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# Effect of Bacterial Strains Isolated from Stored Shiitake (*Lentinula edodes*) on Mushroom Biodeterioration and Mycelial Growth

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**Abstract:** Shiitake (*Lentinula edodes*) is a fungus that attracts the attention of consumers and researchers due to its flavour and bioactive compounds with biological activities. However, it swiftly loses quality because of its short shelf life. The present work evaluated the shelf life of cultivated shiitake carpophores with a weekly monitoring of the physico-chemical, microbiological, and sensory parameters under three different storage conditions. To establish the mycopathogenicity of fruitbody-inhabiting bacterial microbiota, 48 strains were isolated from shiitake throughout its storage time and subsequently inoculated on cap cuticles and slices. Additionally, an antibiogram of these bacterial strains against shiitake mycelium was performed. Shiitake had a total microbial load of 5 log cfu/g, composed mainly of *Pseudomonas* genus, followed by *Enterobacteriaceae* family, and moulds and yeasts. Among the 48 bacterial strains identified, 18 belonged to the *Enterobacteriaceae* family and 21 to the *Pseudomonas* genus. The biodeterioration capacity of the isolated strains increased throughout the conservation period, with slices being more damaged than cap cuticles. We observed a mycopathogenic behaviour of *Burkholderia* sp., *Ewingella americana*, *Serratia* sp., *Microbacterium* sp. and *Rahnella* sp. strains, although only the latter highly inhibited shiitake vegetative mycelium growth.

**Keywords:** shiitake; shelf life; modified atmosphere; antibiogram; mycopathogen; bacteria; *Rahnella*

## 1. Introduction

Shiitake (*Lentinula edodes* (Berk.) Pegler) mushrooms are currently attracting the attention of consumers for their high nutrient value, gastronomic interest, and human health properties [1]. In the last decade, shiitake cultivation has increased exponentially, achieving the second position in the global mushroom industry after *Agaricus bisporus* (J.E. Lange) Imbach. Mushrooms have a short shelf life due to factors such as their high respiration and metabolic rates, rapid weight loss, and high microbiological load, which all together induce quality losses shortly after harvesting [2]. The distribution chain of this fresh product is highly conditioned by its short shelf life. In order to guarantee its sanitary aptitude, a combination of decontamination and preserving technologies—such as irradiation combined with modified atmosphere packaging—has been developed [3]. The use of ionisation technologies combined with refrigerated storage has also been reported to extend the shelf life of different mushrooms, including shiitake [2]. The modified atmosphere limits gas exchanges with the outside air, thus altering the atmosphere composition inside the package and decreasing the

respiration rate of the product. Optimum conditions inside the package can be passively achieved as a result of mushroom respiration [4].

Bacteria, moulds, enzymatic activity and biochemical changes can cause physical spoilage of mushrooms during storage. The *Pseudomonas* genus is a very competitive microbial group in the rhizosphere which takes part symbiotically in the mushroom fruiting body development [5]. However, some species of this bacterial genus also appear to be responsible for mushroom postharvest deterioration, such as *Pseudomonas tolaasii* Paine, which is accountable for the surface browning produced by the action of the tolaasin exotoxin. Yeasts such as *Candida sake* Uden and H.R. Buckley are also able to provoke post-harvest damages [6]. *Ewingella americana* Grimont et al. (*Entobacteriaceae* family) is associated with a browning disorder of the *A. bisporus* stipe, called internal stipe necrosis, which can also occur in other mushroom species [7]. *Burkholderia gladioli* pv. *agaricicola* Yabuuchi is an important pathogen in the mushroom industry, causing soft rotting symptoms over several important commercial mushrooms such as *L. edodes*, *Pleurotus ostreatus* (Jacq.) P. Kumm., *Flammulina velutipes* (Curtis) Singer, *Pholiota nameko* (T. Itô) S. Ito and S. Imai or *Hypsizygus marmoreus* (Peck) H.E. Bigelow [8].

In this work, several physico-chemical, microbiological and sensory parameters of shiitake mushrooms were monitored under three storage conditions until carpophores were completely degraded. Throughout the conservation period, different bacterial strains were isolated, selected and identified from all batches to detect possible changes in the predominant microbiota. These strains were subsequently inoculated on the cuticle, slices and mycelium of *L. edodes* mushroom caps, in order to identify which ones were primarily responsible for bacterial biodeterioration.

## 2. Materials and Methods

### 2.1. Samples and Storage Conditions

Shiitake mushrooms were purchased in Mercazaragoza food market (Zaragoza, Spain). Samples were transported to the laboratory and stored at 4 °C and 95% of relative humidity. For the experiment, mushrooms were stored under three conditions: (1) room temperature (22 °C) as a control (C), (2) refrigerated at 4 °C (R), and (3) packaged in passive modified atmosphere under refrigeration at 4 °C (MAP). For all batches, semi-rigid polypropylene trays (TS250, Linpac) of 250 mL of capacity were used for packaging 6–7 mushrooms per tray (100 g approximately). C and R trays were wrapped with absorbent paper to retain moisture excess. The trays of the MAP treatment were heat-sealed using a manual packaging machine BOV 160 (ORA Constructeur, Trévoux, France) with a microperforated film composed of a double layer of polyethylene (40 µm) and polyester (12 µm), and having two microperforations (50 × 90 µm) per tray (PPLUS, Amcor Flexible, Madrid, Spain). A total of 15 trays per batch were prepared, so that three trays could be analysed in each sampling day.

### 2.2. Shelf-Life Assay

#### 2.2.1. Modified Atmosphere Gas Measurement

The modification of the atmosphere composition inside heat-sealed trays was monitored using a respirometer CheckMate II (PBI Danasensor, Barcelona, Spain). O<sub>2</sub> and CO<sub>2</sub> concentrations were measured in each tray immediately after packaging, 24 and 48 h later, and then every seven days. The value presented for each sampling day is an average of three different trays.

#### 2.2.2. Determination of Weight Loss

The weight of all the trays was measured immediately after packaging, and after that every seven days, with a Sartorius 3716 scale (Sartorius, Göttingen, Germany). The value presented for each batch on each sampling day is an average of three different trays. Weight loss is expressed as a percentage of the initial weight.

### 2.2.3. Microbial Analysis

Microbial analyses included the quantification of aerobic mesophilic microorganisms (AMM; which constitute the total aerobic microbial load), *Pseudomonas* genus, *Enterobacteriaceae* family, lactic acid bacteria (LAB), and moulds and yeasts. The quantifications were done following the corresponding ISO Norm, with slight modifications as indicated in Rivera et al. [6]. The batches were analysed once a week until shiitake was completely degraded. Each sample (25 g) was serially diluted in sterile distilled peptone water 0.1% (Merck, Darmstadt, Germany), and homogenised using a laboratory blender Stomacher 400 Circulator (Seward Laboratory, London, England) during 2 min at 250 rpm according to ISO Norm 6887-1:2017. In order to facilitate the subsequent selection and picking of the microbial colonies on the basis of colony morphology, quantifications were carried out in PCA agar plates by spread plating of 0.1 mL. Culture media and incubation conditions for each microbial group were: (1) AMM: plate Count Agar (PCA) (Merck) during 72 h at  $30 \pm 1$  °C; (2) *Pseudomonas* genus: *Pseudomonas* agar base (Oxoid, Basingstoke, Hampshire, U.K.) supplemented with cephaloridine-fucidin-cetrimide (CFC) during 48 h at  $25 \pm 1$  °C; (3) *Enterobacteriaceae* family: violet red bile glucose (VRBG) (Oxoid) during 24 h at  $30 \pm 1$  °C; (4) LAB: Man, Rogosa and Sharpe agar (MRS) (Merck) during 72 h at  $30 \pm 1$  °C (Oxoid), using anaerobic jars with an atmosphere generation system; and (5) moulds and yeasts: dichloran rose-bengal chloramphenicol agar (DRBC) (Merck), supplemented with 0.1% gentamicin (Carrier, Barcelona, Spain) to avoid *Pseudomonas* spp. growth, during 4 days at  $25 \pm 1$  °C. The value presented for the microbial count of each sampling day is the average of three measurements (one per tray, three trays per sampling day and batch), expressed as log cfu/g.

### 2.2.4. Sensory Analysis

A panel of eight trained tasters evaluated the quality and characteristics of shiitake mushrooms from each storage condition each seven days from day 0 to day 28 of storage. Tasters were previously trained for three sessions of 45 min. The analyses were conducted according to the ISO 11035:1994 and the following sensory parameters were assessed: external aspect, hymenium appearance, texture, aroma, taste and general acceptability. Each parameter was assessed with a nine-point rating scale. The values presented for each sampling day and batch are the average of three measurements (one per tray, three trays per sampling day and batch). The end of shiitake shelf life was established when the score of either of the attributes dropped below 6.0.

### 2.3. Strains Isolation and Identification

The PCA agar plates of the microbiological analysis were used for strain isolation and identification. For each combination of sampling day and batch, we picked six colonies on the basis of their morphology and relative frequency, using a stereomicroscope Leica EZ4 (Wetzlar, Germany). Then, culture plates were incubated for five days—longer than in the counting protocol—to more easily detect morphology differences. DNA isolation was performed following Barghouthi (2011) [9] and using PrepMan<sup>®</sup> Ultra Sample Preparation Reagent (Applied Biosystems, Foster City, CA, USA) as extract agent. Universal primers employed for amplifying V1-V9 region of the *16S rRNA* gene sequence were: 8F (5'-AGAGTTTGATCCTGGCTCAG-3') and 1492R (5'-CGGTTACCTTGTTACGACTT-3'). The amplification reaction was prepared for a 25 µL final volume, containing: 12 µL of sterile double distilled water, 1 µL of each primer, 1 µL of BSA (bovine serum albumin), 2.5 µL of Taq free DNA polymerase (Invitrogen, Carlsbad, CA, USA), 5 µL of PCR reaction buffer including dNTP and MgCl<sub>2</sub> (Invitrogen), and 2.5 µL of template DNA. PCR was performed on MyCycler Thermal Cycler (Bio-Rad, Hercules, CA, USA). The following amplification program was used: 94 °C for 2 min, 30 cycles of 94 °C 30 s, 51 °C 60 s, 72 °C 1 min, and 7 min at 72 °C. Samples were kept at 4 °C until revealing by electrophoresis. 1.5% agarose gel was performed with 30 mL of TAE buffer (Buffer Tris, Acetic Acid, EDTA) and 0.8 µL of Midori Green (NIPPON Genetics GmbH, Düren, Germany); 60 mV of current was used from an electrophoresis source BioRad PowerPac HV (BioRad, Hercules, CA, USA).

Band revelation was carried out in a transilluminator (Chemidoc XRS + BioRad, USA) with GeneSys software (Syngene, Cambridge, UK). Amplicons were sequenced by an external company (STAB VIDA, Caprica, Portugal). Sequences were then checked for quality using 4Peaks (Nucleobytes, Amsterdam, The Netherlands) and identified by highest similarity using BLAST algorithm in Genbank.

#### 2.4. Pathogenicity Test

The strains isolated for identification (Table 3) were revived in Tryptone Soy Broth (TSB) (Sigma Aldrich, Saint Louis, MO, USA). After 24 h of incubation at 30 °C, 200 µL of each isolated strain was inoculated on slices of *L. edodes* and 70 µL on mushroom caps of *L. edodes*. Before the inoculation, the concentration was measured and adjusted to 10<sup>8</sup> cfu/mL according to McFarland scale. For comparison purposes, cultivated *A. bisporus* slices were also included in this test. Mushroom slices were placed in Petri dishes and mushroom caps in plastic trays covered with film to prevent dehydration. Both containers were incubated at 10 °C for two weeks, until damages around the inoculation spots were general. Distilled water was used as a negative control. Biodeterioration was evaluated using a four-level rating scale: 0—no damage; 1—damage in the inoculation area; 2—widespread damage in the inoculation area; 3—severe damage to the carpophore (Supplementary Materials Figure S1).

#### 2.5. Antibioassay

Mycelial shiitake agar disks of 4 mm of diameter were removed using a sterile cork borer, aseptically transferred into PDA agar (Merk) and incubated for 24 h at 25 °C. Then, the bacterial strains (the same ones used for the pathogenicity test) were inoculated with a sterile loop, approximately 2.5 cm away from shiitake mycelium, and further incubated for 2 weeks at 25 °C. Sterile distilled water was used as a negative control. Three inoculations per strain were prepared. The effect was visually evaluated.

#### 2.6. Statistical Analysis

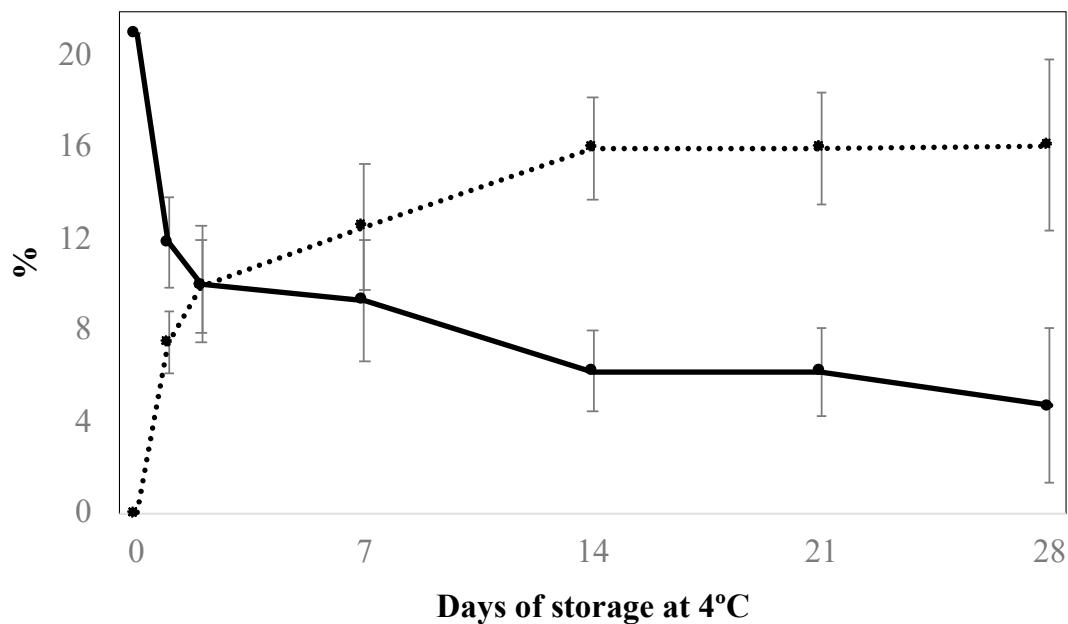
Data were analysed through the *t*-student test and one-way analysis of variance (ANOVA) using SPSS (Windows 2006; Statistical Analysis). It was checked that the models met the assumptions of normal distribution and constant variance. The different treatments and sampling times were compared using post-hoc Tukey's test with a significance level of  $p < 0.05$ . The pathogenicity test was analysed through ordinal logistic regression.

### 3. Results and Discussion

#### 3.1. Modification of Package Atmosphere Composition

During the first 48 h, the CO<sub>2</sub> and O<sub>2</sub> gas concentrations reached values of 10%. The equilibrium was reached after two weeks, with CO<sub>2</sub> and O<sub>2</sub> concentrations of 16% and 7%, respectively (Figure 1). An excessive accumulation of CO<sub>2</sub> could cause physiological lesions to fungi, producing severe browning, or stimulating the proliferation of facultative anaerobic microorganisms. The lack of oxygen could lead to an anoxia situation, accompanied by strange odours due to the production of volatile substances such as ethanol and acetaldehyde [10]. Shiitake has a higher respiratory rate and a greater susceptibility to elevated concentrations of CO<sub>2</sub> than other fungal species [2]. In this respect, previous studies compared MAP packaging with low-density polyethylene (PE) and polypropylene (PP) films [2]. In both cases, O<sub>2</sub> concentration decreased and CO<sub>2</sub> increased rapidly during the first days of storage. However, after that, the CO<sub>2</sub> concentration remained constant at very different levels, 13.6% for PP and 9% for PE. Our results show similar dynamics to those previously obtained with PP films, although with slightly higher levels of CO<sub>2</sub>, likely because of the plastic film characteristics (permeability, size and number of perforations), or of the intra-specific differences among *L. edodes*

strains, as other authors reported [2]. Moreover, our results agree with those obtained for hypogeous mushrooms when using similar storage conditions [4,6].



**Figure 1.** Time trend of CO<sub>2</sub> (--- dashed line) and O<sub>2</sub> (— continuous line) of shiitake mushrooms packaged under modified atmosphere stored at 4 °C. Data expressed as mean ± standard deviation (SD) of three samples.

### 3.2. Time Trend of Weight

The control batch suffered a weight loss of almost 80% after one week of storage (Table 1). This was probably caused by a quick dehydration related to the respiratory activity of *L. edodes* carpophores, which is very high at room temperature [4,6]. Weight loss was reduced to 25% when refrigeration was applied (Table 1). By incorporating MAP packaging, the weight only dropped by 2% during the first week (Table 1). At the end of the storage time, the R batch lost over 80% of the initial weight, whereas in the MAP batch the weight loss did not exceed 5% (Table 1). Previous studies also obtained lower weight losses with microperforated than with macroperforated films (5% and 47%, respectively, after 16 days of storage under refrigeration) [2].

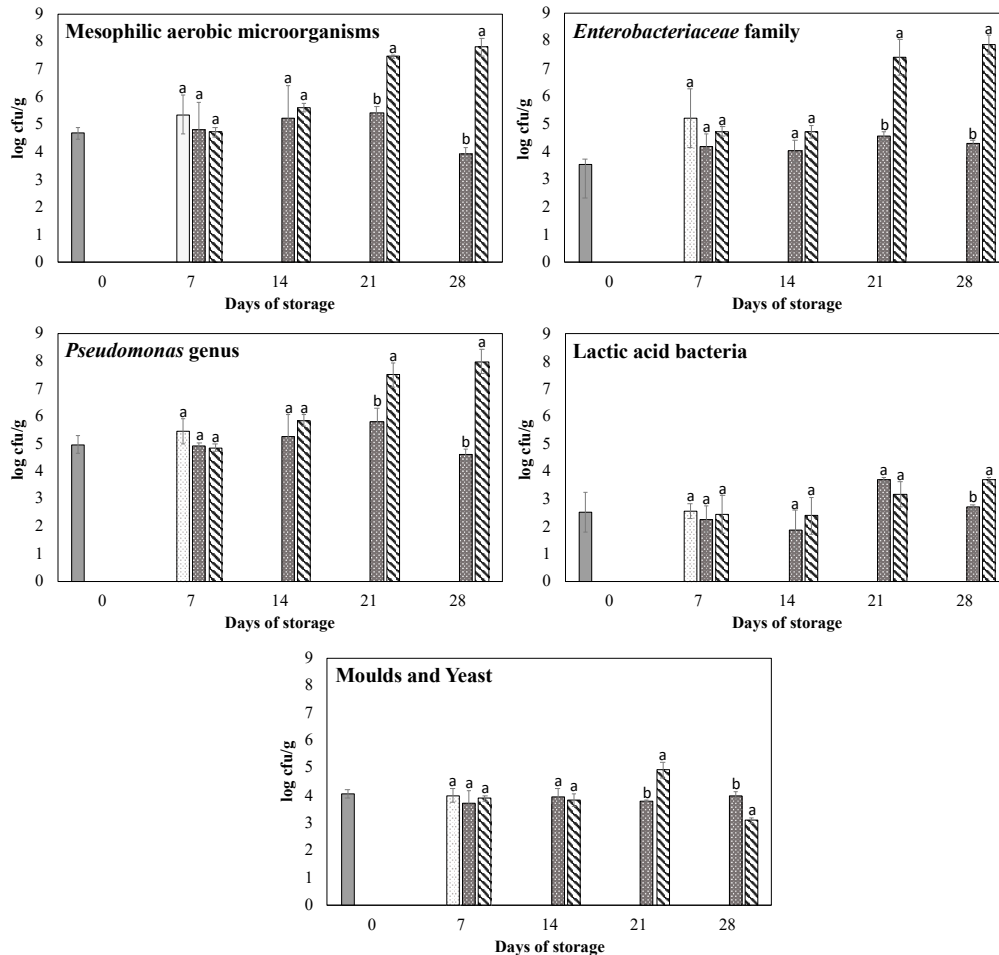
**Table 1.** Weight loss (%) in *L. edodes* under different storage conditions: room temperature (C), refrigerated at 4 °C (R), and modified atmosphere packaging at 4 °C (MAP). Data expressed as mean ± standard deviation (SD) of three samples. Different letters within the same column indicate statistically significant differences at  $p < 0.05$ .

Day	C	R	MAP
0	0.0 ± 0.0 <sup>b</sup>	0.0 ± 0.0 <sup>d</sup>	0.0 ± 0.0 <sup>c</sup>
7	78.1 ± 5.9 <sup>a</sup>	24.7 ± 4.6 <sup>c</sup>	2.3 ± 0.6 <sup>b</sup>
14	-	36.4 ± 4.0 <sup>bc</sup>	2.5 ± 0.4 <sup>b</sup>
21	-	50.5 ± 5.1 <sup>b</sup>	3.0 ± 0.1 <sup>b</sup>
28	-	84.2 ± 5.7 <sup>a</sup>	4.9 ± 0.7 <sup>a</sup>

### 3.3. Time Trend of Microbial Populations

Initial total AMM load was 5 log cfu/g. Genus *Pseudomonas* was the most frequently isolated group (4.7 log cfu/g), followed by moulds and yeasts (4.1 log cfu/g) and *Enterobacteriaceae* family (3.5 log cfu/g). Lactic acid bacteria showed much lower values (2.5 log cfu/g) (Figure 2). Venturini et al. obtained

similar microbial loads for shiitake [11], whereas Jiang et al. showed lower counts for AMM than for *Pseudomonas* [12]. In comparison with other mushroom species, the microbial load of *L. edodes* is lower than those reported in *A. bisporus* (7.5 and 7.2 log cfu/g of AMM and *Pseudomonas*, respectively), possibly due to the antimicrobial potential of this species [11].



**Figure 2.** Time trend of mesophilic aerobic microorganisms, *Pseudomonas* genus, *Enterobacteriaceae* family, lactic acid bacteria and yeasts populations for fresh *L. edodes* stored under different conditions: ■ initial counts, □ room temperature, ▒ refrigerated at 4 °C, and ▨ modified atmosphere packaging at 4 °C. Data expressed as mean ± standard deviation (SD) of three samples. <sup>a, b</sup> Different superscript letters within the same day of storage indicate statistically significant differences at  $p < 0.05$ .

The microbial load increased slightly during storage for all groups, although there were no significant differences until the third week (Figure 2). The C batch was unacceptable after one week due to the great loss of moisture and the resulting changes in organoleptic characteristics (see below). On day 28 of storage, the R batch suffered a slight decrease in AMM, *Pseudomonas*, and LAB contents of almost 1 u.log, which was likely related to dehydration. In contrast, the MAP treatment increased all microbial groups counts on day 28, except for moulds and yeasts. The decrease in moulds count could be explained by the fungicide effect of the high CO<sub>2</sub> concentration (16%), and in the case of yeasts by the longer generational time than bacteria and the increased nutrient competition exerted by bacteria. The CO<sub>2</sub> had no apparent bactericidal effect on *Pseudomonas*, even though this group is strictly aerobic. This could be explained by the high humidity inside the package (and therefore on the surface of the carpophore) facilitating microbial proliferation. It is remarkable that on day 28 MAP differed to R in more than 3 u.log of AMM, *Pseudomonas* and *Enterobacteriaceae*. Finally, although the high levels of CO<sub>2</sub> facilitated LAB growth, there were no significant differences among batches until day 28 of storage.



Jiang et al. applied MAP to shiitake, and reported similar counts of AMM (7.3 log cfu/g), and higher counts of *Pseudomonas* (8.7 log cfu/g), and moulds and yeasts (6.8 log cfu/g) after 20 days of conservation [12]. The MAP of the hypogeous fungi *Tuber melanosporum* Vittad. and *Tuber aestivum* Vittad. at 4 °C barely increased AMM load by 2 u.log after four weeks of storage [6]. This difference might be related to the lower initial microbial load of shiitake, 2 log cfu/g lower than those found in *Tuber* genus, and by the greater availability of nutrients on shiitake surface. These factors could explain that shiitake microbial populations have higher possibilities of proliferation under propitious conditions of high humidity.

### 3.4. Sensory Evaluation and Shelf Life

Externally, the deterioration of shiitake mushrooms was characterised by the presence of superficial white spots, dehydration and darkening of the carpophores. After one week, the C batch showed lower values than R and MAP for all the sensory parameters tested (Table 2). In the C batch, the shelf life was established in less than one week, due to moisture loss and subsequent changes in external and hymenium appearance, with values of 5.8 and 4.7 on day 7, respectively (Table 2). In the R and MAP batches, the shelf life increased by two weeks. The R batch was considered unacceptable on day 21 due to texture changes, value of 5.5, whereas the MAP batch was rejected due to the darkening of the carpophores and taste changes, with values of 5.5 and 4.7, respectively, and the increase of the microbial load (i.e., the visualisation of superficial microbial growth) (Table 2). The use of MAP prevented moisture losses; however, this allowed the microbial proliferation on the mushroom surface during storage. Thus, MAP did not seem to contribute to increase the shelf life, although it significantly reduced dehydration and preserved texture. The observations on day 28 confirmed those of previous samplings.

**Table 2.** Time trend of the sensory parameters of *L. edodes* under different storage conditions: room temperature (C), refrigerated at 4 °C (R), and modified atmosphere packaging at 4 °C (MAP). Each of the sensory attributes was scored on a 1–9 scale (1 = definitely unacceptable, 3 = unacceptable, 5 = may be acceptable, 7 = acceptable, and 9 = definitely acceptable). Data expressed as mean ± standard deviation (SD) of three samples.

Day	Storage Condition	External Appearance	Hymenium Appearance	Texture	Aroma	Taste	General Acceptability
0	C	9.0 ± 0.0 <sup>a</sup>	9.0 ± 0.0 <sup>a</sup>	9.0 ± 0.0 <sup>a</sup>	9.0 ± 0.0 <sup>a</sup>	9.0 ± 0.0 <sup>a</sup>	9.0 ± 0.0 <sup>a</sup>
	R	9.0 ± 0.0 <sup>a</sup>	9.0 ± 0.0 <sup>a</sup>	9.0 ± 0.0 <sup>a</sup>	9.0 ± 0.0 <sup>a</sup>	9.0 ± 0.0 <sup>a</sup>	9.0 ± 0.0 <sup>a</sup>
	MAP	9.0 ± 0.0 <sup>a</sup>	9.0 ± 0.0 <sup>a</sup>	9.0 ± 0.0 <sup>a</sup>	9.0 ± 0.0 <sup>a</sup>	9.0 ± 0.0 <sup>a</sup>	9.0 ± 0.0 <sup>a</sup>
7	C	5.8 ± 1.5 <sup>b</sup>	4.7 ± 0.9 <sup>b</sup>	6.5 ± 0.8 <sup>b</sup>	7.1 ± 0.5 <sup>b</sup>	6.8 ± 0.8 <sup>b</sup>	5.8 ± 1.2 <sup>b</sup>
	R	8.4 ± 0.5 <sup>a</sup>	8.2 ± 0.3 <sup>a</sup>	8.6 ± 0.2 <sup>a</sup>	8.5 ± 0.1 <sup>a</sup>	8.5 ± 0.2 <sup>a</sup>	8.3 ± 0.5 <sup>a</sup>
	MAP	8.3 ± 0.7 <sup>a</sup>	8.6 ± 0.3 <sup>a</sup>	8.8 ± 0.1 <sup>a</sup>	8.7 ± 0.2 <sup>a</sup>	8.6 ± 0.3 <sup>a</sup>	8.5 ± 0.4 <sup>a</sup>
14	C	1.7 ± 0.9 <sup>b</sup>	1.1 ± 0.3 <sup>b</sup>	3.0 ± 0.5 <sup>b</sup>	4.5 ± 0.5 <sup>b</sup>	-	2.1 ± 0.5 <sup>b</sup>
	R	7.5 ± 0.7 <sup>a</sup>	7.9 ± 0.5 <sup>a</sup>	7.2 ± 0.5 <sup>a</sup>	7.3 ± 0.4 <sup>a</sup>	7.7 ± 0.6 <sup>a</sup>	7.5 ± 0.4 <sup>a</sup>
	MAP	7.1 ± 0.9 <sup>a</sup>	7.7 ± 0.6 <sup>a</sup>	8.1 ± 0.4 <sup>a</sup>	7.9 ± 0.6 <sup>a</sup>	7.5 ± 0.7 <sup>a</sup>	7.0 ± 0.6 <sup>a</sup>
21	C	-	-	-	-	-	-
	R	7.0 ± 1.1 <sup>a</sup>	7.1 ± 0.8 <sup>a</sup>	5.5 ± 0.4 <sup>b</sup>	6.6 ± 0.8 <sup>a</sup>	6.6 ± 0.8 <sup>a</sup>	6.5 ± 0.8 <sup>a</sup>
	MAP	5.5 ± 0.7 <sup>a</sup>	5.8 ± 0.9 <sup>a</sup>	7.2 ± 0.8 <sup>a</sup>	5.7 ± 1.1 <sup>a</sup>	4.9 ± 0.3 <sup>b</sup>	5.0 ± 1.1 <sup>b</sup>
28	C	-	-	-	-	-	-
	R	5.4 ± 0.8 <sup>a</sup>	5.1 ± 0.3 <sup>a</sup>	3.8 ± 0.7 <sup>a</sup>	5.9 ± 0.4 <sup>a</sup>	6.0 ± 1.2	5.1 ± 1.0 <sup>a</sup>
	MAP	1.5 ± 1.1 <sup>b</sup>	2.1 ± 0.3 <sup>b</sup>	5.1 ± 0.4 <sup>a</sup>	3.1 ± 0.7 <sup>b</sup>	-	1.3 ± 0.4 <sup>b</sup>

<sup>a, b</sup> Different letters denote significant differences ( $p \leq 0.05$ ) between storage treatments for the same day.

### 3.5. Identification of Bacterial Isolates

A total of 48 different strains were selected on the basis of colony morphology in PCA agar. The bacterial sequences were deposited in GenBank under accession numbers from MT335639 to MT335686 (Supplementary Materials Table S1). Almost all the strains isolated throughout the shelf

life of shiitake were Gram-negative, specifically gamma proteobacteria belonging to the *Pseudomonas* genus and the *Enterobacteriaceae* family (Table 3). Within this last group of facultative aero-anaerobic bacteria, *E. americana* predominated with 18 strains. In addition, 21 different strains of *Pseudomonas* were found. Some of these bacteria are responsible for the degradation of the carpophores of various fungal species, due to the toxins they produce [13] or to their pathogenic nature [7]. Aslani et al. [14] isolated 18 Gram-negative strains from mushrooms with symptoms similar to those of brown blotch disease, and 66 strains from different species of wild mushrooms, in which 58% were Gram-negative and 42% Gram-positive. These authors identified *Pseudomonas* and *Serratia* strains, but also *Bacillus*, *Stenotrophomonas* and *Brochothrix* strains.

**Table 3.** List of the strains isolated during the microbiological analysis, Genbank accession numbers, and results of the biodeterioration evaluation with a pathogenicity test. The genus corresponds to the best BLAST hit in Genbank database for each bacterial strain, as detailed in Table S1. The pathogenicity test was performed on mushroom slices of *L. edodes* and *A. bisporus*, and on cap cuticle of *L. edodes*. The biodeterioration was scored on a 0–3 scale (0: no appreciable damage, 1: slight damage in the inoculation area, 2: extended damage around the inoculation area, 3: severe damage throughout the carpophore sample). Data are disaggregated by the day of storage in which the strain was retrieved (D0–D28) and by the batch from which the strain was retrieved (C: room temperature; R: refrigerated at 4 °C; MAP: modified atmosphere packaging at 4 °C).

Day-Storage	Accession Number	Genus	<i>L. edodes</i>		<i>A. bisporus</i>
			Slices	Cap	Slices
D0	MT335659	<i>Microbacterium</i>	3	3	0
D0	MT335657	<i>Paenibacillus</i>	0	2	2
D0	MT335639	<i>Ewingella</i>	1	1	0
D0	MT335646	<i>Pseudomonas</i>	0	1	1
D0	MT335652	<i>Pseudomonas</i>	0	1	0
D0	MT335647	<i>Pseudomonas</i>	0	1	1
D7-C	MT335641	<i>Ewingella</i>	0	2	2
D7-C	MT335650	<i>Pseudomonas</i>	1	1	1
D7-C	MT335667	<i>Pseudomonas</i>	3	1	0
D7-C	MT335653	<i>Pseudomonas</i>	3	1	2
D7-C	MT335673	<i>Pseudomonas</i>	0	1	2
D7-C	MT335654	<i>Pseudomonas</i>	3	1	3
D7-R	MT335642	<i>Ewingella</i>	1	2	3
D7-R	MT335644	<i>Ewingella</i>	2	1	1
D7-R	MT335648	<i>Ewingella</i>	3	1	1
D7-R	MT335656	<i>Ewingella</i>	3	1	2
D7-R	MT335683	<i>Pseudomonas</i>	3	1	1
D7-R	MT335651	<i>Pseudomonas</i>	1	0	0
D7-MAP	MT335643	<i>Micrococcus</i>	0	2	2
D7-MAP	MT335658	<i>Ewingella</i>	3	1	1
D7-MAP	MT335660	<i>Ewingella</i>	0	2	3
D7-MAP	MT335679	<i>Pseudomonas</i>	1	1	1
D7-MAP	MT335645	<i>Pseudomonas</i>	3	2	1
D7-MAP	MT335661	<i>Serratia</i>	3	2	1
D14-R	MT335662	<i>Ewingella</i>	3	2	1
D14-R	MT335666	<i>Ewingella</i>	3	3	1
D14-R	MT335665	<i>Ewingella</i>	3	2	0
D14-R	MT335684	<i>Pseudomonas</i>	1	2	2
D14-R	MT335664	<i>Pseudomonas</i>	3	1	2
D14-R	MT335669	<i>Serratia</i>	3	0	2
D14-MAP	MT335649	<i>Burkholderia</i>	3	2	3
D14-MAP	MT335663	<i>Ewingella</i>	3	2	2
D14-MAP	MT335671	<i>Ewingella</i>	3	1	3
D14-MAP	MT335675	<i>Pseudomonas</i>	3	2	2
D14-MAP	MT335640	<i>Pseudomonas</i>	3	2	2
D14-MAP	MT335676	<i>Pseudomonas</i>	1	2	1



Table 3. Cont.

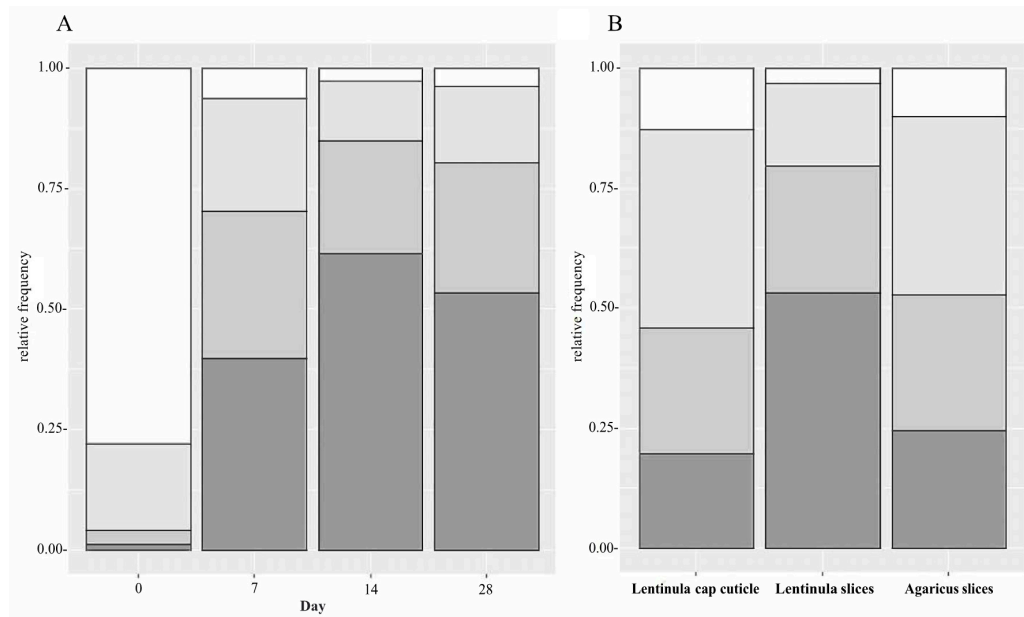
Day-Storage	Accession Number	Genus	<i>L. edodes</i>		<i>A. bisporus</i>
			Slices	Cap	Slices
D28-R	MT335686	<i>Ewingella</i>	3	1	3
D28-R	MT335674	<i>Ewingella</i>	3	3	2
D28-R	MT335680	<i>Ewingella</i>	3	2	3
D28-R	MT335685	<i>Ewingella</i>	3	2	2
D28-R	MT335670	<i>Pseudomonas</i>	3	2	1
D28-R	MT335678	<i>Pseudomonas</i>	3	1	1
D28-MAP	MT335682	<i>Paenarthrobacter</i>	1	1	3
D28-MAP	MT335672	<i>Ewingella</i>	3	1	3
D28-MAP	MT335668	<i>Ewingella</i>	3	1	3
D28-MAP	MT335677	<i>Pseudomonas</i>	3	1	3
D28-MAP	MT335681	<i>Pseudomonas</i>	0	1	2
D28-MAP	MT335655	<i>Rahnella</i>	3	2	3

### 3.6. Biodeterioration Capacity of the Isolated Strains

Almost all the strains of *E. americana* and *Pseudomonas* used in the pathogenicity test caused severe damages on shiitake slices (Table 3). *Microbacterium* sp., *Burkholderia* sp., *Serratia* sp. and *Rahnella* sp. strains also produced severe damages scoring three in damage scale. In contrast, one strain of *Pseudomonas* and one of *Serratia* were the only ones that did not apparently damage shiitake caps (score 0). The slices of *A. bisporus* were not damaged by *Microbacterium* sp., two strains of *E. americana* and three of *Pseudomonas* (Table 3).

Among the isolated species, *P. tolaasii*, *E. americana* and some species of the *Burkholderia* genus are usually the cause of different mushroom diseases [5–7]. The reservoir of *E. americana* is unknown, but it has been isolated as an opportunistic pathogen from carrots [15] and vacuum-packed meat [16]. *Ewingella americana* has been reported as a mycopathogenic agent causing a brown lesion and necrosis in mushroom stems [7]. *Burkholderia gladioli* is an important pathogen causing the softening and deterioration of various commercial mushrooms, such as *L. edodes*, *A. bisporus* and *P. ostreatus* [8]. However, as far as we know, this is the first time that *Rahnella* sp. has been isolated from mushroom carpophores.

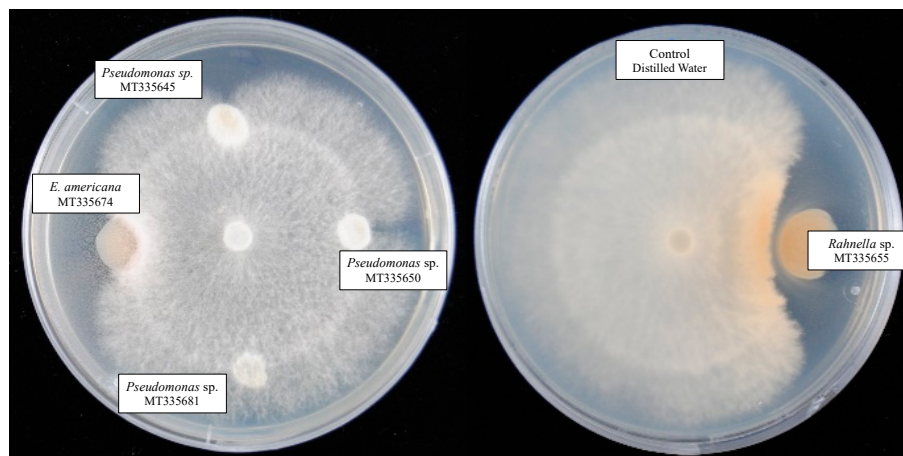
The ability of the strains isolated on days 7 to 28 to deteriorate *L. edodes* caps was significantly higher than that of the strains isolated on day 0 (z-ratio = −5.7, −8.7 and −7.7, with *p*-value < 0.001 in all cases) (Figure 3A). The slices of *L. edodes* were significantly more damaged than its cap cuticle (z-ratio = −3.6, *p*-value < 0.001) (Figure 3B). This clearly indicates that the cuticle serves as a barrier against microbial deterioration [14]. The isolated bacterial strains produced a higher damage level on shiitake slices than on *A. bisporus* slices (z-ratio = 2.9, *p*-value = 0.011) (Figure 3B). This could be due to differences in the enzymatic repertoire of the various bacterial strains or to bacterial strains having higher deterioration ability on the mushroom species from which they are isolated [5,8,14]. Regarding the latter hypothesis, it would be interesting to test whether bacterial strains develop pathogenic mechanisms specific to the mushroom in which they grow.



**Figure 3.** Predicted damage severity caused to *L. edodes* tissues by the microbial strains inoculated in the pathogenicity test, according to the day of storage in which the strains were isolated (A). Mean damage levels provoked by the inoculated microbial strains (all time points pooled) on *L. edodes* and *A. bisporus* tissues (B): □ 0: no appreciable damage, ◻ 1: slight damage in the inoculation area, ◻ 2: extended damage in the inoculation area, ◼ 3: severe damage throughout the carpophore sample.

### 3.7. Inhibition of Mycelium Growth

In the antibiogram test, most of the inoculated strains did not affect mycelial growth, although some strains showed a slight inhibition around the colony, preventing *L. edodes* mycelium from growing above them. Only *Rahnella* sp. had a clear effect on the mycelium of shiitake, inhibiting its growth and provoking mycelium browning (Figure 4). *Rahnella* sp. could be releasing compounds to the media that might be responsible for mushroom browning and mycelium growth inhibition. However, Rao et al. tested an aqueous extract of shiitake against *Rahnella*, obtaining low levels of inhibition [17]. It is remarkable that we did not find any inhibition effect for any of the tested *Pseudomonas* strains. This contrasts with previous research describing that *P. tolaasii* produces volatile compounds such as methanethiol, dimethyl disulfide and 1-undecene that inhibit in vitro mycelium growth of several fungi and causes mycelium browning [18].



**Figure 4.** Inhibition of *L. edodes* mycelium (in the center of the plate) by the isolated strains, and negative control (sterile distilled water). Strains were incubated for two weeks at 10 °C.

#### 4. Conclusions

We found that weight loss, sensory parameters, and the increase of microbial load were the main constraints to shiitake shelf life under different conservation conditions. The application of low temperatures reduced weight loss and extended the shelf life for two weeks. Although MAP was not able to extend the shelf life in comparison to refrigeration, it could be extremely recommendable to prevent weight loss. The microbial analysis of shiitake carpophores showed that *Pseudomonas* (4.7 log cfu/g) and *Enterobacteriaceae* (3.5 log cfu/g) were the most frequently isolated microbial groups. A total of 48 isolated strains were identified. Among them, *E. americana*, *Burkholderia* sp., *Serratia* sp., *Rahnella* sp. and some *Pseudomonas* strains were able to damage shiitake carpophores. Only *Rahnella* sp. showed an inhibitory effect on shiitake mycelium growth. The study demonstrated the major role of Gram-negative bacteria on *L. edodes* deterioration. Furthermore, the biodeterioration capacity of the isolated strains increased throughout the conservation period, with cap slices being more affected than the cap cuticle, and with bacterial strains isolated from *L. edodes* showing higher damaging ability on the mushroom species from which they were isolated than on *A. bisporus*.

**Supplementary Materials:** The following are available online at <http://www.mdpi.com/2073-4395/10/6/898/s1>, Figure S1: Levels of damage severity on *L. edodes* tissues by the microbial strains inoculated in the pathogenicity test analysed on experimental day seven; Table S1: Genbank Accession Numbers of Bacterial strains isolated from shiitake mushroom (*Lentinula edodes*) in this work and best BLAST hit sequences used for identifications.

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