

Pathogenic Fungi: an unacknowledged risk at coastal resorts? New insights on microbiological sand quality in Portugal

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ABSTRACT

Whilst the potential impact on beach users from microorganisms in water has received considerable attention, there has been relatively little investigation into microbial contaminants in sand. 33 beaches across Portugal were analyzed during a five year period (2006-2010) to determine the presence of yeasts, pathogenic fungi, dermatophytes, total coliforms, *E. coli* and intestinal enterococci in sand.

Our results showed that 60.4% of the samples were positive for fungi and that 25.2% were positive for the bacterial parameters. The most frequent fungal species found were *Candida sp.* and *Aspergillus sp.*, whereas intestinal enterococci were the most frequently isolated bacteria. Positive associations were detected among analyzed parameters and country-regions but none among those parameters and sampling period.

Regarding threshold values, we propose 15 cfu/g for yeasts, 17 cfu/g for potential pathogenic fungi, 8 cfu/g for dermatophytes. 84 cfu/g for coliforms, 250 cfu/g for *E. coli*, and 100 cfu/g for intestinal enterococci

Highlights:

- 60.4% of the analyzed marine beach sand samples were positive for the studied fungi
- *Candida sp.* and *Aspergillus sp.* were the most frequent fungal species found
- 25.2% of the analyzed marine beach sand samples were positive to the studied bacteria

- Positive association was found between the presence of yeasts and coliforms
- Proposal of microbiological threshold values for marine beach sands

Keywords: Sand Quality, Beach, Fungi, Bacteria, Bioindicators, Microbiological Thresholds

Introduction

Beaches, both coastal and inland, are amongst the most significant sites for water based recreation. Bathing, contact waters sports and appreciation of the general ambience are all valuable uses made of these areas. The importance of good water quality, particularly microbiological quality is reflected in the statutory limits now in place in most of the developed world. The influence of water quality on bathers' health was extensively reviewed by the World Health Organisation (WHO, 2003). The European Community Bathing Water Directive (2006) places greater emphasis on measures for the protection of the (bathing) public. Development of quality of life and health of the population is a complex function of different conditions of the living environment. Exposure to organic and inorganic pollutants as well as to the wide spectrum of microorganisms is part of this problem (Matavulj et al., 2005). Potentially pathogenic microorganisms in sands include faecal bacteria as found in the water but also mycological agents including fungi, yeasts and moulds (Vieira et al., 2001; Mancini et al., 2004; Abdallah et al., 2005; Sato et al., 2005; Gomes et al., 2007). Several American (Vieira et al., 2001; Sato et al., 2005; Gomes et al., 2007) and European studies (Brandão et al., 2002; Mancini et al., 2004; Abdallah et al., 2005; Matavulj et al., 2005) have been performed on this issue and although the World Health Organisation report considered sand contamination it made no recommendations as to standards, neither do they feature in current European or US legislation.

With this emphasis on public protection, it is perhaps surprising that there have been no similar bacterial or mycological limits for sand, though more beach users come into contact with the sand than the water. Bathers and those engaged in water sports will invariably also be in contact with the sand, yet there are many beach-goers who never enter the water (LaLiberte and Grimes, 1982; Obiri and Jones, 2000; Alm et al. 2003; Whitman, 2006).

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To date, no correlation has been shown between health and pathogenic fungi in beach sands (Prado et al., 1994; Brandão et al., 2002) but it may be expected that the public exposed to sand contaminated with fungal pathogens are at an increased exposure risk through direct contact with their skin and mucous membranes or by inhaling spores. Beach users, both people and animals, may themselves be partially responsible for contaminating beach sands, either by carrying microorganisms that are left on the sand or by discarding organic litter, including refuse from fishing, which provides the food and humidity necessary for the preservation and development of yeasts, bacteria, and moulds (Brandão et al., 2002). Recently Heaney et al., 2009, performed an epidemiological study on beach users and observed strong association between sand contact and enteric illness at marine beaches.

The Blue Flag scheme (FEE) is one of the principal mechanisms for the maintenance of beach quality standards and it currently monitors 29 different indicators which cover Environmental Education and Information, Water Quality, Environmental Management, Safety and Services (Associação Bandeira Azul da Europa, 2009). One matter insufficiently addressed to date is whether contact with beach sand should be of concern to public safety. In order to address this concern, a project was created within Portugal, as a joint venture between the Portuguese Blue Flag Association, the National Institute of Health and the Environmental Protection Agency. The present work reports on the microbiological quality of the sand at selected beaches, including the distribution of bacteria and fungi collected from beach sands and suggest threshold values for selected microbiological indicators.

MATERIALS AND METHODS

Study Design

The Portuguese coast was divided into five regions (North, Central, Lisbon and Tagus Valley, Alentejo and Algarve) and a total of 164 different sand beaches were analyzed during the period 2006-2010. All the selected beaches are classified as good quality beaches, holding the Blue Flag award. Thirty three were analyzed every year during the referred sampling period and those were the ones selected for the present study. From the 33 beaches, four were from the North, four from the Central, nine from Lisbon and Tagus Valley (LVT), one from Alentejo and 15 from Algarve. Three sand samples were collected from each beach during the summer season (June, July and August).

Sample collection

Samples were collected at a depth of about 10 cm, with sterile gloves into a sterile plastic container, from the middle of the dry sand section of the shore, as three equidistant sub-samples which mixed into one composite, represented each beach as a whole. The first sampling event took place at start of the bathing season, the second in the peak of the summer (highest concentration of users), and the last before the end of the bathing season. Samples were processed within 18 hours and transported refrigerated to the laboratory.

Sample analysis

Mycological

Each sand sample (40 g - not oven-dried prior to processing, retaining thus its natural water content), was diluted in 40mL of sterilized distilled water, agitated for 30 min at 100 rpm and 0.2 mL of this suspension was spread, in triplicate, onto Petri dishes containing Mycobiotic agar for dermatophytes (up to 3 weeks incubation at 27.5°C) and malt extract agar (2%) with cloramphenicol (0.05g/L) for non-dermatophyte fungi (5-7 days at 27.5°C of incubation). Fungal identification was carried out by macroscopic and microscopic (using lactophenol blue staining) observation of colonies for filamentous fungi, using identification atlases and using the biochemical identification galleries ID32C (bioMérieux SA, Marcy-l'Etoile, France) for yeasts. Results were reported as the average count of the three replicas, in colony forming unit per gram of sand (c.f.u/g) when detecting the fungal species more frequently associated to human infections, as indicated in Table I. Other moulds are reported when being the predominant species, or with extremely high counts – over 500 colony forming unit per gram of sand (cfu/g). *Histoplasma spp.*, *Coccidioides spp.*, *Exophiala sp.*, *Fonsecae spp.*, *Phialophora spp.* were always reported. This methodology is accredited by the ISO17025 (IPAC L0425).

Bacteriological

Detection of total coliforms/*Escherichia coli* (*E. coli*) and enterococci was accomplished through the use of Colilert and Enterolert with Quanti-Tray (IDEXX Laboratories, Maine, USA) respectively. Extraction fluid resulting from 50g of sand extracted with 500 ml of sterile distilled water and agitating for 30 min at 100 rpm was processed according to manufacturer's instructions, both for Colilert and Enterolert and for each sample. Bacteria detected and reported individually, as indicated in Table 1. The methods complied with HPA NHS W18 and ASTM D6503-99 respectively.

Statistical analyses

Descriptive analysis for parameters was performed, using means, 95% confidence intervals for mean, median, minimum and maximum values for continuous variables. Counts and percentages are categorical variables. Pearson correlation coefficient was used to measure the association between parameters. The Mean values of parameters were compared between years, months and regions of samplings using ANOVA. The Brown-Forsythe robust tests of equality of means were used to fortify the results of ANOVA when the variances of groups were significantly different. Fisher's Least Significant Difference (LSD) or the Dunnett post hoc tests for multiple comparisons were used assuming homogeneity of variances or not, respectively. 90, 95, 97.5 and 99 percentiles were calculated to establish thresholds for the parameters, using the distributions of values for the five years of sampling. Values of $p \leq 0.05$ were accepted as statistically significant. IBM SPSS version 18.0 software was used to perform all statistical analysis.

Results

During a five year period 33 beaches distributed along the Portuguese coast (Figure 1) were analyzed. A total of 495 sand samples were collected and processed in mycological and bacteriological laboratories.

From the 495 analyzed sand samples, 299 (60.4%) were positive for the investigated fungal species and 125 samples (25.2%) were positive for the three bacterial groups analyzed. One hundred and sixty five samples (33.3%) were negative for all the parameters analyzed whereas 100 (20.2%) of them were positive for both parameters (Table 2).

The number of fungal colony-forming units of yeasts and dermatophytes tends to increase along the sampling periods, whereas the bacterial number of colony forming units tends to decrease from the first to the second sampling period and is maintained from the second to the third one (Figure 2). Nevertheless, no positive association was found between the the three sampling periods in yeasts ($p=0.321$), potential pathogenic fungi ($p=0.056$), dermatophytes

($p=0.171$), coliforms ($p=0.976$), *E. coli* ($p=0.90$), and intestinal enterococci ($p=0.332$).

Concerning the distribution of the number of fungal colony-forming units along the Portuguese coast (Table 3), the highest value was observed in beaches belonging to the Central region of Portugal (average of 498.9 cfu/g) whereas the lowest values were detected in beaches from Algarve (average of 62.5 cfu/g).

Associations among some of the analyzed parameters and country regions were found. In fact, significant differences were found between LVT plus Alentejo and Algarve regarding potential pathogenic fungi ($p=0.010$) and dermatophytes ($p=0.033$). Each one of these parameters had higher counts in LVT and Alentejo regions. Significant differences ($p=0.004$) were also found between North plus Central and Algarve in coliforms distribution, with North and Central having tenfold the counts of the Algarve.

Yeasts were detected in 25.4% of the sand samples (Table 2). Of the detected yeasts, 67.5% belonged to *Candida* species, with an average of 5.8 cfu/g (range 0-1934). A positive association was found between the presence of *Candida* and other yeasts ($p=0.00$). Potential pathogenic fungi were found in 47.9% of the samples (Table 2). From those, *Aspergillus* sp. was the predominant genus, detected in 153 samples, with an average of 0.87 cfu/g (range 0-250). In fact, a positive association was found between the presence of *Aspergillus* sp. and potential pathogenic fungi group ($p=0.000$). *Fusarium* sp. was detected in 115 samples (23.2%), followed by *Scedosporium* sp. (30 samples, 6.1%), *Scytalidium* sp. (22 samples, 4.4%), *Chrysosporium* sp. (17 samples, 3.4%), and *Scopulariosis* sp. (13 samples, 2.6%). Regarding dermatophytes, they were found in 14.3% of the analyzed samples and the most predominant genus was *Trichophyton* sp., detected in 69 sand samples, with an average of 1.5 cfu/g (range 0-152) (Table 2). *Microsporum* sp. was detected in three samples (0.61%) whereas no *Epidermophyton* isolates were recovered.

For bacteria (Table 2), intestinal enterococci were detected in 100 samples, followed by coliforms in 96 samples and *E. coli*, detected in 34 sand samples. Coliforms were, however, the bacteriological parameter with highest average of colony forming units per gram of sand (36.1).

To assess possible relationships between the fungal and bacterial colonization of sands, data regarding total coliforms, *E. coli*, intestinal enterococci, and fungi were examined for

1 correlations. A positive association was found between yeasts and coliforms ($p=0.005$). All the
2 other parameters did not show positive associations ($p>0.05$).

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4 Interesting positive associations were found among the analyzed fungi, namely between
5 *Aspergillus* sp. and *Chrysosporium* sp. ($p=0.00$), *Chrysosporium* sp. and *Fusarium* sp. ($p=0.00$),
6 *Fusarium* sp. and *Scytalidium* sp. ($p=0.04$), and also between *Scopulariopsis* sp. and
7 dermatophytes ($p=0.00$).
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11 As there is no current values for health impacts of fungi and bacteria on beach sand, we
12 herewith aim to propose thresholds values for the six parameters studied, based on the
13 microbiological quality of sand beaches during the five years of study, considering that, as
14 stated above, the batch of beaches used in this study is considered of good quality and hence
15 meeting the blue flag award criteria (Associação Bandeira Azul da Europa, 2009). The threshold
16 values we propose are the ones obtained when 95% of the samples analyzed are below those
17 values. Thus, 15 cfu/g is the calculated threshold for yeasts, 17 cfu/g for potential pathogenic
18 fungi, 8 cfu/g for dermatophytes and may serve as a national or regional references.
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30 **Discussion**

31 Quality of life and the health of people is a complex function of different conditions in the living
32 environment. Sand beaches may act as a passive element of cumulative pollution and can
33 become contaminated by garbage, animal waste or water itself carrying pathogenic
34 microorganisms and parasites (Sato et al., 2005). The microbiological context of sediments at
35 the sediment–water interface in bathing waters is receiving increased attention (Arakel, 1995).
36 There is evidence that faecal indicator and pathogenic bacteria survive for longer periods in
37 sediments than in the overlying water and it has been proposed that sediments serve as sinks
38 for fecal bacteria with the potential to pollute the overlying bathing waters (Ashbolt et al., 1993;
39 Nix et al., 1993; Ghinsberg et al., 1994; Howell et al., 1996). Nevertheless, and despite the well
40 established cause-effect relationship between faecal pollution of recreational waters and
41 gastroenteritis outbreaks (Prüss et al., 1998), no epidemiological evidence of transmission
42 between microbiological contamination of the sand and human infection has been found (World
43 Health Organization, 2003). Persons with different immunological status are all beach users but
44 immunosuppressed individuals are more susceptible in acquiring infections caused by
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1 microorganisms that were found in sand beach (Burton et al., 1987; Mendes et al., 1993).
2 Moreover, a recent epidemiological study published by Heaney et al., 2009, report that sand
3 contact activities were associated with enteric illness at beach sites. Variation in beach-specific
4 results suggests that site-specific factors may be important in the risk of illness following sand
5 exposure.
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10 This study provides original and comprehensive data on the microbiological content of sand
11 along the Portuguese coast during a five year period. Our study was performed only in summer
12 and using dry sand. Results of a study performed by Brandão et al., 2002 showed that
13 microbiological contamination is higher in dry sand. Several other authors also showed that
14 sand contamination during summer was higher than during spring and highest in dry sand
15 (Mancini et al., 2005; Sato et al., 2005). A high proportion of the samples analyzed in our study
16 showed zero or low colony-forming counts, for both fungi and bacteria. Nevertheless potentially
17 harmful fungi and bacteria can be found in 66.5% of all the samples.
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26 There is some evidence of a lower occurrence of microorganisms in areas with a hot and dry
27 climate. In fact, Algarve was the region with the lowest 5 year, all beaches, average number of
28 colony-forming units per gram of sand (62.5) when compared with beaches from the North and
29 Central regions of the country (466.3 and 498.9, respectively).
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34 The potential health risk associated with the exposure to the contaminated sand should be
35 higher in summer, especially for the people, usually children and youngsters, who tend to stay
36 longer. Higher levels of bacteria in dry sand, that it is not under the influence of the tides, may
37 indicate that the main source of faecal contamination is not the seawater, but instead the
38 heavily polluted water of creeks and runoff (Vidal and Lucena, 1997; Sato et al., 2005). Our
39 results showed that the first sampling period (prior to the bathing season) was the one with
40 higher number of colony forming units of bacteria, decreasing along the summer season. This
41 fact may be explained by the decrease of surface runoff and the increase of solar duration and
42 intensity along the sampling periods. The presence of, higher levels of yeasts and
43 dermatophytes in dry sand may indicate human contamination. Inversely to the trend for
44 bacteria, the number of colony forming units increases from the first to the third sampling period.
45 This may be explained by the increase of beach users in the third sampling period as well as by
46 the higher resistance of fungi to solar exposure (Anderson et al., 2000). In fact, fungal spores
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can survive between 25 and 360 days in the environment whereas the survival of enteric bacteria on the surface of dry sand may be less. (Carillo-Muñoz et al., 1990; Papadakis et al., 1997). Thus, dry sand is the chosen matrix for fungi analysis whereas wet sand or water are better suited for quality assessment using enteric bacteria (Brandão et al., 2002),

The diversity of fungal species was high and 299 (60.4%) of the samples were positive to species considered as potentially harmful, especially in immunocompromised persons. In fact, Lacaz, 1991, states that the variety of yeasts found in the soil and in seawater play an important role in the medical pathology, causing cutaneous mycosis. *Candida* species were found in 7.1% of samples whereas *C. albicans* was detected in only 0.8% of the samples. Absence or low incidence of *C. albicans* has also been recorded by other researchers (Roses Codinachs et al., 1988; Figueras et al., 1992). Sato et al., 2005, on the contrary, obtained a higher value (17%) in the 96 sand samples analyzed. These differences could be explained due to the fact that our samples are from beaches with good quality (with Blue Flag award) and probably also due to the discrepancy in the number of analyzed samples.

Potential pathogenic and allergenic fungi (like *Aspergillus* species) were the most common microorganisms found in our samples (in 30.9%). The same situation was reported by Mendes 2 et al., 1998, also in Portuguese beaches, and by Sarquis and Oliveira (1996) and by Sharaf (2005) in Brazilian and Egyptian beaches, respectively.

Our samples revealed 14.3% of samples contaminated with dermatophytes. Studies performed by Sousa, 1990, also in the Portuguese central coastal area showed a much higher prevalence of dermatophytes (in 42% of the samples analyzed). In our study other filamentous fungi with keratinophilic activity were also found. In fact, *Scytalidium sp.*, *Chrysosporium sp.* and *Scopulariopsis sp.* were isolated from 4.4%, 3.4% and 2.2% of the analyzed samples. Positive associations found among *Chrysosporium sp.*, *Fusarium sp.*, *Scytalidium sp.*, *Scopulariopsis sp.* and dermatophytes could be explained by its keratinophilic nature. Keratinolytic fungi such as *Chrysosporium sp.* and *Trichophyton,sp.* are known as agents of human and animal infections and have been isolated from beach sand of European countries like Sweeden (Vidal et al., 1966), Spain (Ulfig et al., 1997) and Italy (Salvo and Fabianno, 2007), among others.

In our study, we found positive associations between coliforms and yeasts, and also between coliforms and *E. coli*. This association could be explained by possible sand contamination with

1 surface runoff from surrounding areas polluted with organic material and human and animal
2 waste. In an epidemiological study carried out on two beaches in Malaga, Spain, faecal index
3 microorganisms were highly significantly correlated with dermatophyte fungi on one of the
4 beaches and only *E. coli* showed a significant correlation with *Candida albicans*. At other
5 studied beach, intestinal enterococci showed the best correlation with dermatophyte fungi
6 (Borrego et al., 1991).
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11 The current study provides a useful baseline for assessment of sand beach microbiological flora
12 and as a warning to the occurrence of potentially harmful fungi and bacteria in these
13 environments. Few studies have been performed to determine guideline values and in the
14 absence of any specific guidelines based on epidemiological study, it is not possible to set firm
15 limit levels. Preventive measures and health risk assessments have not been considered. This
16 study with a high number of samples from Blue Flag beaches analyzed over a five year period
17 will contribute towards establishing those values. Thus, for fungi, we propose 15 cfu/g for
18 yeasts, 17 cfu/g for potential pathogenic fungi and 8 cfu/g for dermatophytes. The calculated
19 threshold values for bacteria were much lower than the ones recommended for coastal bathing
20 waters by Directive 2006/7/EC of the European Parliament and of the Council of 15 February
21 2006. Moreover, bacteria survive less time in dry sand than water. Thus, we recommend the
22 adoption of the threshold limits for coastal bathing waters (<100 cfu/100 mL for intestinal
23 enterococci and, <250 cfu/100 mL for *E. coli*). The derived thresholds merely set a result in a
24 country wide context but do help in ranking sand beaches. Moreover, threshold limits have
25 proved to be useful to alert beach managers to the danger of potential deterioration in beach
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10 11 12 **DECLARATION OF INTEREST**

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14
15 The authors report no conflicts of interest. The authors alone are responsible for the content
16 and writing of the paper.

17 18 19 **AUTHORS' CONTRIBUTIONS**

20
21 CV, LF, LR, JB conceived and designed the study. RS, FCF, MAC, BW, RR, HP, CV, CP, and
22 JB performed the experiments; RS, CV, LF, LR, EP, JB analyzed the data, EP being the
23 statistician; RS, CV, LF, JB wrote the manuscript. All authors have read and approved the final
24 version of the manuscript.

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Fig.1. Collection points along the Portuguese coast.

Fig. 2 Microbiological quantification of bacteria (a) and fungi (b) along three sampling periods (1st -June, 2nd -July,3rd - August).

Table 1. Microbiological parameters analyzed in sand samples.

Mycological parameters			
Yeasts	Potential pathogenic/allergogenic moulds	Dermatophytes	Bacteriological parameters
<i>Candida albicans</i>	<i>Aspergillus fumigatus</i>	<i>Trichophyton sp</i>	Coliforms
<i>Candida sp</i> (Other species)	<i>Aspergillus niger</i>	<i>Microsporum sp</i>	
<i>Cryptococcus neoformans</i>	<i>Aspergillus sp</i> (Other species)	<i>Epidermophyton sp</i>	<i>Escherichia coli</i>
Other yeasts	<i>Chrysosporium sp</i>		Intestinal enterococci
	<i>Fusarium sp</i>		
	<i>Scytilidium sp</i>		
	<i>Scedosporium sp</i>		
	<i>Scopulariopsis sp</i>		
	Others		

Table 2. Microbiological characterization of the analyzed sand samples during the five years of study.

	Negative	Positive to fungi	Positive to bacteria	Positive to bacteria and fungi	Positive only to fungi	Positive only to bacteria
2006	35	61	21	18	44	3
2007	28	64	27	24	44	3
2008	29	63	32	25	30	7
2009	31	63	18	13	50	5
2010	43	48	27	20	36	6
Total	166	299	125	100	204	24

Table 3. Mycological and bacteriological parameters analyzed in the 495 sand samples

	No. of positive samples (%)	CfU/g average <i>per sample</i> (range)
Yeasts	126 (25.4)	8.7 (0-1997)
Potential pathogenic fungi	237 (47.9)	4.5 (0-267)
Dermatophytes	71 (15.3)	1.7 (0-152)
Coliforms	96 (19.4)	36.1 (0-2420)
<i>E. coli</i>	34 (6.9)	11.8 (0-2420)
Intestinal enterococci	100 (20.2)	10.1 (0-2420)

Legend: CfU/g – colony forming units per gram of sand

Figure(s)
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