Yeast Microflora of Some Aquatic Habitats in El-Minia Governorate, Egypt

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ABSTRACT: 269 yeast strains were isolated from water samples collected from different sites in Minia governorate. These included 126 strains from fresh water, 108 strains from sewage and 35 strains from wastewater from sugar-cane factory. On the basis of 23 different physiological and morphological merkmals, the isolated strains were assigned to 16 species belonging to 11 genera. Total yeast cell counts as well as spectra of yeast species were highly variable in tested water. Total yeast cell counts ranged between $3.0 \times 10^3 / l$ and $1.8 \times 10^6 / l$ for fresh water, $3.0 \times 10^4 / l$ and $3.0 \times 10^7 / l$ for sewage and $1.5 \times 10^6 / l$ and $2.6 \times 10^7 / l$ for wastewater from sugarcane factory. Debaryomyces hansenii, Rhodotorula mucilaginosa and Torulaspora delbrueckii were the dominant species in fresh water, whereas Debaryomyces hansenii, Thrichosporon beigelii, Rhodotorula mucilaginosa and Kluyveromyces marxianus were the dominant species in sewage and Saccharomyces cerevisiae, Kluyveromyces marxianus and Trichosporon beigelii were the dominant species in wastewater from sugar-cane factory. Yeast human pathogens, Trichosporon beigelii, Rhodotorula mucilaginosa and Candida albicans were encountered in water samples indicating that water in El-Minia governorate is also polluted by some pathogenic yeasts.

KEYWORDS: Yeast, aquatic habitats, Egypt.

The occurrence of yeasts in aquatic habitats was reported by several investigators(Kriss et al., 1952, 1967; Spencer et al., 1969, 1970, 1974a; Simard and Blackwood, 1971a, b: Viviani and Tortorano, 1976; Hinzelin and Lectard, 1978 and Hagler and Mendoca-Hagler, 1981). Data of these investigations support the idea that most yeasts of aquatic habitats have terrestrial origin and some species, such as Debaryomyces hansenii and Rodotorula spp., are physiologically adapted for survival and increase in population in water environment with increased nutrients. The correlation between yeast counts and pollution of water has been previously observed by Wollett et al.(1970), Viviana and Tortorano(1976), Vaatanen(1980a, b) and Hagler et al., (1981). This pollution is reflected on the chemical, physical and microbiological characteristics of water. This correlation is indicated from the observation that yeast cell densities in polluted water and sewage exceed that of nonenriched water. Also, yeast counts in sewage were reported to be higher than those recorde for fresh water(Cooke et al., 1960; Cooke & Matsuura, 1963, 1969; Cooke, 1965, 1970; Meyers et al., 1967; Crow et al., 1976 and Hinzelin and Lectard, 1979). Along the same line, pollution of fresh water streams receiving sugar-cane wastes from sugar-cane factories by yeasts was reported by Ahearn et al.(1968). Different yeast species including Saccharomyces exiguus and its anamorph Candida holmii as well as other Saccharomyces spp. have been reported to prevail in fresh water receiving sugar-cane waters(Ahearn et al., 1968).

Due to health hazard which might arise as a result of pollution of water with pathogenic yeasts, it was of interest to study yeast flora of water samples from different habitats in El-Minia governorate as a part of an extensive survey aiming to study yeast flora in different habitats in this governorate in order to evaluate its role in pollu-

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ting the environment.

Materials and Methods

Collection of water samples: 112 fresh water samples were collected from water streams of Ibrahimea canal and its irrigation branches at different localities in El-Minia governorate. These sites were located in Maghagha, Beni-Mazar, Matay, Samalot, El-Minia, Abo-Korkas, Malawy and Dermowas. 27 sewage samples were gathered from 2 different sewage drainage stations(A, B). Station A is found outside El-Minia City(at El-Minia University) where a limited number of population work, whereas sewage of station B(at Ard-Soltan) comes from different areas with a high number of population and receives wastewaters of different industries. 12 wastewater samples from sugar-cane factory(at Abo-Korkas) were collected from a canal receiving this wastewater. 25 ml proteins of each water samples were collected in sterile conical flasks and transported directly to the laboratory. Sampling was performed monthly for one year period.

Isolation and identification of yeast strains: 0.5 ml portions from the prepared decimal dilutions of water samples were spread on plates containing YM-agar medium(Lodder, 1970) adjusted to pH 3.5 and the plates were incubated at 28°C for 2-3 days. In case of sewage samples, an additional set of plates was incubated for the same period at 37°C. Developing yeast colonies were counted and after microscopic examination, yeast strains were isolated, purified, preserved on YM-agar slants and stored at 4°C. Yeast identification was performed on the basis of the various physiological and morphological characteristics as indicated by Lodder(1970), Barnett *et al.*(1983) and Kreger van Rij(1984).

Results and Discussion

Fresh water: Results of table 1 showed that, yeasts were detected in 59(ca. 53%) out of 112 fresh water samples collected from 8 different localities in El-Minia governorate and that, total

veast counts in the investigated fresh water samples ranged between $3.0\times10^3/l$ and $1.8\times10^6/l$. Results also showed that,f relatively high yeast total counts as well as more yeast species were recorded in water samples collected from Maghagha, Dermowas, Abo-Korkas and El-Minia, whereas low yeast counts and relatively few yeast species were found in fresh water from Beni-Mazar, Matay, Samalot and Malawy(Table 1). Other investigators reported that, clean fresh as well as marine waters usually contain below 100 yeasts/l(Van Uden & Ahearn, 1963; Ahearn et al., 1968, 1969; Cooke, 1970; Meyers et al., 1970; Ahearn, 1973; Spencer et al., 1974a; Colwell et al., 1976 and Hagler & Mendoca-Hagler, 1981), whereas yeast cell densities in polluted water frequently exceed 10³/l(Van Uden & Fell, 1968 and Ahearn, 1973). Moreover, a correlation was found between yeast counts and chemical, physical and microbiological parameters at various sites with different levels of pollution (Wollett et al., 1970; Simard & Blackwood, 1971b; Noel & Simard, 1973; Viviana & Tortorano, 1976; Vaatanen, 1980a,b and Hagler et al., 1981). In the light of the results presented in Table 1, it could be concluded that fresh water in El-Minia governorate is polluted by yeasts and the level of pollution varies among the different localities. The highest level of water pollution was observed in water samples collected from Maghagha, whereas the lowest was recorded in Samalot.

On the basis of morphological and physiological merkmals, 126 yeast strains isolated from fresh water samples were assigned to 12 species belonging to 9 genera(Table 1). It is clear that, spectra of yeast species as well as the dominant ones varied in fresh water samples collected from different localities. Debaryomyces hansenii, Rhodotorula mucilaginosa and Torulaspora delbrueckii occurred in high numbers in 6 different localities but were missed in two localities. It is possible that these species are physiologically adapted for survival in aquatic habitats than the others. This observation was previously reported for Debaryomyces hansenii and *Rhodotorula spp.* by other investigators(Cooke et al., 1960; Capriotti, 1962a,b; Fell, 1967, 1974; Hoppe, 1972, Ahearn, 1973 and Hagler & Ahearn,

Table 1. Yeast species isolated from aquatic habitats.

	Waste-	water from sugar- cane	lactory		12	100	1.5×10 ⁻⁶	2.0 × 10	1 1	8(2) 8(3)	12(5)	4(1) 2(1)	t	1 1	1 1	1 :
	Sewage			В	15 15		$1.9 \times 10^{\circ}$		14(6) 14(1)	14(2) $10(2)$	8(1) 8(3)	4		1	-	-
				Y	12 12	100	3×10^{-4} 1.8 × 10 ⁶		6(5) 10(2)	4(2) 6(3)		11,		1 1		1
		Эеппомая			10	- 1	1.2×10^{-4} 2.0×10^{4}	3	4(4) 2(1) 2(3)	7			4(3)			
r of Fresh water from locality			yweie	K Z	1	36	1.3×10^5	,	2 2(1)	4(2)	7 2	2(2)		5	; ;	
	ocality		4po-Korkas	7 14	10	3.6×10 ⁻⁴	3.8×10^{5}						2(2) 2(2)			
	ret trom l		El-Minia	14	8 57	1.8×10^{-4}	4.8×10⁴						(T)			inant.
	DA TO	Samalot			э 36	3×10-3	4.2×10 ⁴	; §	(Z)	2(2)	1 1	1 1 1	2(1)	1 1	1	ies is dom
			Katay	14	38	1.2×10^{-4}	7.1 \ r	6(4)		1 1		_			1 1	yeast speci
		9t.	Beni-Maz	14	43	1×10^{-4} 5.4 × 10 ⁵		4(4)	1	2(2)	11	1 1	11,		11	in which
	ed St ()	В	Kaghagh	14	12	1.4×10^{-4} 1.8×10^{6}		$8(4)^{**}$ $2(1)$	8(4) 4	2(1)	4	7	121	1		of samples in which yeast species is dominant.
Number	strains (269)			ig yeasts			0,4		%				4 8		٠ ا	* * = Number
				Number of samples tested Number of samples containing yeasts Percentage of veget occurrents	Total count of yeast cells//	410	Species Debaryomyces hansenii	Khodotorula mucilaqinosa* Trichosporon beiaelii*	Kluyveromyces marxianus Torulaspora delbrueckii	Saccharomyces cerevisiae Candida blankii	Hansenula polymorpha Cryptococcus albidus	Clavispora lusitaniae Rhodotorula aurantiaca	Debaryomyces vanriji Cryptococcus laurentii Candida alkicaas	Candida steelytica) cast numan pathogen, **

Table 2. Physiological and morphological properties of the isolated yeast species.

ಹ	ələilləq	54	0	100	100	100	09		0	0	100	0	20	100	100	100	100
Building of	pseudomycelium	42	0	100	100	100	09		0	0	100	0	20	100	100	100	100
<u> </u>	true mycelium	0	0	100	0	0	0	100	0	0	0	0	0	100	0	0	0
	secoebies	100	0	0	100	100	100	100	100	0	100	0	100	0	0	0	100
rth	೨,7₹	0	0	32	22	0	20	001	100	0	75	0	0	0	100	0	100
Growth at	3,2,2	33	48	63	82	20	09	0	100	0	100	0	100	100	100	100	100
	citrate	100	33	89	65	0	0	100	29	80	22	0	20	100	100	100	100
	succinate	100	71	100	82	25	40	80	20	80	75	33	0	100	100	100	100
	lotinnsm	100	92	001	88	28	20	001	100	100	100	33	100	001	100	100	100
	lactose	25	0	001	47	0	0	001	0	100	0	0	0	100	0	100	0
	maltose	100	100	8	001	92	001	001	100	100	100	0	100	100	100	100	100
ation	sncrose	100	100	96	001	83	001	001	001	100	100	33	001	001	001	100	100
Assimilation	гратпове	88	10	28	0	0	0	100	0	100	100	0	0	100	0	100	0
Ä	arabinose	54	52	001	47	0	0	100	29	100	0	0	100	100	100	100	100
	xylose	100	95	. 62	82	83	0	100	83	100	100	0	20	100	100	001	100
	ribose	96	52	00]	22	0	0	100	83	80	75]	0	20	100	100	100	100
	sorbose	29	62	90	65	22	0	100	29	100	100	0	001	0	1000	100	100
	galactose	100	06	001	001	001	80	100	83	100	1000	001	100	001	100	100	100
	lactose	0	0	0	9	0	0	0	0	0	0	0	0	0	0	0	0
uo	sncrose	75	0	0	53	33	20	99	29	0	25	0	20	0	8	100	0
Fermentation	maltose	54	0	0	41	33	40	28	0	0	22	0	0	0	100	100	0
Ferm	galactose	58	0	0	59	33	20	99	0	0	20	0	0	0	100	100	0
<u>.</u>	gncose	*96	0	0	100	100	100	68	100	0	100	0	20	0	100	100	0
umber of strains tested (269)			2	αc				18	12 1	0	8 1	9	₩	2	1 1	1 1	1
Number of strains tested (269)		48	4	38	34	24	20	H	-	10			·				
Species		Debaryomyces hansenii	Rhodotorula mucilaginosa	Trichosporon beioelii	Kluyveromyces marxianus	Torulaspora delbrueckii	Saccharomyces cerevisiae	Candida blankii	Hansenula polymorpha	Cryptococcus albidus	Clavispora Iusitaniae	Rhodotorula aurantiaca	Debaryomyces vanriji	Cryptococcus laurentii	Candida albicans	Candida steolytica	Pichia quilliermondii

1987).

Table 1 shows that the red yeast species *Rhodotorula mucilaginosa*, *Rhodotorula aurantiaca* and *Cryptococcus laurentii* were isolated from fresh water samples in low frequency(21% of the total isolates). Their occurrence in low frequency in fresh water could be used as an indicator for water pollution as previously shown by Spencer *et al.*(1970, 1974b); Simard(1971) and Hagler and Mendoca-Hagler(1981). Also, *Trichosporon beigelii*, a filamentous arthro-spore-forming yeast of basidiomycetious affinity, which has been suggested as a pollution indicator(Hinzelin & Lectard, 1976, 1978), was isolated frequently in this study particularly from fresh water samples collected from Maghagha.

Sewage water: Results presented in table 1 showed that, all the tested sewage samples contained yeast cells and that, yeast cell counts ranged between $3\times10^4/l$ and $1.6\times10^7/l$. It is also clear from table 1 that, the total count of yeast cells as well as the spectrum of yeast species in sewage from station B is higher than those from station A. Saccharomyces cerevisiae and Torulaspora delbrueckii were isolated in considerable numbers from sewage of station B but were missed in sewage of station A. These variations are expected since station B receives various wastewaters coming from different sources of higher populated areas indoor the city, whereas station A receives wastewater from low population areas. Cooke et al.(1960), Cooke & Matsuura(1963, 1969), Spencer et al.(1974 b) and Hinzelin and Lectard(1979) have shown that yeast counts in sewage, although highly variable, are much higher in sewage than in fresh waters. Counts up to $2\times10^8/l$ yeasts/l have been reported form sewage and 105/l were observed in domestic sewage.

Debaryomyces hansenii, Rhodotorula mucilaginosa, Trichosporon beigelii and Kluyveromyces marxianus were dominant and were represented by high number of yeast cells. Except for Kluyveromyces marxianus, the remaining 3 species were previously recorded as prevalent species in sewage by Hagler and Ahearn(1987).

Wastewater from sugar-cane factory: Our inves-

tigations on yeast flora of wastewater from sugarcane factory show that, yeast cell counts ranged between $1.5\times10^6/l$ and $2.6\times10^7/l$ (Table 1). Yeast counts were much higher than those of sewage. Wastewater from sugar-cane factory contains manu residues of sugar industry and is acidic with pH value between 4.8 and 5.8 which promote flurishing of yeasts. Saccharomyces cerevisiae was the most dominant species followed by Kluyveromyces marxianus and Trichosporon beigelii(Table 1). Saccharomyces spp. were reported by Ahearn et al. (1968) as dominant species in fresh water receiving sugar-cane wastes.

The physiological and morphological properties of the ioslated yeast species are presented in table 2. It is clear that, glucose fermenting ability, building of ascospores, true mycelium, pigmented colonies and arthrospores represented differential merkmals in our choosen merkmal set for the most dominant species.

It is worthmentioning that *Trichosporon beigelii*, *Rhodotorula mucilaginosa* and *Candida albicans*, which are known to be human pathogens(Rosalinde *et al.*, 1987), were isolated in this study from some fresh water samples, sewage water and wastewater of sugar-cane factory. Their occurrence in fresh water samples may represent health hazard to human and care should by taken to avoid this pollution.

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