

Studies on Fungal and Bacterial Population of Air-conditioned Environments

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ABSTRACT

In tropical countries such as Brazil, there is not enough information about microbial contaminants in indoor environments with air conditioning systems. Microbial monitoring of such environments is important for the quality of human life. The aim of this work was to assess the fungal genera and bacterial morphotypes occurring in such environments. Air samples were taken indoors and outdoors from a public auditorium, a hospital, a company and a shopping center during the 2001 winter by using a six-stage impactor Millipore M air T[®]. Twenty-one fungal genera were identified. Bacterial morphological groups found were Gram positive and negative rods and Gram positive coccus.

Key words: Sick building syndrome, fungi, bacteria, air conditioner, indoor air quality, air ecology

INTRODUCTION

The aim of air conditioning systems is to provide occupants with a more comfortable environment. Nevertheless, such artificial environments may be favorable to fungi, bacteria, protozoan and mites growth, which may bring health risks to users, either by hypersensitivity or infections (Morey et al., 1986; Vincent et al., 1997; Cooley et al., 1998). According to the American Society of Heating, Refrigerating and Air Conditioning Engineers (ASHRAE, 2000), the health effects caused by microorganisms that are in indoor environments with air conditioning systems can be infective or immunological.

The term sick building syndrome (SBS) is commonly used for health disturbances related to

indoor air quality (IAQ). Lack of cleaning and checking out of the heating, ventilation and air conditioning systems (HVAC) may allow microbial growth, which causes rhinitis, bronchitis, pharyngitis, pneumonia, conjunctivitis and keratitis in the users. According to ASHRAE (2000), if 20% or more users of such environments present the above-mentioned symptoms, these places are considered under SBS. It must be stressed that several other factors may aggravate SBS, including outdoor air quality, air insufficiency or bad air distribution, inefficient control of air temperature and humidity, improper project or changes in the original building project and lack of monitoring, checking out and cleaning of HVAC systems.

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SBS probably has a multi-factorial etiology with chemical, physical, biological and psychosocial factors that interact with each other, resulting in symptoms and discomfort (Baker, 1989; Norback et al., 1990). Thus, SBS may influence public health and economy negatively, because some of these diseases decrease workers' productivity and cause their absence at work.

Indoor air quality effects on health and the large use of HVAC systems in indoor environments reinforce the importance to reproduce and standardize methods for microbial monitoring in such indoor environments. For tropical countries, microbial assessment of environments with HVAC systems is even more important due to the high microbial biodiversity associated with high temperature and relative air humidity, favoring microbial growth. The objective of this study was to determine the fungal community and bacterial levels in four sites that use HVAC systems.

MATERIAL AND METHODS

Experimental design

The present work had the following experimental design: four sites (a public auditorium, a hospital, a company and a shopping center), two types of air samples indoor air (AIM) and outdoor air (AOM), both collected by impactor M air T[®] in five replicates for each sample, distributed as fixed point in each collecting place.

Samples

The collecting of the samples was conducted in the winter of 2001, in the city of Londrina, PR, Southern of Brazil. During this season, there is wide use of environment insulation, which probably increases the number of bioaerosols in indoor air, coinciding with high levels of respiratory infections. A six-stage impactor Millipore M Air T[®] collected the samples, so that the air was impacted in Petri dishes containing specific medium for fungi and bacteria, as well as appropriate design for equipment use (cassettes M Air T[®]). During the collection, the air sampler remained about one meter above floor level.

Each sample collected with the impactor Millipore M Air T[®] contained 100 L of air for fungi and 1000 L for bacteria. All the collection patterns (time of air impactation, culture medium, agar concentration, season of the year, sites and

collection points) have been determined in previous experiments.

Isolation and taxonomic study of fungi

For inoculation through air sample impactation, the culture medium was semi-solid malt extract agar (malt extract 20 g, glucose 20 g, peptone 2 g, agar 7 g, dH₂O 1000 mL, pH 5.5), being that the dishes were incubated for 5 days at 28°C. The number of fungi colonies present was assessed in the third and fifth days of incubation, and the number of colony forming units was determined per m³ of air (CFU m⁻³ air). After that, the colonies considered morphologically distinct were isolated and again compared with each other, so that only one isolated strain representative of each morphological group found was selected. The representative strains were preserved at room temperature in sealed glasses with paraffin, containing sterile saline (0.85% NaCl) and agar plates obtained from the culture of each fungus.

The taxonomic identification of fungi was performed considering the morphological characteristics of the vegetative mycelium and the reproductive structures (Barnett and Hunter, 1972; Koneman and Roberts, 1992; Hoog and Guarro, 1995; Larone, 1995; Ainsworth et al., 1995; Silveira, 1995; Alexopoulos et al., 1996; Lacaz et al., 1998).

Some fungi strains had poor growth or did not showed any reproductive structures in the initial conditions of cultivation, this strains were submitted to different culture conditions, such as temperature, exposure to black light UV, sun light and incubation time and others culture media such as Sabouraud Dextrose agar (casein peptone 5 g, meat peptone 5 g, glucose 40 g, agar 15 g, dH₂O 1000 mL, pH= 5.6), Potato Dextrose agar (potato extract 4 g, glucose 20 g, agar 15 g, dH₂O 1000 mL, pH= 5.6) and agar V8 (juice V8 200 mL, CaCO₃ 3 g, agar 15 g, dH₂O 800 mL, pH= 6.2) and incubated at temperatures 20, 22, 24, 26 e 28°C and at room temperature from 7 to 10 days.

Isolation and morphological study of bacteria

For bacteria study was used Tryptic Soy agar plus Benlate (casein peptone 17 g, soy peptone 3 g, dextrose 2.5 g, sodium chloride 5 g, dibasic potassium phosphate 2.5 g, agar 15 g, Benlate 20 mg; dH₂O 1000 mL, pH= 6.2). After the air samples inoculation, the Petri dishes were incubated for three days at 28°C. The number of

bacteria colonies were assessed in the first and third days of incubation, and the number of CFU m^{-3} air was determined. Following that, the colonies considered morphologically distinct were isolated and compared with each other again, being that only one isolated strain representative of each morphological group found was selected. The representative strains were stored at $-70^{\circ}C$ in glycerin 20%.

For the study of the morphological groups, the strains was stained by Gram method and examined by microscopic and morphological characteristics of the bacterial cells was evaluated, ordering them in positive and negative Gram rods and coccus.

RESULTS

Fungi

The number of CFU of fungi in the samples of AIM was smaller in the public auditorium and in the hospital and bigger in the shopping center and in the company. On the other hand, for the AOM samples, the number of CFU showed differences only on the air of hospital, was smaller than other sites. It has been observed that in all the sites the outdoor air had a bigger number of CFU of fungi when compared with the indoor air. The ratio AIM:AOM of CFU of fungi for the different sites analyzed also showed differences, being smaller in the public auditorium and bigger in the company (Table 1).

Table 1 - Ratio between indoor and outdoor number of fungi colony forming unit (CFU m^{-3} air) collected by stage air impactor Millipore M Air TTM. AIM = air indoor sample; AOM = air outdoor sample; I:O = ratio between indoor and outdoor samples. The values represent means of five replications and the Standard Error (SE) was estimated by sample from each place.

Samples	Public Auditorium	Hospital	Company	Shopping Center
	CFU fungi m^{-3} air	CFU fungi m^{-3} air	CFU fungi m^{-3} air	CFU fungi m^{-3} air
AIM	178	194	312	338
SE	58.86	39.45	36.80	38.00
AOM	507	338	454	526
SE	85.32	29.90	43.66	29.43
I:O ^a	0.35	0.57	0.69	0.64

^a The values from indoor and outdoor ratio of number of fungi colony forming unit (CFU m^{-3} air). I:O = < 1.5 environment in proper condition; I:O = 1.5 a 2 environment regular condition; I:O = >2 environment not proper condition.

The genera *Aspergillus* and *Epicoccum* were found in all the AIM and AOM air samples of the four sites. The genera *Penicillium* was found in all AIM samples but in the AOM samples this genera was not found in the shopping center samples. Among the three most common fungi found, the population of *Aspergillus* was biggest in all the AIM and AOM samples, except for the public auditorium. The biggest number of the *Penicillium* population was observed in the AIM air samples, and the genera *Aspergillus* and *Epicoccum* in the AOM air samples (Table 2). In the AIM samples the genus *Alternaria* was found only in the public auditorium and AOM in all the samples of the four sites. Others genera found such as *Botrytis*, *Chrysosporium*, and *Cladosporium*, *Leptosphaeria* [doubtful], *Mucor*, *Neurospora*, *Phoma*, *Rhizopus*, *Trichoderma* and *Verticillium* were found only in

AOM samples. On the other hand, some fungi were found in a few AIM samples, in this way, the *Microsporum* was observed only in the AIM air of the hospital. The genus *Fusarium* was not found either in the public auditorium or in the hospital. The genera *Emmonsia* did not occur in the company and the *Myriodontium* [doubtful] in the hospital. The genus *Curvularia* was found in AOM samples of the public auditorium and in the AIM of the company and the shopping center. The genus *Acremonium*, occurred in the AIM and AOM samples of the public auditorium and in the AOM samples of the shopping center. Non-identified fungi were found in the AIM and AOM samples of the four analyzed sites, because it was not possible to obtain reproductive structures in the several conditions tested (Table 2).

Table 2 - Number of colony forming unit (CFU m⁻³ air) of fungi isolated from air of indoor (AIM) and outdoor (AOM) at different locals discriminated between genera. Air samples were collected by stage air impactor Millipore M Air T™.

Genera	Public Auditorium		Hospital		Company		Shopping Center	
	AIM	AOM	AIM	AOM	AIM	AOM	AIM	AOM
<i>Acremonium</i>	2	1	-	-	-	-	-	4
<i>Alternaria</i>	6	50	-	8	-	30	-	22
<i>Aspergillus</i>	20	76	100	78	186	198	186	234
<i>Brotrytis</i>	-	3	-	2	-	-	-	-
<i>Chrysosporium</i>	-	3	-	8	-	-	-	8
<i>Cladosporium</i>	-	3	-	-	-	6	-	6
<i>Curvularia</i>	-	2	-	-	2	-	2	-
<i>Drechslera</i>	-	-	4	-	4	-	18	-
<i>Emmonsia</i>	2	-	4	-	-	-	6	-
<i>Epicoccum</i>	10	98	2	44	4	86	12	70
<i>Fusarium</i>	4	-	2	-	-	-	-	-
<i>Leptosphaeria</i> (doubtful)	-	153	-	116	-	64	-	70
<i>Microsporum</i>	-	-	6	-	-	-	-	-
<i>Mucor</i>	-	24	-	14	-	20	-	42
<i>Myriodontium</i> (doubtful)	4	-	-	-	16	-	2	-
<i>Neurospora</i>	-	17	-	2	-	34	-	2
<i>Penicillium</i>	102	45	62	4	64	2	68	-
<i>Phoma</i>	-	7	-	48	-	8	-	24
<i>Rhizopus</i>	-	-	-	-	-	-	-	10
<i>Trichoderma</i>	-	18	-	8	-	6	-	32
<i>Verticillium</i>	-	4	-	6	-	-	-	-
Not identified	28	3	14	-	36	-	44	2
Total (CFU fungi m ⁻³ air)	178	507	194	338	312	454	338	526

The relative frequency of fungi varied according to the genera, sites and samples. The genera that occurred most in the public auditorium were, in decreasing order, in the AIM samples *Penicillium* (57.3%), *Aspergillus* (11.24%) and *Epicoccum* (5.63%); as for the AOM samples, *Leptosphaeria* [doubtful] (30.18%), *Epicoccum* (19.33%) and *Aspergillus* (14.99%). In the hospital, in the AIM samples the genera were *Aspergillus* (51.56%), *Penicillium* (31.96%), *Microsporum* (3.09%), *Emmonsia* (2.06%), and *Drechslera* (2.06%); and in the AOM samples the genera *Leptosphaeria* [doubtful] (34.31%), *Aspergillus* (23.08%), *Phoma* (14.2%) and *Epicoccum* (13.02%), *Mucor* (4.14%), *Alternaria* (2.37%), *Trichoderma* (2.37%) and *Chrysosporium* (2.37%). In the company, the most frequent fungi genera in the AIM were *Aspergillus* (59.62%), *Penicillium*

(20.51%), and *Myriodontium* [doubtful] (5.13%), and in the AOM samples, *Aspergillus* (43.61%), *Epicoccum* (18.94%), *Leptosphaeria* [doubtful] (14.1%), *Neurospora* (7.49%), and *Alternaria* (6.61%). In the shopping center there was predominance of the genera *Aspergillus* (55.02%), *Penicillium* (20.12%), *Drechslera* (5.33%), and *Epicoccum* (3.55%) in the AIM samples, and the genera *Aspergillus* (44.5%), *Epicoccum* (13.31%), *Leptosphaeria* [doubtful] (13.31%), *Mucor* (7.98%), *Trichoderma* (6.08%), *Phoma* (4.56%), and *Alternaria* (4.18%) in the AOM samples. The relative values for the strains of genera represented in brackets were calculated based on the total number of fungi of both the AIM and AOM samples (Table 3).

Table 3 - Fungal community isolated from indoor air samples (AIM) and outdoor (AOM) in different locals, represented by percentage of each genera. Air samples were collected by stage air impactor Millipore M Air T™.

Genera	Public Auditorium		Hospital		Company		Shopping Center	
	AIM	AOM	AIM	AOM	AIM	AOM	AIM	AOM
<i>Acremonium</i>	1.12	0.2	-	-	-	-	-	0.76
<i>Alternaria</i>	3.37	9.86	-	2.37	-	6.61	-	4.18
<i>Aspergillus</i>	11.24	14.99	51.56	23.08	59.62	43.61	55.02	44.5
<i>Brotritis</i>	-	0.59	-	0.59	-	-	-	-
<i>Chrysosporium</i>	-	0.59	-	2.37	-	-	-	1.52
<i>Cladosporium</i>	-	0.59	-	-	-	1.32	-	1.14
<i>Curvularia</i>	-	0.4	-	-	0.64	-	0.59	-
<i>Drechslera</i>	-	-	2.06	-	1.28	-	5.33	-
<i>Emmonsia</i>	1.12	-	2.06	-	-	-	1.78	-
<i>Epicoccum</i>	5.63	19.33	1.03	13.02	1.28	18.94	3.55	13.31
<i>Fusarium</i>	2.25	-	1.03	-	-	-	-	-
<i>Leptosphaeria</i> (doubtful)	-	30.18	-	34.31	-	14.1	-	13.31
<i>Microsporum</i>	-	-	3.09	-	-	-	-	-
<i>Mucor</i>	-	4.73	-	4.14	-	4.41	-	7.98
<i>Myriodontium</i> (doubtful)	2.25	-	-	-	5.13	-	0.59	-
<i>Neurospora</i>	-	3.35	-	0.59	-	7.49	-	0.38
<i>Penicillium</i>	57.3	8.88	31.96	1.18	20.51	0.44	20.12	-
<i>Phoma</i>	-	1.38	-	14.2	-	1.76	-	4.56
<i>Rhizopus</i>	-	-	-	-	-	-	-	1.9
<i>Trichoderma</i>	-	3.55	-	2.37	-	1.32	-	6.08
<i>Verticillium</i>	-	0.79	-	1.78	-	-	-	-
Not identified	15.72	0.59	7.21	-	11.54	-	13.02	0.38
Total (%)	100	100	100	100	100	100	100	100

Table 4 - Ratio between indoor and outdoor number of bacteria colony forming unit (CFU m⁻³ air) collected by stage air impactor Millipore M Air T™. AIM = air indoor; AOM = air outdoor; I:O = ratio between indoor and outdoor. The values represent means of five replications and Standard Error (SE) was estimated by sample from each place.

Samples	Public Auditorium	Hospital	Company	Shopping Center
	CFU bacteria m ⁻³ air	CFU bacteria m ⁻³ air	CFU bacteria m ⁻³ air	CFU bacteria m ⁻³ air
AIM	16.20	54.60	67.60	84.40
SE	4.82	6.73	7.65	8.51
AOM	38.60	41.00	93.00	104.60
SE	8.71	4.84	4.39	4.38
I:O	0.42	1.33	0.73	0.81

^aThe values from indoor and outdoor ratio of number of bacteria colony forming unit (CFU m⁻³ air), I:O = < 1.5 environment in proper condition; I:O = 1.5 a 2 environment regular condition; I:O = >2 environment not proper condition.

Bacteria

The number of CFU of total bacteria in the AIM and AOM samples from the different sites was

smaller in the public auditorium and in the hospital, and bigger in the shopping center and in the company. When compare the number of CFU

indoor and outdoor, was observed that only in the hospital, AOM population was smallest than AIM air samples. The ratio AIM:AOM concerning the number of bacteria CFU in the different sites analyzed also showed differences, being smaller in the public auditorium and bigger in the hospital (Table 4). In the indoor air samples there was

predominance of Gram positive rods for the public auditorium, hospital and shopping center, except for the company, where Gram positive coccus was predominant. For the outdoor air samples, predominance of Gram negative rods was observed in all the sites (Table 5).

Table 5 - Number of colony forming unit (CFU m⁻³ air) of bacteria isolated from air of indoor (AIM) and outdoor (AOM) at different locals discriminated between morphological group. Air samples were collected by stage air impactor Millipore M Air TTM.

Morphological group	Public Auditorium		Hospital		Company		Shopping Center	
	AIM	AOM	AIM	AOM	AIM	AOM	AIM	AOM
Gram-positive coccus	3.60	3.10	19.00	6.20	34.40	17.20	32.60	21.80
Gram-positive rods	9.60	3.40	22.00	9.00	16.20	11.00	37.80	13.20
Gram-negative rods	3.00	32.10	13.60	25.80	17.00	64.80	14.00	69.60
Total (CFU m ⁻³ air)	16.20	38.60	54.60	41.00	67.60	93.00	84.40	104.60

Table 6 - Bacteria community isolated from indoor air samples (AIM) and outdoor (AOM) in different locals, represented by percentage of each morphological group. Air samples were collected by stage air impactor Millipore M Air TTM.

Morphological group	Public Auditorium		Hospital		Company		Shopping Center	
	AIM	AOM	AIM	AOM	AIM	AOM	AIM	AOM
Gram-positive coccus	22.22	8.03	34.80	15.12	50.89	18.50	38.63	20.84
Gram-positive rods	59.26	8.81	40.30	21.95	23.96	11.83	44.78	12.62
Gram-negative rods	18.52	83.16	24.90	62.93	25.15	69.67	16.59	66.54
Total (%)	100	100	100	100	100	100	100	100

The relative frequency varied according to the morphological group, sites and samples. In the AIM samples from the public auditorium there was predominance of Gram positive rods (59.26%), followed by Gram positive coccus (22.22%) and Gram negative rods (18.52%). However, in the AOM samples there was predominance of Gram negative rods (83.16%), followed by Gram positive rods (8.81%) and Gram positive coccus (8.03%). In the AIM samples of the hospital Gram positive rods predominated

(40.30%), followed by Gram positive coccus (34.80%) and Gram negative rods (24.90%), whereas in the AOM samples the Gram negative rods were more frequent (62.93%), followed by Gram positive rods (21.95%) and Gram positive coccus (15.12%). In the company, the most frequent morphological groups in the AIM samples were Gram positive coccus (50.89%), followed by Gram negative rods (25.15%) and Gram positive rods (23.96%). In the AOM samples the highest frequency was of Gram

negative rods (69.67%), followed by Gram positive coccus (18.50%) and Gram positive rods (11.83%). In the AIM samples from the shopping center, there was predominance of Gram positive rods (44.78%), Gram positive coccus (38.63%) and lower occurrence of Gram negative rods (16.59%). At the same time, in the AOM samples, the highest frequency was of Gram negative rods (66.54%) and the lowest frequency was of Gram positive coccus (20.84%) and Gram positive rods (12.62%) (Table 6).

DISCUSSION

In the public auditorium, company and shopping center, both indoor and outdoor air samples showed contamination fungi below the acceptable levels, when compared with many health organization standardization. The World Health Organization (WHO, 1988) recommends until 500 CFU m⁻³ air, the Healthy Building International (HBI) (Rao et al., 1996) concentration less than 750 CFU m⁻³ air, the National Institute of Occupational Safety and Health (NIOSHI-USA) (Jensen and Schafer, 1998) up to 1000 CFU m⁻³ air, and the Brazilian Health Ministry (ANVISA, 2000) 750 CFU m⁻³ air. Nevertheless, according to Rao et al. (1996), for qualitative indoor air evaluation, levels ranging from 100 to 1000 CFU of fungi m⁻³ air could be considered. However, Jensen and Schafer (1998) point out that the NIOSHI-USA, in its manual of analytical methods, reinforces the fact that a low concentration of microorganisms in indoor air environments does not mean they are healthy, but needs to identify the microorganisms to evaluate the conditions of the air-conditioning systems.

For the hospital environments, the WHO (1988) recommends not more than 50 CFU of fungi m⁻³ air. In this study, nevertheless, the fungal contamination levels were 194 CFU m⁻³ air for indoor samples. This level for fungal contamination is four times the WHO limit, which could result in a transmission of several diseases especially in immune-depressive patients.

In the indoor air evaluated in England and USA, the CFU of fungi were three times high than in this work, but the fungi genera were the same (Jones, 1999). The ratio AIM:AOM found in all samples was lower than 1.5, is controversy for Brazilian regulations that consider this level as safety air, however other authors (Rao et al., 1996)

considered the ratio level up to 1 the air with potential risk to occur an amplification of microbial populations such as fungi and bacteria.

Probably the most common fungi isolated from indoor air environment belong to the Deuteromycota division (imperfect fungi) and the most frequently genera found are *Penicillium*, *Aspergillus*, *Eurotium*, *Wallemia*, *Cladosporium* and *Alternaria* (Wanner et al., 1993). In the present study, we found the same results, most of them were belong Deuteromycota division and the most common genera were also the same.

The *Penicillium* population was biggest in indoor air and *Aspergillus*, *Epicoccum* and *Alternaria* in outdoor air. The same results were found in a hotel (Mcgrath et al., 1999) and a school (Cooley et al., 1998) with HVAC system in the USA.

The number of bacteria in the four sites evaluated remained within the maximum acceptable microbiological limits for the quantitative patterns considered. According to Morey et al. (1986) the American Conference of Governmental Industrial Hygienists (ACGIH) has as pattern that the concentration of *Bacillus sp.* and Gram negative bacteria should not over 500 CFU m⁻³ air. According to The HBI (Rao et al., 1996), they considers that concentrations lower than 750 CFU m⁻³ air of viable bacteria indicate that the building is free contamination if there are not infectious or allergenic species.

These values, however, are controversial, since there is a great variation as to what is considered adequate by different authors. Toth (1992) suggests that the counting of human normal flora bacteria above 200 CFU m⁻³ air be considered high. On the other hand, Hood (1990) states that 500 CFU m⁻³ air of Gram negative bacteria is sufficient to suspect of problems in the indoor air quality. For hospital environments, the maximum number of bacteria CFU allowed by the WHO (1988) is 100 CFU m⁻³ air. In the indoor air samples, the values do not surpass 100 CFU.

The air samples evaluated from home and school in England and USA (Jones, 1999) had nine times more bacteria than found in this study in the places evaluated. But (Li et al., 2001) evaluating many shopping centers in Hong Kong found different bacteria levels on the weekend when the number of people inside the building increased.

Aeroecology of indoor air environments among different countries around the world showed that the fungi genera found in these places generally are the same. However the number of CFU is high

in template zone when compared with tropical zone. The homogenized community that are living in the indoor air certainly does not correspond that what occur in the outdoor air environments, when the focus is biodiversity and size of population in a tropical and template zones, this fact should indicate that the human activity is building an artificial ecosystem, that can be favoring some genera of hazardous microorganisms and increase the risk of health inside the building.

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RESUMO

Em países tropicais como o Brasil, não há informação suficiente sobre contaminantes microbianos em ambientes internos com sistemas de ar condicionado. Monitoramento microbiano em tais ambientes é importante para a qualidade de vida humana. O objetivo deste trabalho foi identificar os gêneros de fungos e morfotipos de bactérias que ocorrem em tais ambientes. Amostras de ar foram coletadas dentro e fora de um auditório público, um hospital, uma empresa e um shopping center durante o inverno de 2001 utilizando um impactador de ar de seis estágios Millipore M air T[®]. Vinte e um gêneros de fungos foram identificados. Foram encontrados grupos morfológicos de bactérias bastonetes Gram positivos e negativos e cocos Gram positivos.

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