

Research Paper

## Survey of molds, yeast and *Alicyclobacillus* spp. from a concentrated apple juice productive process

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### Abstract

Bacteria and molds may spoil and/or contaminate apple juice either by direct microbial action or indirectly by the uptake of metabolites as off-flavours and toxins. Some of these microorganisms and/or metabolites may remain in the food even after extensive procedures. This study aim to identify the presence of molds (including heat resistant species) and *Alicyclobacillus* spp., during concentrated apple juice processing. Molds were isolated at different steps and then identified by their macroscopic and microscopic characteristics after cultivation on standard media at 5, 25 and 37°C, during 7 days. Among the 19 isolated found, 63% were identified as *Penicillium* with 50% belonging to the *P. expansum* specie. With regards to heat resistant molds, the species *Neosartorya fischeri*, *Byssoschlamys fulva* and also the genus *Eupenicillium* sp., *Talaromyces* sp. and *Eurotium* sp. were isolated. The thermoacidophilic spore-forming bacteria were identified as *A. acidoterrestis* by a further investigation based on 16S rRNA sequence similarity. The large contamination found indicates the need for methods to eliminate or prevent the presence of these microorganisms in the processing plants in order to avoid both spoilage of apple juice and toxin production.

**Key words:** concentrated apple juice, *Alicyclobacillus acidoterrestis*, heat resistant mold, *Penicillium expansum*.

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### Introduction

Apple juice belongs to the most frequently consumed types of fruit juices worldwide (53). The necessity to implement and optimize fast and efficient methods for quality control appears as a consequence of this use, and should be done throughout the whole processing procedure, beginning in the orchard (Silva *et al.*, 1997).

Some species of molds may cause serious postharvest diseases in apples (Kupferman 1986). Among them, *Penicillium expansum* is reported as being responsible for major decay on apples. Studies have shown that this mold is found in soil, on plant surfaces, in dump tank or flume water (Spotts and Cervantes, 1993), in contaminated wooden

bins (Sanderson and Spotts, 1995) and in the atmosphere (Amiri and Bompeix, 2005). Furthermore, this mold has the ability to produce patulin (Doores, 1983), a mycotoxin reported to cause oxidative damage to the DNA in human cells, which plays a role in mutagenesis and cancer initiation (Liu *et al.*, 2003).

Heat resistant molds (HRM) are among the microorganisms of great importance in the spoilage of heat-processed fruit juices, such as apple juice. Representative species are found in the genera *Byssoschlamys*, *Neosartorya*, *Eupenicillium*, *Talaromyces*, (Murakami *et al.*, 1998; Suresh *et al.*, 1996; Tournas, 1994) *Eurotium* (Splittstoesser *et al.*, 1989; Yildiz and Coksoyler, 2002) and *Paecilomyces* (Peña *et al.*, 2004; Piecková and Sam-

son, 2000). Some of the heat resistant molds can cause both spoilage of fruit products and produce toxic and sometimes carcinogenic compounds (Tournas, 1994; Ugwuanyi and Obeta, 1999).

*Alicyclobacillus acidoterrestris* is a thermoacidophilic spore-forming bacterium (ATSB) which is able to spoil acidic juices (Bahçeci and Acar, 2007; Chen *et al.*, 2006; Eguchi *et al.*, 1999; Groenewald *et al.*, 2009; McKnight *et al.*, 2010; Yamazaki *et al.*, 1996). Spoilage by *Alicyclobacillus* has become a problem for the apple juice industry and effective solutions should be found to control its development. The spoilage generally is manifested as an off-flavour and an off-odour of a medicinal or chemical nature due to the formation of guaiacol and halophenols (Chang and Kang, 2004; Chen *et al.*, 2006; Yamazaki *et al.*, 1996), leading to consumer rejections (Zierler *et al.*, 2004).

Apples rejected by the rigid selection criteria for the fresh fruit consumption market are used for juice processing. Microorganisms mentioned above may be present leading to a health risk for consumers and/or high economic losses due to juice deterioration. This study aims to survey molds (including heat resistant ones), yeast and *Alicyclobacillus* spp. in a concentrated apple juice manufacturing process.

## Materials and Methods

### Sampling

Samples were obtained at different stages from a concentrated apple juice processing plant in Brazil. The samples (collected in the off-season period) were taken at the following steps: apples in the reception (A); washed apples (WA); wash water (W) (recycled in a close system); must (M); bagasse (B); before pre-concentration (BPC); after pre-concentration (APC); enzymatic treatment (ET); before ultra-filtration (BUF); after ultra-filtration (AUF); before concentration (BC); after concentration (AC) and final product (FP). All samples were collected (approximately 250 mL per step) in sterile sampling containers, on the same day. Then they were stored under refrigeration using reusable ice for transport during 6 h. In the laboratory, the containers were placed in a refrigerator and the samples were analyzed within 24 h. Thirty samples of apples were collected randomly to represent the real state of the fruits. During the analysis, pieces of different apples were mixed to maintain the randomness of the tests.

Samples from different part of the process were tested in order to provide comprehensive data on the occurrence of molds and yeasts, heat resistant molds (HRM) and *Alicyclobacillus* at different stages of processing. Two visits were made in the processing plant and the aforementioned samples were taken in each one. Figure 1 presents the flow diagram of the production process indicating the sample localizations.

### Molds and Yeast enumeration

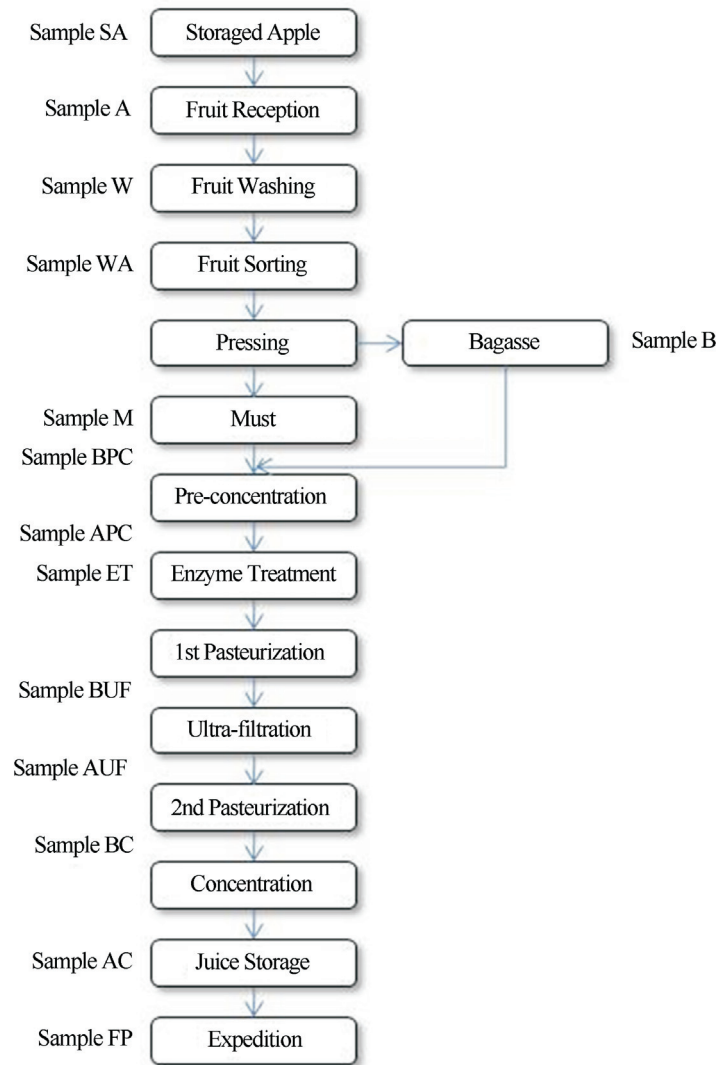
Molds and yeasts were detected by diluting 25 g of apples and bagasse samples in 225 mL of 0.1% peptonate water, followed by a two minute homogenization in a stomacher (ITR, Model 1204). Juice and liquid samples were just diluted in water (10 mL of the sample in 90 mL of the diluent). Subsequently, serial dilutions were made using 0.1% peptonate water and then samples were plated in duplicate, using Potato Dextrose Agar (PDA, Biolife®, Milan, Italy) acidified to pH 3.5 with 10% tartaric acid solution. All plates were then incubated at 25 °C and colonies were counted after 3-5 days. The results were expressed as CFU/mL or CFU/g, depending on the kind of sample (Silva *et al.*, 1997).

### Heat resistant molds enumeration

Apples (previously mixed in a sterile blender) and samples of bagasse were placed each one in a sterile plastic bag and homogenized (100 g of the samples plus 100 mL of sterile distilled water) in a stomacher (Pro-analise®, Model Boit-Sto1) for 4 min. Two 50 mL portions of homogenized samples were then transferred to sterile test tubes and heat shocked in a water bath (Tecnal®, Model 184 ± 0.1 °C) at 80 °C for 30 min. Concurrently, 50 mL samples of concentrated apple juice (70°Brix) were diluted with 50 mL of sterile distilled water and then also heated shocked. Liquids samples (35° Brix or less) were analyzed without any previous dilution and two 50 mL aliquots of each sample were subjected to the same heat treatment previously described. Samples Brix was measured in a refractometer (AR200 Reichert, USA). After heating, duplicate samples of 50 mL were cooled and each one was combined with 100 mL of acidified (pH 3.5) PDA (double agar concentration) supplemented with 50 mg/L of rose bengal and 4 mg/L of cloranfenicol. Subsequently, samples were distributed in eight Petri plates which were placed in plastic bags and sealed to prevent drying. The incubation was performed at 30 °C for up to 30 days. Most viable ascospores germinated and formed visible colonies within 7 to 10 days (Beuchat and Pitt, 2001). The results were expressed as cfu/mL or cfu/g, depending on the kind of sample.

### Mold identification

The mold identification was made by macroscopic and microscopic observations using identification keys as described by Pitt and Hocking (Pitt and Hocking, 1999). Each strain isolated on PDA was transferred to MEA (Malt Extract Agar) and CYA (Czapek Yeast Extract Agar) and G25N (25% Glycerol Nitrate Agar) media in duplicate trials. Cultures were grown on these three standard media at 25 °C for 7 days. Those inoculated on CYA were additionally incubated at 5 °C and 37 °C for the same period. Furthermore, Czapek agar with 20% sucrose (CY20S) was prepared to help in the identification of genera suspected to be xerophilic. After incubation, the diameters of macro-



**Figure 1** - Concentrated apple juice flow diagram and sampling localizations.

scopic colonies from the underside were measured and macroscopic characteristics such as color, texture and exudates, were analyzed for each colony. Microscopic structures were investigated with a microscopic (Bioval®) by using 0.1% lactofuchsin stain or lactophenol fungal stain (Cotton blue). Yeasts were not identified in this study; only molds were identified (heat resistant or not).

In addition, this investigation identified the mold present in a storage apple (SA) with visual signs of blue mold invasion. For this, a colony of mold from an apple were scraped and transferred to PDA and incubated for 5 days at 25 °C. After the isolation, the strain was sent to André Tosello Foundation (Campinas, SP, Brazil) for further identification and to be deposited at the Collection of Tropical Cultures (CCT).

#### *Alicyclobacillus* isolation and enumeration

The isolation and enumeration of *Alicyclobacillus* spp. (ATSB) was realized according to Eguchi *et al.* 1999.

In order to count ATSB, samples of concentrate juice were diluted prior to analysis (10 mL of juice in 90 mL of sterile distilled water). The other liquid samples (10 mL) were diluted in 90 mL of BAT (*Bacillus alicyclosterrestris*) broth. Samples of apples and bagasse (25 g) were blended with 225 mL of BAT broth in a stomacher for 2 min. The diluted samples were heated shocked at 80 °C for 10 min, cooled at 40-45 °C and their aliquots were plated on BAT agar by the pour plate technique. All plates were incubated at 50 °C for 4 days (plates were monitored for up to 10 days). After incubation time, the colonies were enumerated and the results were expressed in cfu/g or cfu/mL.

Concurrently, a technique for detecting was also performed with the same samples. In the detection method, the samples were prepared as described above and heated shocked at 80 °C for 10 min. Subsequently, the samples were incubated at 50 °C for 24 h to enrich the cultures. An aliquot of 1 mL of the enrichment culture was directly pour plated with BAT agar and incubated at 50 °C for 4 days

(plates were monitored for up to 10 days). This result was expressed as the absence or presence of *Alicyclobacillus* (ATSB).

### *Alicyclobacillus* morphological characterization and identification

Single colonies present at the surface of BAT agar were picked off and streaked onto the same medium used for isolation. Plates were incubated at 50 °C for 48 to 72 h and then, the overall microscopic morphology and the presence or characteristics of spores were investigated. Strains with characteristic morphology were inoculated onto Nutrient agar (Biolife®, Milan, Italy) at pH 7 and incubated at 50 °C to confirm the acidophilic nature of the isolates and discard the presence of acid tolerant bacilli.

The isolated *Alicyclobacillus* strains were sent to André Tosello Foundation (Campinas, Brazil) for further identification on basis of standard biochemical and morphological tests and also based on 16S rRNA sequence similarity. Also, the isolated strains were deposited at the Collection of Tropical Cultures (CCT).

## Results

### Molds and Yeast

The counts of molds and yeast in this study are shown in Table 1. In order to express the results, each strain was encoded according to the initial letters of the sample which was isolated and preceded by a number that represents the sequence of isolation.

A total of 13 strains of molds were isolated of which 12 were identified as belonging to the genus *Penicillium*: 1 strain from sample SA (1SA), 1 strain from sample A (1A),

2 strains from sample W (1W; 2W), 2 strains from sample WA (1WA; 2WA), 3 strains from sample B (1B; 2B; 3B), 1 strain from sample M (1M) and 2 strains from sample BPC (1BPC; 2BPC). From this total, the species was encoded as 1SA, 1A, 1W, 1WA, 1B and 2BPC was identified as *Penicillium expansum*. The strain 1SA was deposited in the Tropical Collection Cultures and re-encoded as *Penicillium expansum* CCT 7549. The strain isolated from sample AC was identified as *Talaromyces* sp. (1AC). Only yeasts (not identified in this study) were verified in the samples from ET, BUF, AUF and BC.

The macroscopic and microscopic description of molds isolated in this study is shown below. Furthermore, some pictures of their structures are exhibited in Figure 2.

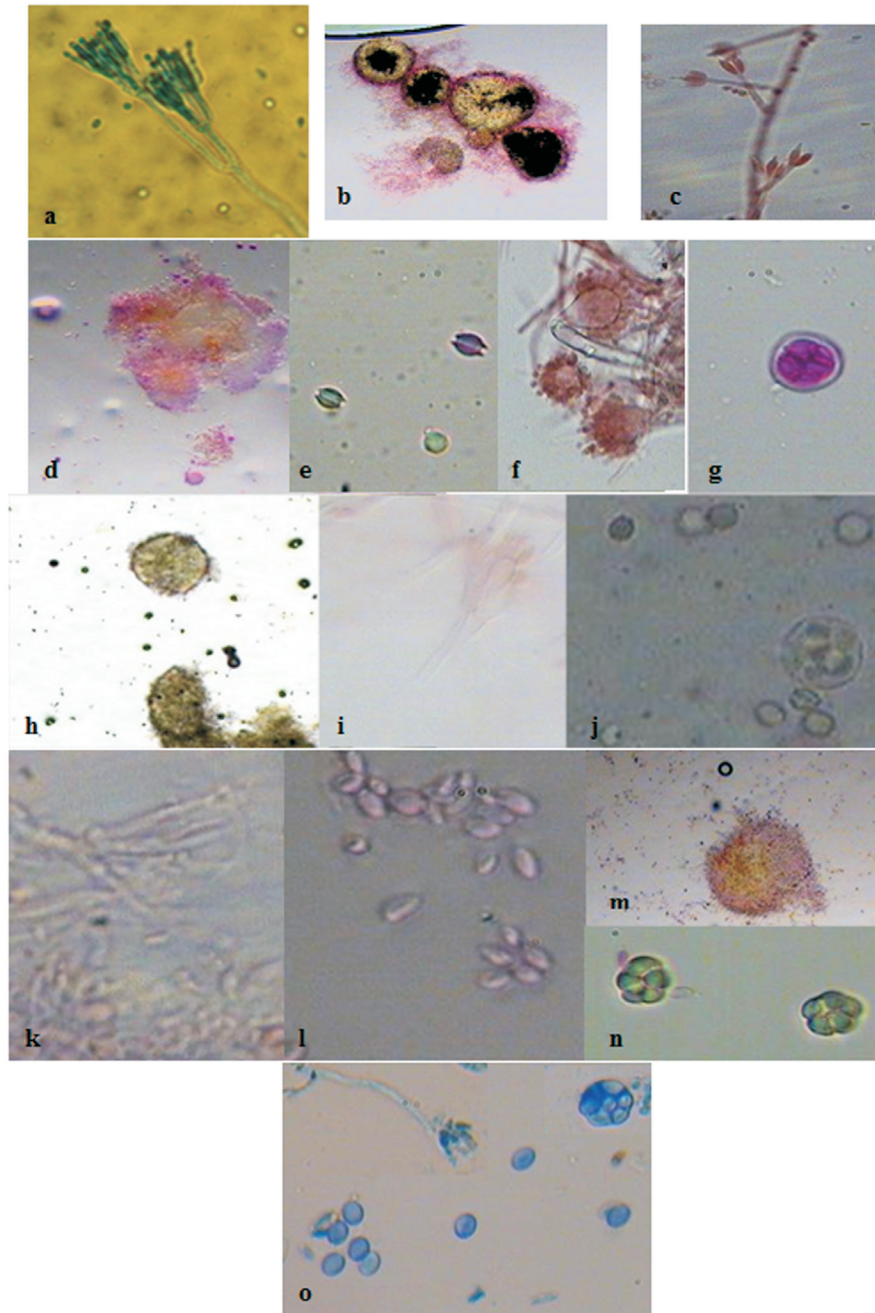
The isolates identified as *Penicillium* sp. (2W, 2WA, 2B, 3B, 1M and 1BPC) showed macroscopically colonies with different characteristics of color, texture and size, depending on the strain. It was observed that all strains grow on CYA, MEA and G25N at 25 °C but not on CYA at 37 °C and 5 °C. Microscopic observation revealed the presence of different types of penicilli and spherical or ellipsoidal conidia.

The isolates 1SA, 1A, 1W, 1WA, 1B and 2BPC were identified as *P. expansum*. Their colonies on CYA (25-35 mm diameter) were colonies centrally colored from dull green to slightly brown, surrounded (annularly) by a thick external layer of white color, clear to pale yellow exudates, tufted surface velutinous to floccose, moderate sporulation and a deep brown underside (caramel). The colors of colonies on MEA (20-35 mm diameter) ranged as seen on CYA to slightly greyer with orange exudates and a pale underside. Colonies on G25N (15-20 mm diameter) were centrally cream colored, surrounded by a thin white external layer with a dull brown underside. No growth was observed at 5 °C and 37 °C. Microscopically, conidiophores with smooth walls bearing terminal penicilli terverticillate (The strain 1SA, encoded as *P. expansum* CCT 7549, is represented in the Figure 2a) and smooth walled ellipsoidal conidia (4 to 5 µm long) were observed. In addition, 14.5-19.0 µm long metula and 10.0-14.0 µm long phialides were observed.

The strain codified as 1AC showed colonies on CYA of 60-70 mm of diameter, a floccose appearance, uniformly pale olive brown color and a pale underside. Colonies on MEA (covering the whole Petri dish) were very similar to those in CYA. Colonies on G25N (15-17 mm of diameter) were similar to those on CYA. At 37 °C on CYA, colonies covering the whole Petri dish were similar to those at 25 °C. No germination was observed at 5 °C. Microscopically (Fig. 2 k, l, m, n) the strain developed the characteristic of a HRM showing yellow gymnothecia with closely interwoven hyphae and ascus characteristic of the teleomorph *Talaromyces*. The conidia (around 5 µm long) were strictly cylindrical to pyriform. *Talaromyces* is associated with an anamorphic state characteristic of *Penicillium*,

**Table 1** - Mean values of molds and yeast in different stages of a concentrated apple juice process line.

Sample (code)	Mean values of molds and yeast molds and yeast
Apples at the reception (A)	1.3 x 10 <sup>5</sup> cfu/g ± 0.57
Wash water (W)	6.6 x 10 <sup>5</sup> cfu/mL ± 0.71
Washed apples (WA)	1.3 x 10 <sup>7</sup> cfu/g ± 0.63
Bagasse (B)	4.5 x 10 <sup>5</sup> cfu/g ± 0.57
Must (M)	9.3 x 10 <sup>4</sup> cfu/mL ± 1.16
Before pre-concentration (BPC)	7.0 x 10 <sup>5</sup> cfu/mL ± 0.37
After pre-concentration (APC)	< 10 cfu/mL ± 0.00
Enzymatic treatment (ET);	1.5 x 10 <sup>2</sup> cfu/mL ± 0.64
Before ultra-filtration (BUF)	3.0 x 10 <sup>2</sup> cfu/mL ± 0.80
After ultra-filtration (AUF)	2 cfu/mL ± 0.55
Before concentration (BC)	5 cfu/mL ± 0.67
Concentrate apple juice (AC)	1.7 x 10 <sup>2</sup> cfu/mL ± 0.87
Final product (FP)	< 10 cfu/mL ± 0.00



**Figure 2** - Microscopic structures: (a) *P. expansum* CCT7549; (b) *Eupenicillium* sp. 1BPFr (cleistotecium structure); (c) *Eupenicillium cinnamopurpureum* 1WAr (*Penicillium* anamorph state); *Eurotium* sp. 1WAr structures: (d) cleistotecia, (e) ascospores, (f) *Aspergillus* (anamorph state), (g) asci; *Neosartorya fischeri* 2WAr structures: (h) cleistotecia, (i) *Aspergillus* (anamorph state), (j) asci and ascospores; *Talaromyces* sp. 1AC structures: (k) *Paecilomyces* (anamorph state), (l) conidia, (m) gymnotecium, (n) ascus; (o) *B. fulva* 1FP structures (asci, ascospores and *Paecilomyces* (anamorph state)). All microscopic observations were made in 400 X, except to for pictures of cleistotecium and gymnotecia (b; d; h and m) which were made 40 X.

*Paecilomyces* or *Geosmithia*. The observation of the anamorph showed phialides alone characteristically swollen at the base and gradually narrowing into a long beak. Based on this, considering the characteristic of the conidiophores borne and conidia shape, the anamorph state was identified as *Paecilomyces*.

On CYA at 25 °C, the strain of *Neosartorya fischeri* (2WAr) showed colonies (around 70 mm of diameter) col-

ored white to pale yellow (centrally) with plane and sparse surface with a floccose texture and pale underside. Similarly on MEA, colonies (75 mm of diameter) were white to pale cream showing a plane surface with a floccose texture and pale underside. Colonies on G25N (15 mm diameter) were colored white with a pale underside. At 37 °C on CYA, colonies covering the whole Petri dish were white to pale cream, sulcate, floccose and a pale underside. No

growth was observed at 5 °C. Microscopic observations showed (Fig. 2 h, i, j) clear cleistothecia, ascus and ornamented ascospores (7.5 µm long) with two longitudinal flanges. The anamorph *Aspergillus* with the head formed from phialides was alone observed.

*Eupenicillium cinnamopurpureum* (1W<sub>r</sub>) presented colonies on CYA (10-20 mm of diameter) of dense texture, sulcate, clear exudate of white color and pale cinnamon underside. Colonies on MEA (12-17 mm of diameter) were colored white with a pale underside. Colonies on G25N (12-15 mm of diameter) were colored white with a cinnamon-colored central region and a pale cinnamon underside. At 37 °C on CYA, colonies with a diameter of 5-7 mm were colored white. No growth was observed at 5 °C. Microscopic observations showed a strictly monoverticulate penicillin with ampulliform phialides (Figure 2 c) and ellipsoidal spores. It was not possible to observe cleistothecia, however, Pitt and Hocking (Pitt and Hocking, 1999) affirmed that some isolates fail to produce cleistothecia at all.

Colonies of *Eupenicillium* sp. (1BPC<sub>r</sub>) on CYA (45-50 mm of diameter), were colored white, radially sulcate, with a dense texture and pale yellow underside. Colonies on MEA (50-55 mm of diameter) were similar to CYA. On G25N (5-7 mm of diameter) colonies were colored white with a floccose texture and pale yellow underside. At 37 °C, white colonies with a diameter of 25-30 mm were radially sulcate with a pale yellow underside. No germination was observed at 5 °C. Microscopic observations showed yellow cleistothecia (Figure 2b), however it was difficult to observe ascospores.

The colonies of *Eurotium* sp. (1W<sub>Ar</sub>) on CYA (17-20 mm of diameter) were dense, sulcate with an intense yellow color and an orange underside. Both the colonies and the underside on MEA (5-10 mm of diameter) were orange. Colonies on G25N (30-35 mm of diameter) were pale yellow in color with a yellow underside. Colonies on CY20S (30-35 mm) were sulcate, with white mycelium at the margins, becoming yellow with an intense orange center and an orange underside. No growth was observed at 5 °C. Microscopically (Figure 2 d, e, f g), ascospores were not evident within 7 days, however after 14 days of incubation, ascospores (around 5 µm long) showing smooth walls with two prominent, parallel longitudinal flanges were observed. Furthermore, yellow cleistothecia, ascus and the anamorph *Aspergillus* producing only phialides was found.

*Eurotium* sp. (3W<sub>Ar</sub>) showed colonies on CYA (20-22 mm of diameter), sulcate with yellow at the margins, becoming orange brown in the central area with a very intense orange underside. Colonies on MEA (27-30 mm of diameter) were sulcate, showing an intense orange color and the same color on the underside side. No growth was observed at 5 °C or 37 °C. Colonies on CY20S showed a diameter of 20 mm after 7 days of incubation reaching 40 mm after 14 days. These colonies showed a texture plane, sul-

cate with a clear color at the margins becoming intensely orange at the center with the same color on the underside. Microscopic observations showed (after 14 days of incubation) ellipsoidal ascospores without double flanges. The anamorph *Aspergillus* with the head formed from phialides alone and also cleistothecium enveloped in yellow hyphae was observed.

*Byssochlamys fulva* (1FP) showed colonies on CYA and MEA covering the whole Petri dish with sparse, low floccose, heavy conidial production and were colored olive brown on MEA and on CYA with white filaments on CYA. The underside color was pale brown. Colonies on G25N (5-8 mm of diameter) were olive brown. No growth was observed at 5 °C. Microscopically (Figure 2 f), there were not any kind of bodies typically present in most of ascomycetes. Furthermore, ellipsoidal ascospores (5 µm long) with smooth walls and also asci (10 µm) were observed. Anamorphic reproductive structures of penicillin were observed.

#### *Alicyclobacillus* sp.

The incidence of *Alicyclobacillus* was only verified in those samples that were subjected to the detection method, so that the result was expressed based only on the presence or absence of *Alicyclobacillus* (ATSB). The strains ET and BUF were identified as *Alicyclobacillus acidoterrestris*.

## Discussions

The survey of mold and yeasts from a concentrated apple juice process showed a high contamination mainly on the steps before heat treatment (Table 1). After pre-concentration stage (110-115 °C for 30 s) the number of molds and yeast decreases from > 10<sup>7</sup> cfu/mL to < 10 cfu/mL. However, enzyme treatment stage increased their number to > 10<sup>2</sup> cfu/mL and even after pasteurization stages the number of molds and yeast was kept. However, in the end of the process, the final product decreased their count to < 10 cfu/mL, indicating that raw apple juice is not homogeneous concerning its mold and yeast contamination and the various stages of manufacture are really necessary in order to obtain a concentrate apple with no contaminant.

The mold *Penicillium* was the prevalent genera found. However, the pre-concentration step was enough to completely destroy *Penicillium* spores since they were not found in the following stages. Labuda *et al.* (2005) investigated the incidence of toxinogenic fungi in fruits and also reported *Penicillium* as the major contaminant. Other researchers observed that *P. expansum* (30-62%) was the most prevalent species on apples in storage rooms (Amiri and Bompeix, 2005). In this present study, half of the *Penicillium* strains were identified as *P. expansum* representing a concern for apple juice processing since it is the typical contaminant of apple brown rot having the ability to grow at low temperatures and to produce patulin on de-

cayed fruits (Amparo *et al.*, 2012; Doores, 1983; Jackson *et al.*, 2003; Salomão *et al.*, 2009a). During the off-season, juice companies are supplied with apples stored in packing houses. Therefore, the presence of *P. expansum* in apples should be considered a possible consequence of postharvest handling of fruits associated with the extension and conditions of storage. During the present investigation, as cited by other authors (Sydenham *et al.*, 1995), it was observed that at several times the apples are not processed as soon as received in the processing plant. This situation leads to deck storage (under no refrigerated conditions) for sometimes more than 5 days, resulting in serious implications in the level of patulin (Sydenham *et al.*, 1997). Therefore, monitoring the quality of apple lots under a rigid selection criterion during the reception step and minimizing storage at room temperature should be considered important measures to control this toxigenic mold (Fao, 2003).

The sample of wash water also showed a high contamination by molds and yeast ( $> 10^5$  cfu/mL), including *P. expansum* (Codex, 2003). In the wash step, the high-pressure water sprayed against apples helps in the removal of rotten parts which contain high levels of toxin (Acar *et al.*, 1998). However, spores will be suspended in the water causing a possible cross contamination and also increasing the risk of mold growth during bulk storage (Fao, 2003). The increase of the counts of molds and yeast observed in the washed apples ( $> 10^7$  cfu/mL), including *P. expansum*, proof that cross contamination is real and possible. Therefore is important to use a sanitizer, such as chlorine, during this step in order to reduce *P. expansum* spores (Salomão *et al.*, 2008a, 2009b). Besides, the apple sorting should be rigorous enough to remove, as far as possible, rotten fruits, even those with only small areas of rot (Codex, 2003).

*Talaromyces* sp. (1AC) was the mold isolated from the concentrated juice. This mold is a HRM that has been isolated on several occasions in screening fruit juices (Pitt and Hocking, 1999). The sequence of heat shocks applied during the process probably caused the activation of its spores from the asexual to a sexual state. The presence of *Talaromyces* species in the concentrated apple juice implies a concern because they cause spoilage in heat-treated products and members of this genus are known to produce the mycotoxin talaromycin (Enigl *et al.*, 1993). Their ascospores are extremely heat resistant and can survive 5 to 12 min of heating at 100 °C (Tournas, 1994; Pitt and Hocking, 1999).

The genus *Eupenicillium* was isolated from wash water (1Wr) and before concentration treatment (1BPCr). Another study also found this HRM from a concentrated apple juice process in a step after pasteurization (Salomão *et al.*, 2008b).

*N. fischeri* was detected in washed apples. *N. fischeri* can be introduced in the process through contaminated apples and its spores can be suspended in the water causing a possible cross contamination. Other study also related the

presence of this same HRM species in apple products (Salomão *et al.*, 2008b). *N. fischeri* ascospores showed high resistance in juices and survived to a heat treatment of 94 °C for 20 min in apple juice (Salomão *et al.*, 2004, 2007, 2008b). Some strains of *N. fischeri* are capable of producing toxins such as fumitremorgins and verruculogen (Tournas, 1994).

*Eurotium* species also were detected among the HRM and their presence in the apples was unusual. However their presence was detected in washed apples and wash water. Some species of *Eurotium* are a risk to the thermal process since they also produces ascospores which would include them to the list of resistant fungi (Splittsoesser *et al.*, 1989; Yildiz and Coksöyler, 2002). Their xerophilic characteristic is probably the major reason for their stability during dormancy (Eicher and Ludwig, 2002). The heat resistance of *Eurotium haerberiorum*, isolated from a spoilage outbreak involving grape preserves, showed  $D_{70\text{ °C}}$  of 2.5 min in 5°Brix grape juice and  $D_{71\text{ °C}}$  of 5.2 min in 65°Brix juice (Splittsoesser *et al.*, 1989).

The presence of *B. fulva* in the concentrated juice demonstrated the ability of its ascospores to remain viable after being submitted to the high temperatures applied during processing. A study demonstrated that even at a very low initial contamination, clarified apple juice can be easily spoiled by *B. fulva* (Sant'Ana *et al.*, 2010a). A strain of *B. fulva* IOC 4518, isolated from apples, survived heating at 95 °C for 20 min (Salomão *et al.*, 2008b) and its D values in apple juice at 85 °C, 90 °C, 92 °C and 95 °C were 64.5, 16.6, 6.3 and 3.1 min, respectively (Sant'Ana *et al.*, 2009). A study showed that from 16 samples of concentrate apple juice analyzed, 25% belonged to the *Byssochlamys* genera. Three strains were identified as *B. fulva* and one strain as *B. nivea* (Welke *et al.*, 2009). In addition to patulin, other metabolites have been reported to be produced by *B. fulva* such as byssochlamic acid and byssotoxin A (Tournas, 1994; Sant'Ana *et al.*, 2010b).

Although the incidence of HRM in the concentrate apple juice was low (Table 2), all heat resistant genus cited in the literature were detected in this investigation. Based on this, control measures should be studied in order to minimize the contamination by these microorganisms.

The presence of *A. acidoterrestis* in the samples is in agreement with other studies that report its incidence in a wide range of fruit juices as well as processing facilities. The contamination of fruit surfaces probably is from soil during production and harvesting (Chen *et al.*, 2006; Grande *et al.*, 2005; Murakami *et al.*, 1998). Therefore, considering that *Alicyclobacillus* are soil-borne, its control should start in the fields along with proper cleaning of fruits at the beginning of processing (Groenewald *et al.*, 2009). Groenewald *et al.* also reported the isolation of *A. acidoterrestis* from wash water and flume water, which increases the risk of possible recontamination by this bacterium through the water.

**Table 2** - Mean values of heat resistant molds (HRM) and the identified species in different stages of a concentrated apple juice process line.

Sample (code)	Mean values of HRM	Identified species of HRM (code)
Wash water (W)	1 cfu/100 mL ± 0.00	<i>Eupenicillium cinnamopurpureum</i> (1Wr)
Washed apples (WA)	3 cfu/100 g ± 1.41	<i>Eurotium</i> sp. (1WAr) <i>Neosartorya fischeri</i> (2WAr) <i>Eurotium</i> sp. (3WAr)
Before pre-concentration (BPC)	1 cfu/100 mL ± 0.70	<i>Eupenicillium</i> sp. (1BPCr)
Final product (FP)	2 cfu/100 mL ± 1.41	<i>Byssoschlamys fulva</i> (1FPr)

In this present survey, *A. acidoterrestris* was detected in the step of enzymatic treatment and step before ultra filtration. These facts showed the high resistance of its spores in remaining viable even after submitted to the heat treatments applied in the pre-concentration stage (110-115 °C for 30 s) and in the first pasteurization stage (85-90 °C for 30 s). The D values for *A. acidoterrestris* in apple juice at 85, 90 and 95 °C were 56, 23 and 2.8 min, respectively (Splitstoesser *et al.*, 1994). Since *A. acidoterrestris* is a thermoacidophilic and spore-forming bacterium they can survive in acid media (such as apple juice) and grow at temperatures higher than 20 °C, thereby having the potential to spoil the shelf stable products during storage (Chen *et al.*, 2006; Chmal-Fudali and Papiewska, 2011). Although *A. acidoterrestris* is not pathogenic, it is recognized as a problem in the juice industries worldwide since spore-related juice spoilage may result in bad smelling compounds (Jensen and Whitfield, 2002; Splitstoesser *et al.*, 1998; Walls and Chuyate, 2000).

The presence of *P. expansum* in the raw material represents a likely risk of patulin in the final product, since this toxin shows relatively high heat stability in an acidic environment. The detection of *B. fulva* and *A. acidoterrestris* CCT 7548 in the stages after heat treatment demonstrated their high heat resistance and also the possibility of toxins and off-flavors remaining in the apple juice. Besides, it was demonstrated that *Byssoschlamys* spp. are able to produce patulin in apple juice packages during storage and distribution (Sant'Ana *et al.*, 2010b). Therefore, this research concluded that *P. expansum*, *B. fulva* and *A. acidoterrestris* are important biologic hazards for apple juice products. Furthermore, the application of more severe thermal treatments to inactivate resistant spores and toxins is impracticable for fruit juices. So, surveys should find new strategies to reduce the target microorganisms in raw material, controlling the initial contamination of fruits.

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