S*2				-	
20.	D. hansenii	-	-	х		-	-	K^{+3}	-	-	-	-	S^{+2}				S^{+1}	
21.	D. nepalensis	-	-	-	х	-	-	-	-	-	-	-	-	-	-	-	-	-
22.	D. vanrijii		-		-	x	-	-	-		-	-	-		-		-	-
23.	D. yamadae	-	-		-	-	х		-	-		-	-		-		-	
24.	Fibulobasidium inconspicuum	-	-	S+3	-	-	-	х	-	-	-	-	-		-	K^{+1}	-	-
25.	Filobasidiella neoformans	-	K^{+1}	-	-	-	-	-	x	-	-	-	-		-	-	-	-
26.	Filobasidium uniguttulatum		-		-		-	-	-	х	-	S^{+3}	-		-		-	
27.	Kluyveromyces polysporus	-	-		-	-	-	-	-		х	S^{+2}	-		-	-	-	-
28.	Lipomyces lipofer	-	-		-	-	-	-	-	K^{+3}	K^{+2}	х	S ⁺¹	-	-	K^{+4}	-	-
29.	L. starkeyi	K^{+2}	K^{+2}	K^{+2}		-	-	-	-			K^{+2}	х		-	K^{+2}	-	
30.	Phaffia rhodozyma	-	-	-	-	-	-	-	-	-	-	-	-	Х	-	-	-	-
31.	Pichia angusta	-	-	-	-	-	-	-	-	-	-	-	-	-	х	-	-	-
32.	P. anomala	S^{+3}	-			-	-	S^{+1}	-		-	S^{+3}	S^{+1}	S^{+1}	-	х	S^{+1}	
33.	P. euphorbiiphila	-	-	K^{+1}	-	-	-	-	-	-	-	-	-	-	-	K^{+1}	х	-
34.	P. guilliermondii	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	х
35.	P. heimii	S*1	S ⁺²	S*3	S*2	-	K^{+2}	$K^{+1}S^{+3}$	S*4	S*3	K+2S+3	S*3	-	S*3	-	K*3S*2	S*3	S*3
36.	P. jadinii	-	-	-	-	-	-	-	-	-	K^{+1}	S^{+1}	-	-	-	-	-	-
37.	P. lynferdii	-	-	-	-	-	-	S^{+1}	-	S^{+1}	S^{+1}	-	-	-	-	-	K^{+2}	-
38.	P. methanolica	-	K^{+1}	•	-	-	-	-	-	-	K^{+2}	-	-	•	-	K^{+1}	-	-
39.	P. mexicana	-	-	-	-	-	-	S ⁺¹	S^{+1}	-	S ⁺¹	-	S ⁺¹	K+2S+1	-	-	-	-
40.	P. ofunaensis	-	-	-	-	-	-	S ⁺¹	-	$K^{+2}S^{+1}$	S ⁺³	-	S ⁺³	-	-	-	-	-
41.	P. ohmeri	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
42.	P. strasburgensis	-	K^{+1}	S^{+1}	-	-	-	K*2	-	-	-	-	-	-	-	K^{+1}	-	-
43.	P. sydowiorum	-	-	K ⁺¹ S ⁺¹	-	-	-	K^{+1}	-	-	K^{+1}	S ⁺¹	-	-	K^{+1}	-	-	-
44.	Saccharomycodes ludwigii	-	-	-	-	-	-	-	-	-	-	-	S ⁻¹	-	-	-	-	-
45.	Sporidiobolus ruineniae	-	-	-	-	-	-	S-1	-	K**S*1	S*3	S*2	S'2	-	•	-	•	-
46.	S. salmonicolor	-	-	K**	-	-	-	K**	-	K**	K**	ST	S**	-	-	-	-	-
47.	Sporobolomyces Isugae	8.	8	-	-	-	-	K	8	K	K	-	8	8	-	-	-	K 'S'
48.	Stephanoascus čiferrii	-	-	-	-	-	-	-	-	-	-	- c +1	e ⁺²	-	-	-	-	-
49.	Tremena encephala Willionsis californica	-	s+2	s+3	s+1	-	-	- K ⁺¹ S ⁺²	s+2	-	-	5	5	-	-	-	-	-
50.	manopsis canjornica	-			3	-	-	N 2	3									

Table 1. (Cont'd.).

		35	36	37	38	39	40	41	Streak 42	strains 43	44	45	46	47	48	49	50
		. heimii	, jadinii	lynferdii	ethanolica	mexicana	ofunaensis	. ohmeri	asburgensis	ydowiorum	mycodes Indwigii	bolus ruineniae	dmonicolor	lomyces tsugae	oascus ciferrii	lla encephala	sis californica
No.	Seeded strains	1		Ρ.	P. 11	Ρ.	P. (1	P. st	P. 5	Saccharo.	Sporidio	S. 50	Sporoba	Stephan	Treme	Williop
1.	Arxula adeninovorans	K*2		S ⁺¹	-		S ⁺¹				-			S*1		-	K*2
2.	Bensingtonia intermedia	K ⁺⁴	-	-	-	K ⁺²		-			-	-	-	K+3		-	K ⁺²
3.	B. naganoensis	K*3	-	S^{+1}	-		-	-	-	S ⁺¹	-	S^{+2}	S^{+2}	S*2	-	-	K ⁺² S ⁺¹
4.	Bullera pseudoalba	K+2S+4	-	S^{+2}	K ⁺² S ⁺²	-	S^{+3}	-	S*2		S*2	S^{+2}	S^{+4}	S*2		-	K^{+1}
5.	B. pyricola	S*3	S^{+1}	S*3	S^{+2}	S ⁺¹	-	S^{+1}			S*1	S^{+1}	S ⁺¹	K*2S*2	S*2	S ⁺¹	S ⁺¹
6.	Candida diddensiae	K*3	-	-	-	-	-	-	K^{+1}	-	-	-	-	-	-	K^{+1}	$\mathbf{K}^{+2}\mathbf{S}^{+1}$
7.	C. etchellsii	-	-	-	-	-	-	-	-	-	S^{+2}	-	-	K^{+3}	-	-	-
8.	C. friedrichii	-	-	-	-	-	-	S^{+2}	-		-	-	$\mathrm{K}^{+1}\mathrm{S}^{+1}$	-	K^{+2}	-	S^{+1}
9.	C. haemulonii	K^{*2}	-	-	-			-			-	-	K^{+1}	K^{+3}		-	-
10.	C. membranifaciens	K*3	-	S^{+2}	-	-	$\mathrm{K}^{+1}\mathrm{S}^{+1}$	-	-	-	-	-	-	K^{+1}	-	-	-
11.	C. pseudointermedia	K^{+2}	-	S^{+2}	-	-	K^{+2}	-	-	K^{+3}	-	S^{+1}	K^{+1}	$K^{+1}S^{+1}$	-	-	K*2
12.	C. shehatae	$\mathrm{K}^{+2}\mathrm{S}^{+2}$	K^{+1}	-	-	K^{*3}	-	-	-	-	K^{+1}	-	K^{+1}	-	-	S^{+1}	-
13.	C. succiphila	-	-	-	-	-	-	-	-	-	K^{+1}	-	K ⁺²	-	-	-	-
14.	C. valdiviana	K*3S*3	-	-	-	-	-	-	-	•	-	-	S^{+1}	S*2	-	-	S*2
15.	C. xestobii	S ⁺³	-	-	-	-	-	-	-	-	-	-	-	S ⁺³	-	-	S ⁺²
16.	Clavispora lusitaniae	K ⁺² S ⁺²	-	-	-	-	-	-	-	-	K^{+1}	S ⁺²	-	-	-	K^{+1}	-
17.	Cryptococcus albidus	K*2S*3	-	S^{+2}	-	-	$K^{+1}S^{+2}$	-	-	-	-	-	K^{+2}	-	-	-	-
18.	C. gastricus	K*2	-	-	-	-	-	-	-	-	-	-	-	K*3	-	-	-
19.	Debaryomyces castellii	K*2	-	-	S*1	-	-	-	S*1	-	-	-	-	K*3	-	-	K*2
20.	D. hansenii	K*3	-	-	-	-	-	-	K^{*1}	$K^{+1}S^{+1}$	-	-	S^{+2}	-	-	-	K*3
21.	D. nepalensis	K^{+2}	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
22.	D. vanrijii	-	-	-	-	•	-	-	•	•	-	-	•	•	•	-	-
23.	D. yamadae	S ⁻¹	-	-	-	-	-	-	-	-	-	-	-	- ***1e#1	-	-	-
24.	Fibulobasidium inconspicuum	K**S**	-	K	-	K	K	-	S	S	-	-	S	K 'S''	-	-	K**S**
25.	Filobasidiella neoformans	K **	-	-	-	-	- rr+1e+2	-	-	-	-	-	- e+1	K**	-	-	K** V*2
26.	Filobasidium uniguttulatum	K -	- c+1	K -	-	K *	K 'S -	-	-	- c+1	-	- c+2	S ·	S -	-	-	K -
27.	Lingweromyces polysporus	K 5	5 V ⁺¹	ĸ	5	K 1	K	-	•	5 · v+1	-	S	K -	5	•	- v+1	K S
28.	Lipomyces iipojer	к	к	-	-	K 1	- V ⁺³	-	-	K	- V ⁺²	K +3	K V ⁺²	- v=+2	-	c+2	-
29.	L. starkeyi Phaffia vhodozvma	- V*2	-	-	-	к v+1e+2	ĸ	-	-	-	ĸ	ĸ	к	N V ⁺¹	-	5	V*2
31	Pichia angusta			-	-								-				ĸ
32	P anomala	K ⁺⁴			S ⁺¹				S*1	S ⁺¹							
33.	P. euphorbiaphila	K*3	-	S ⁺²	-			-			-	-	-			-	K*2
34.	P. guilliermondii	K ⁺²	-	-	-	-	-	-	-		-	-	-	K+3S+3		-	
35.	P. heimii	x	S ⁺²	S*2	S+4	K+2S+1		S*3	S ⁺⁴	K+2S+4	S*1	S*1	S ⁺³	K*3S*2		S ⁺³	-
36.	P. jadinii	K*2	х	-	-	K ⁺¹	K ⁺³	-			-	S^{+2}	S ⁺²	K ⁺²		-	K*2
37.	P. lynferdii	K ⁺³	-	х	-	-	-	-	-	K^{+2}	-	-	-	K^{+3}	-	K^{+2}	K+2S+3
38.	P. methanolica	K^{*4}	-	-	х	K^{*2}	-	-	-		-	-	-	K^{*3}	-	-	K^{*2}
39.	P. mexicana	$\mathrm{K}^{+1}\mathrm{S}^{+3}$	S^{+1}	-	S^{+2}	х			S^{+1}				S^{+1}				
40.	P. ofimaensis	-	S^{+3}	-	-		x	-	-	K^{+2}	K^{+1}	-	-		-	S*2	-
41.	P. ohmeri	K^{+4}	-	-	-	-	-	х	-	-	-	S^{+1}	K^{+2}	K^{+3}	-	-	-
42.	P. strasburgensis	K^{+4}	-	-	-	K^{+1}	-	-	х	-	-	-	-	K^{+3}	-	-	K^{+2}
43.	P. sydowiorum	$\mathrm{K}^{+4}\mathrm{S}^{+2}$	-	S^{+2}	-	-	K^{+2}	-	-	х	-	-	K^{+1}	S^{+1}		-	K ⁺²
44.	Saccharomycodes ludwigii	K^{*3}		-	-	-	S^{+2}		-	-	х	S^{+2}	S^{+2}	K^{+3}	-	-	
45.	Sporidiobolus ruineniae	K^{+2}	S^{+3}	-	-	-	-	K^{+1}	-	K^{+2}	K^{+1}	х	K^{+3}	K^{+2}	-	-	-
46.	S. salmonicolor	K^{*3}	-	-	-	K^{*1}	-	S^{+1}	-	S ⁺¹	K^{+3}	S^{+2}	х	K^{+2}	-	K^{+2}	K*2
47.	Sporobolomyces tsugae	$\mathrm{K}^{*3}\mathrm{S}^{*2}$	S^{+2}	-	S^{+3}	-	-	S^{+2}	S^{+3}	K^{+1}	S^{+2}	S^{+2}	S^{+2}	х	-	-	-
48.	Stephanoascus ciferrii		-	-	-	-	-	-	-	-	-	-	-	-	х	-	-
49.	Tremella encephala	K^{*3}	-	S^{+2}	-	-	K^{+2}	-	-	-	-	-	S^{+2}	-		х	-
50.	Williopsis californica	K+3S+2	S ⁺²	-	S ⁺²	-	-	-	S ⁺³	S ⁺²	-	-	S ⁺²	-	-	-	х

Table 1. (Cont'd.).

C No	Voost mooing	Phenotypes of seeded yeast strains (%)							
5.NO.	Y east species	Killer	Sensitive	Neutral					
1.	Arxula adeninovorans	10.2	10.20	79.59					
2.	Bensingtonia intermedia	16.32	6.12	77.55					
3.	B. naganoensis	10.2	14.29	7.55					
4.	Bullera pseudoalba	12.25	51.02	42.86					
5.	B. pyricola	14.29	77.55	18.37					
6.	Candida diddensiae	26.53	2.04	73.47					
7.	C. etchellsii	10.2	4.08	85.71					
8.	C. friedrichii	20.40	8.16	73.47					
9.	C. haemulonii	12.25	2.04	85.71					
10.	C. membranifaciens	10.2	12.25	79.59					
11.	C. pseudointermedia	22.45	6.12	73.47					
12.	C. shehatae	26.53	8.16	67.35					
13.	C. succinhila	6.12	0.00	93.88					
14	C valdiviana	6.12	22.45	73 47					
15	C restobii	4 08	16 33	79 59					
16	Clavispora lusitaniae	14 29	10.33	77.55					
10.	Cryptococcus albidus	10.2	20.4	75.51					
18	C gastricus	10.2	2 04	71.43					
10.	Debaryomyces castellii	12.24	10.2	77.55					
20	D bansenii	14 29	16.33	71.43					
20.	D. nanalansis	4.08	0.00	95.92					
21.	D. vanriiji	4.00 2.04	0.00	07.06					
22.	D. vanniju D. vanadao	2.04	0.00	97.90					
23.	D. yumuuue Fibulohasidium inconspicuum	2.04	2.04	93.92					
24.	Filohagidialla u coformana	20.4	20.4	07.33					
23.	Filobasiaiena neojormans	16.20	0.00	89.80 77 5					
20.	Filodasiaium uniguilulaium Kluuwanamuaaa nahanamua	10.52	0.10 26.52	11.5					
27.	Kiuyveromyces polysporus	14.29	20.55	05.20 45.2					
20.	Lipomyces lipojer	30.01	4.08	05.5					
29. 20	L. STARKEYI Di affi a al a da muna a	40.93	2.04	51.02					
30. 21	Phaffia rhodozyma Bishin swasta	14.29	2.04	85.71					
31.	Picnia angusta	0.00	0.00	100.00					
32.	P. anomala	10.2	28.52	61.22					
33.	P. euphorbuphila	12.25	2.04	85.71					
34.	P. guilliermondii	8.16	2.04	91.84					
35.	P. heimii	28.57	77.50	18.37					
36.	P. jadinii	14.29	8.16	77.55					
37.	P. lynferdu	26.53	8.16	67.35					
38.	P. methanolica	18.37	2.04	81.63					
39.	P. mexicana	6.12	26.53	71.43					
40.	P. ofunaensis	12.25	18.37	75.51					
41.	P. ohmeri	8.16	2.04	89.8					
42.	P. strasburgensis	16.32	4.08	79.59					
43.	P. sydowiorum	18.37	12.25	73.47					
44.	Saccharomycodes ludwigii	10.20	10.20	79.59					
45.	Sporidiobolus ruineniae	20.4	14.29	67.35					
46.	S. salmonicolor	30.61	20.4	48.98					
47.	Sporobolomyces tsugae	26.53	36.37	42.86					
48.	Stephanoascus ciferrii	2.04	0.00	97.96					
49.	Tremella encephala	8.16	12.25	79.59					
50.	Williopsis californica	14.29	42.86	50.02					

Table 2. Percentages of killer, sensitive and neutral phenotypes in yeast species isolated from dairy products.

Whereas, other yeasts that showed lesser killing activity were *Candida diddensiae* (26.53%), *C. friedrichii* (20.40%), *C. pseudointermedia* (22.45%), *C. shehatae* (26.53%), *Fibulobasidium inconspicuum* (20.40%), *Lipomyces lipofer* (30.61%), *Pichia heimii* (26.53%), *P. lynferdii* (26.53%), *Sporidiobolus ruineniae* (20.40%), *S. salmonicolor* (30.61%) and *Sporobolomyces tsugae* (26.53%). Several strong (K^{+3}) and very strong (K^{+4})

killing zones were produced by *Lipomyces starkeyi*, and other killer strains against sensitive strains.

A number of yeasts which, neither showed killing nor sensitive reactions even against the supper sensitive and killer strains are considered as neutral or resistant strains. The phenomenon of insensitivity towards killer yeasts generally occurs at the cell wall level. It is known that resistant (neutral) yeasts lack receptors necessary for the

formation of the link and thus for the action of the killer toxin (Marquina et al., 2002; Golubev, 2006). Taxonomically, different cell wall chemical compositions are used for classification of organisms, hence resistance causing insensitivity towards killer yeasts could be a taxon-related property as well (Golubev, 1998, 2006). In some studies Golubev (Golubev, 1992; Golubev et al., 1997) inferred that killer toxin effectiveness is inversely related to phylogenetic affinity (e.g. ascomycetous yeasts are usually insensitive to toxins produced by basidiomycetous species and vice versa). However, in the present studies, we observed a mixed effectiveness of killer yeasts against the neutral (insensitive) yeasts (Table 1). In this context, we emphasize (which Golubev (2006) also emphasized in his studies) that the use of killer toxins as a taxonomic tool should be preceded by a careful study of their KSP. Broad-spectrum killer toxins should be used for overall phylogenetic evaluation, while those characterized by a narrow range of activity may be used for clarifying relationships between more closely related species, or for grouping phenotypically similar strains before using molecular techniques [e.g. nucleotide composition in the D1/D2 domains and ITS regions of the ribosomal DNA (r-DNA)].

Similarly, a number of yeast strains showed strong killing activity against sensitive yeasts during cross reactions. In nature certain strains of killer yeasts dominate only in particular niches (Zorg *et al.*, 1988). Killer activity is one of the mechanisms of antagonism among yeasts during spontaneous fermentations and because of this mechanism killer strains may be used to avoid contamination by sensitive spoilage yeasts (Starmer *et al.*, 1987; Bussey *et al.*, 1988; Jacobs *et al.*, 1988; Longo *et al.*, 1990; Vagnoli *et al.*, 1993; Hidalgo & Flores, 1994).

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(Received for publication 7 September 2011)