

Upgrading of Mixed Food Industry Side-Streams by Solid-State Fermentation with *P. ostreatus*

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Abstract: In the frame of efforts to exploit agroindustrial side-streams and wastes (AISS) for added-value products that are based on single cell protein (SCP), mixed substrates consisting of brewer's spent grains (BSG), malt spent rootlets (MSR), cheese whey, molasses, orange, and potato pulps, were used for growth of the edible mushroom *Pleurotus ostreatus*. The substrates were mixed in various combinations, and were used for *P. ostreatus* growth at various conditions. The substrate, for which the highest sugar consumption, protein increase, and mycelium yield were observed, consisted of 20 mL molasses (4° Baume density), 20 mL potato pulp, 5 mL whey, 5 mL orange pulp, 30 g BSG, and 5 g MSR (at 25 °C and substrate pH 4). The mycelium-enriched product was analyzed for protein, fat, minerals (Ca, Mg, Fe, Cu), and aroma volatile compounds, indicating the potential for use as nutritious supplement for food, feed, or microbiology uses. The product was also autolyzed, freeze-dried, powdered, and analyzed for total ribonucleic acid content, showing the potential for use as a commercial natural food flavor enhancer.

Keywords: Agroindustrial side-streams; *Pleurotus ostreatus*; protein; volatiles; autolysis; RNA content; natural food additive

1. Introduction

Agro-industrial wastes and side-streams (AISS), which are produced in large amounts worldwide, are usually rich in fermentable components. Specifically, food industry side-streams, such as brewer's spent grains (BSG) and malt spent rootlets (MSR) from breweries, wasted fruits and vegetables from agricultural production, landfills, restaurant wastes, etc., whey from the dairy industries, and molasses from the sugar industries, have the high potential for biotechnological utilization. However, apart from molasses, they are used in limited amounts as livestock feed or are discarded in the environment since their high organic load makes biological treatment impossible [1–10]. Bio-utilization of these materials and the conversion to products of added-value is, therefore, of great importance from both economical and environmental points of view. Single cell protein (SCP) production has been the subject of extensive research mainly for the production of animal feeds that are enriched in protein by microbial treatment of various AISS using both submerged (SmF) and solid-state fermentation (SSF) processes. The most common examples of food grade SCP are baker's, brewer's, and distiller's yeasts, as well as food and feed supplements (as protein and vitamin sources and flavor enhancers). Molasses is an ideal carbon source that is commonly added to other substrates that contain low amounts of fermentable sugar, in order to facilitate their bioconversion [7,9,10]. Whey also has a huge potential for bioconversion to added-value products,

such as food additives, protein, SCP, organic acids, enzymes, bioethanol, etc. [6,9,10]. The development of efficient and cost effective processes for its exploitation is a great scientific and industrial challenge, since it is generated in large amounts and has a high organic load. BSG and MSR can both be treated with cellulolytic, amylolytic, and/or ligninolytic species, to produce enriched livestock feeds, or other products of added-value. The use of BSG for such applications is limited but with promising results, while MSR have not been very widely exploited [3–5,9,10]. The exploitation of the above AISS for SCP production of various species, such as *Kluyveromyces marxianus*, *kefir*, and *Saccharomyces cerevisiae*, was recently reported [9,10]. A promotional effect of addition of orange pulp was shown during a SmF process using mixed substrates of orange pulp, cheese whey, molasses, and potato pulp. Specifically, a 2- to 3-fold increase of cell mass was observed in the presence of orange pulp. Likewise, the promotional effect of BSG on cell growth during SSF of mixtures containing BSG, whey, molasses, potato pulp, MSR, and orange pulp, was also shown. The *K. marxianus* and *S. cerevisiae* growth was enhanced by three-fold and two-fold, respectively, in these substrates, thus proving their suitability for SCP production without need for synthetic nutrients [9,10]. Therefore, the mixed AISS can be used for low cost SCP production, achieving at the same time waste valorization and minimization.

The most studied food flavor enhancer is monosodium glutamate (MSG), although it has been accused for undesirable health effects, including the so-called *Chinese restaurant syndrome* [12,13]. Substitution of MSG with a safer natural flavor enhancer has been the subject of many research efforts based on the use of low cost biotechnological products, such as autolyzed microbial extracts [14,15]. Autolysis is a degradation process that produces soluble peptides, amino acids, nucleotides, and amino acid derivatives. Autolyzed SCPs can also be used as low cost extracts to replace the expensive commercial ones that are used in synthetic growth media. In this study, the enrichment of six AISS (BSG, MSR, molasses, whey, potato, and orange pulp) by SSF with *Pleurotus ostreatus* is described, as well as the autolysis and the proximate analysis of the produced SCP for potential use as multipurpose nutritious extract.

2. Materials and Methods

2.1. Materials

Cheese whey was obtained from the Agricultural Cooperatives Union of Kalavryta (Achaia province, Greece). Molasses was supplied by the B.G. Spiliopoulos S.A. alcohol distillery (Patras, Greece). BSG and MSR were supplied by the Athenian Brewery S.A. (member of the Heineken NV group, Patras, Greece). Their proximate composition has been reported in previous studies [5,9,10]. BSG were homogenized by blending and MSR were sieved to produce a fine powder. Sterilization of the substrates was done at 120 °C for 15 min. The potatoes and oranges were obtained from a local market. The external yellow exocarp of orange skins was removed and the fruit were blended to produce the whole orange pulp. In this pulp water was added at a ratio of 1:1 (by weight). In the same manner, the potato pulp was produced and diluted with two parts water (by weight).

2.2. *P. ostreatus* Growth on Mixed AISS by SSF

The edible fungus *P. ostreatus* was obtained by a local market and was grown on Malt Extract Agar (MEA). Specifically, cuttings with basidiospores were used to inoculate the medium in Petri dishes, which were then incubated at 25 °C for 5–7 days until white, fluffy mycelia appeared. For the study of *P. ostreatus* growth by SSF, 25 different mixtures of the AISS were used (Table 1).

Table 1. Composition of the mixed agroindustrial side-streams and wastes (AISS) substrates before and after solid-state fermentation (SSF) by *P. ostreatus* at 25 °C.

Substrate	Substrate Composition							Fermented Substrate Composition						
	Molasses	MSR	BSG	Whey	Potato Pulp	Orange Pulp	Water	Initial Protein	Final Protein	Protein Increase	Initial Sugar	Final Sugar	Sugar Reduction	Moisture
	mL	g	g	mL	mL	mL	mL	%w/w	%w/w	%	g/Kg	g/Kg	%	%
S1	5	5	50	5	5	5	20	19.3	44.2	24.9	8.9	0.4	95.5	86.7
S2	5	5	50	5	5	5	20	15.5	39.2	23.7	9.3	1.2	87.1	78.9
S3	5	12.5	5	5	5	5	20	15.5	32.6	17.1	17.1	0.4	97.7	84.7
S4	5	13	5	5	5	50	20	25.7	39.8	14.1	30.6	0.6	98.2	84.6
S5	5	5	5	5	50	5	20	40.4	58.8	18.4	22.4	1.5	93.3	90.5
S6	5	5	30	5	5	20	20	27.8	49.3	21.5	15.6	3.0	80.5	64.7
S7	20	5	30	5	20	5	20	16.8	43.2	26.4	34.0	0.0	100.0	58.4
S8	-	-	30	-	-	30	20	33.3	43.8	10.5	4.9	2.6	46.8	88.0
S9	-	30	-	-	-	30	20	32.9	34.6	1.7	5.5	0.9	83.3	87.3
S10	30	-	-	-	-	30	20	12.8	28.9	16.1	24.5	0.0	100.0	68.5
S11	-	-	-	10	-	30	20	30.7	35.2	4.5	9.2	3.3	64.0	88.0
S12	-	-	-	-	30	30	20	37.2	59.8	22.6	8.9	0.2	98.2	82.2
S13	30	-	30	-	-	-	20	13.7	13.8	0.1	30.6	17.6	42.4	72.1
S14	30	20	-	-	-	-	20	17.7	17.8	0.1	33.2	21.3	35.6	82.2
S15	-	-	30	-	30	-	20	31.0	31.6	0.6	3.6	0.0	99.4	82.5
S16	-	20	30	-	-	-	20	24.2	45.0	20.8	1.1	0.9	20.0	82.9
S17	20	-	45	-	-	-	20	24.3	31.5	7.2	22.9	0.0	100.0	77.2
S18	-	-	50	15	-	-	20	40.5	56.0	15.5	2.0	0.6	72.0	81.3
S19	-	20	-	-	10	-	20	40.3	42.6	2.3	0.2	0.2	8.7	81.9
S20	20	20	-	-	-	-	20	29.7	30.9	1.2	11.2	3.1	72.2	85.5
S21	-	20	-	60	-	-	20	27.9	34.4	6.5	5.7	2.1	62.3	81.1
S22	-	-	30	10	30	-	20	23.8	45.3	21.5	2.0	1.1	44.4	81.0
S23	-	20	-	70	35	-	20	32.3	53.7	21.4	3.6	0.4	88.8	81.8
S24	-	20	20	70	-	-	20	42.6	58.5	15.9	2.2	1.2	48.7	79.8
S25	-	20	20	-	30	-	20	32.8	51.4	18.6	1.2	0.6	48.7	82.2

All of the substrates were analyzed for initial crude protein content. The substrates were sterilized at 120 °C for 15 min. Amounts of 30 g of each substrate were spread in pre-weighed Petri dishes. From plates containing *P. ostreatus* mycelia grown on MEA, 3 pieces of 10 × 10 mm of agar with mycelia were cut and were vortexed into 30 mL Ringer's solution to remove the basidiospores. Aliquots of 1 mL of this suspension were used to inoculate the substrates. The cultures were incubated at 25 °C for 7–12 days. The growth time was defined as the time that was needed for the white mycelia to cover the surface of each substrate. After that the substrates covered with mycelia were oven-dried at 100 °C for 10 h and the final protein content was determined. The growth of *P. ostreatus* was subsequently studied on the best substrate at 37, 25, and 15 °C, and then at the optimum substrate, optimum incubation temperature, and at different pH values (4.0, 5.5, and 7.0). All of the experiments were performed in triplicate. Finally, *P. ostreatus* growth at larger scale was carried out at the optimum conditions, using different SSF set-ups: (a) polyethylene bags; (b) trays of 1 Kg capacity; and, (c) plastic perforated cylindrical bioreactors. Amounts of 1 Kg of sterilized substrate were placed into the three bioreactor types (in triplicate), and were inoculated, as described above, for the plate experiments. All systems were incubated at 25 °C for 30 days. The substrates were checked daily to detect mycelium appearance, and then small holes were opened in the bioreactors in order to allow for the possible occurrence of fruiting.

2.3. Autolysis of the SSF Product

Autolysis of the *P. ostreatus* SSF product (on the optimum substrate and growth conditions) was carried out by suspending in distilled water, adjusting the pH to 6.0, and incubating at 50 °C in a water bath for 24 h. The process was terminated by heating at 80 °C for 30 min. The suspension was centrifuged to remove the cell debris. The supernatant was concentrated in a water bath at 80 °C and the concentrate was then freeze-dried to produce the extract powder. Freeze-drying was carried out by cooling to −40 °C (3 °C/min) and freeze-drying overnight at $5\text{--}15 \times 10^{-3}$ Bar and −50 °C in a Freeze Dry System, Freezone 4.5 (Labconco, MO, USA).

2.4. Assays

Moisture, pH, ash, and total fat were determined, as previously described [9,10], with minor modifications. In brief, moisture (%w/w) was determined by drying at 80 °C overnight. pH was determined after suspending 10 g of sample in 100 mL water. Ash (%w/w) was determined by burning accurately weighed (1–2 g) amounts of the samples in porcelain crucibles on a burner flame, followed by ashing at 550 °C for at least 3 h. Cheese whey was evaporated before burning. Total fat (%w/w) was determined by the extraction of 5 g of dried sample (105 °C; 3 h) with diethyl ether in a Soxhlet apparatus for about 2.5 h. The extract was condensed in a rotary evaporator, dried at 105 °C for 0.5 h, and weighed [9,10]. In the case of cheese whey, the determination of fat was carried out according to the Gerber method for milk analysis, which involves the treatment with sulfuric acid and amyl alcohol for separation of fat, and measurement of the volume of fat after centrifugation.

Fermentable sugars (sucrose, fructose, lactose, glucose, galactose) and ethanol, were analyzed by HPLC on a Shimadzu (Kyoto, Japan) chromatograph that was equipped with a SCR-101N stainless steel column, a LC-9A pump, a CTO-10A oven (set at 60 °C), and a RID-6A refractive index detector. Water was used as mobile phase (0.8 mL min^{−1}) and a 1% v/v 1-butanol solution (Merck) was used as internal standard.

Crude protein was analyzed according to the Kjeldahl's method for the determination of total nitrogen in milk [9]. The real protein increase (RPI) of the samples that was attributed to the microbial biomass was calculated using the formulas (Eq. 1 and Eq. 2) [16]:

$$RP = \frac{(M_{final} \times P_{final}) - (M_{initial} \times P_{initial})}{M_{initial}} + P_{initial} \quad (\text{Eq. 1})$$

$$RPI = \frac{(RP - P_{initial})}{P_{initial}} \times 100 \quad (\text{Eq. 2})$$

in which $M_{initial}$ is the weight (g dry matter) before SSF, $P_{initial}$ is the initial protein content (% dry matter), M_{final} is the weight (g dry matter) after SSF, and P_{final} is the final protein content (% dry matter).

The determination of total phenolics in both organic (methanol) and the aqueous extracts of the samples was carried out spectrophotometrically, according to the Folin-Ciocalteu procedure [9,17]. Gallic acid solutions in methanol were used as standards. In brief, methanolic extracts of the samples were prepared after alkaline hydrolysis with 1 N NaOH, followed by centrifugation, acidification with conc. HCl to pH 2, extraction with 1:1 diethyl ether/ethyl acetate, evaporation of the solvents, and finally, the dilution of the dry residue with 12 mL methanol. The total phenolics were determined by measuring the absorbance of the extract/Folin-Ciocalteu reagent mixture at 760 nm [17].

The determination of the antioxidant capacity in both organic (methanol) and the aqueous extracts was performed by the DPPH radical scavenging method, based on the decrease in absorbance at 517 nm of a methanolic solution of DPPH against aqueous methanol solution as blank [9,17].

The determination of COD was carried out by both a colorimetric test [reaction with HgSO_4 , $\text{H}_2\text{SO}_4/\text{AgSO}_4$, and $\text{K}_2\text{Cr}_2\text{O}_7$, and titration of excess $\text{K}_2\text{Cr}_2\text{O}_7$ with $\text{FeSO}_4(\text{NH}_4)_2\text{SO}_4 \cdot 6\text{H}_2\text{O}$ and ferroin indicator], according to standard methods [9,18], and a photometric test (using an electronic digestion device HACH DRB-200, a spectrophotometer HACH DR/2400, and digestion reagent vials HACH COD Digestion Vials, Low Range 3–150 mg/L COD) [9]. Metals Ca, Mg, Fe, and Cu were determined on a AA-6500 Series Atomic Absorption Spectrometer (Shimadzu) after ashing at 550°C and dissolving in condensed nitric acid [10].

Major volatile compounds were determined by gas chromatography on a Shimadzu GC-8A system (Japan) with a stainless steel column TRB-624 (Teknokroma, Barcelona, Spain) (30 m, 0.32 mm i.d.; temperature 40–200°C by 10° C/min), He as carrier gas (23 cm³/s), and a Flame Ionization Detector (FID). The injection port and the detector temperatures were 260°C. Samples of 2 µL were injected directly into the column and quantification was done using calibration curves. The composition of headspace volatile compounds of the treated substrates after SSF was analyzed by solid-phase micro-extraction (SPME) gas chromatography mass spectrometry (GC/MS) on a Shimadzu GC-17A gas chromatograph (capillary column Supelco CO Wax-10 60 m, 0.32 mm i.d., 0.25 µm film thickness) coupled to a Shimadzu MS QP5050 mass spectrometer (70 eV ionization energy; 30–400 m/z mass range) [10]. The fiber that was used for absorption of volatiles was a 50/30 µm DVB/Carboxen/PDMS StableFlex for manual holder (Supelco, Bellefonte, PA, USA). For the SPME sampling, 5 g of sample were transferred to a sealed 20 ml glass vial, stirred for 5 min at 60°C, and the fiber was then exposed to the headspace for 45 min. Injection in the GC was performed in splitless mode. The oven temperature program was: 40°C for 5 min; rise to 110°C by 10°C/min; rise to 180°C by 2 °C/min; rise to 250°C by 10 °C/min; and, held for 6 min. Identification was carried out by comparing the retention times and mass spectra of volatiles to those of pure compounds (Sigma–Aldrich, Poole, UK), by mass spectra that were obtained from NIST107, NIST21, and SZTERP libraries, and by determining the Kovats' retention indexes and comparing with those that were reported in the literature [10].

The extract produced after autolysis and freeze-drying of the *P. ostreatus* SCP was analyzed for RNA content, using the orcinol method, as described by Herbert et al. [19], with some modifications. In brief, the samples were extracted with 0.5 M perchloric acid for 90 min at 38°C with shaking, and centrifuged; the supernatant contains all of the RNA that can be determined by the orcinol reagent (aqueous 0.1 g/L orcinol and 0.9 g/L $\text{FeCl}_3 \cdot 6\text{H}_2\text{O}$ in conc. HCl; ratio 1:4; used immediately after

preparation). RNA gives a color with the orcinol reagent at the standard heating time of 20 min, which is exactly equivalent to its content of purine-bound ribose. The absorbance was measured at 670 nm and the RNA content was calculated from a curve that was based on the reaction of the orcinol reagent with standard D-ribose solutions.

3. Results and Discussion

3.1. Composition of the AISS

The sugar, protein, fiber, and mineral content of the six AISS samples, as well as the presence of minor compounds, such as vitamins and trace elements, makes them suitable for microbial conversion. Previous analysis of these exact substrates showed that BSG, MSR, and cheese whey had a considerable protein content of about 31, 24, and 14 %w/w (on dry matter), respectively [9]. In brief, molasses, orange, and potato pulps had lower amounts of protein (av. 7.3, 2.9, and 2.5 %w/w on dry matter, respectively) [9]. Molasses contained the highest amount of fermentable sugar (42.5 %w/w), while orange pulp had the highest amount of phenolics content (15.8 mg/g dry matter in aqueous extract, and 5% mg/g DM in organic extract) [9]. On the other hand, despite the total phenolic content, cheese whey presented the highest antioxidant capacity (4.35 mg/mL IC₅₀) [9]. In this study, the analysis of Ca, Mg, Fe, and Cu (Table 2) showed that molasses had the highest minerals content (Ca and Mg) and orange pulp had the lowest. MSR also contained significant amounts of Mg, as reported in previous studies [20]. BSG also had a considerably high content of these minerals. Whey was found to be rich in Mg (2 g/kg), however, when compared with the concentrations that were reported in other publications, there seem to be discrepancies. The results that were obtained in this study are similar to those reported by Glass and Hendrick [21] with respect to Fe and Cu, but significantly higher than other references [22–24] regarding Mg. These differences are expected due to the different raw materials (milk) and different production process. BSG were also found to be rich in Mg. Also, the amount of Cu contained was much higher than that required daily by the human body [25], while the content of Fe was significantly lower [26]. It has been reported that BSG contain significant amounts of minerals and trace elements and are therefore suitable substrates for the production enzymes by SSF [27]. In the case of the other two metals, there are differences in the levels, which may be due to the way barley that is cultivated, the different harvest time, and the beer production process [27]. Table 2 shows that the orange pulp has a high Ca content, the second largest among the six AISS, and at the same time, the lowest Mg content. Still, the percentage of Cu that is contained in orange pulp is low (1 Kg of pulp would be required to cover the average daily diet requirements). Wide variations in the orange composition have also been reported in the literature, which are affected by several factors, including availability in soil, soil pH, uptake ability of the plant, use of fertilizers, irrigation water, etc. [28]. Finally, potato pulp seems to be another good source of Mg since its percentage is quite high. On the other hand, Fe was not detected, which is not in line with the common knowledge that potatoes are rich sources of iron [29,30]. However, there are also discrepancies in the literature, which may also be attributed to the different soil composition, plant physiology and agricultural practices.

Table 2. Metal elements contained in the used AISS.

Metal (mg/Kg)	MSR	BSG	Whey	Molasses	Orange Pulp	Potato Pulp
Mg	3970	605.6	2000	14,200	0.19	320.63
Ca	20.30	4.39	21.0	1636	285	1.10
Fe	96.02	4.56	0	70.9	-	-
Cu	19.38	24.62	3.66	6.03	3.21	3.34

3.2. *P. ostreatus* Growth in the Mixed AISS

In Table 3, the protein and sugar composition of the mixed AISS substrates before and after the growth of *P. ostreatus* by SSF at 25°C, are presented. The moisture content of the substrates after fermentation was 62.6–88.7%, which is suitable for *P. ostreatus* growth. The fungus consumed most of the fermentable sugar in most cases. The reduction of sugars and the increase of protein are illustrated in Figure 1. The highest protein increase can be observed for the S7 substrate (5 mL whey, 20 mL molasses, 5 mL orange pulp, 20 mL potato pulp, 30 g BSG, 5 g MSR, and 20 mL water), which also had the highest sugar consumption. Generally, *P. ostreatus* was able to grow on all of the substrates, which indicates the suitability of the mixed AISS as substrates for edible mushroom growth.

Table 3. Effect of temperature (15, 25, and 37°C) and pH (4.0, 5.5, and 7.0) on *P. ostreatus* growth in S7 by SSF.

pH	Temp.	Initial Sugar	Final Sugar	Final Protein	Yield	Final ash	Moisture
	°C	g/Kg	g/Kg	%w/w	g _{mycelium} /Kg _{substr}	%w/w	%
4.0	25	30.04	0.15	27.96	59.44	1.10	63.2
5.5	25	29.85	-	33.94	57.71	0.83	59.5
7.0	25	31.17	0.20	38.35	39.50	0.82	62.6
4.0	15	21.04	8.75	27.96	41.84	0.70	87.5
4.0	25	21.04	0.20	32.65	71.84	0.76	88.7
4.0	37	21.04	7.13	30.58	37.65	0.94	85.3

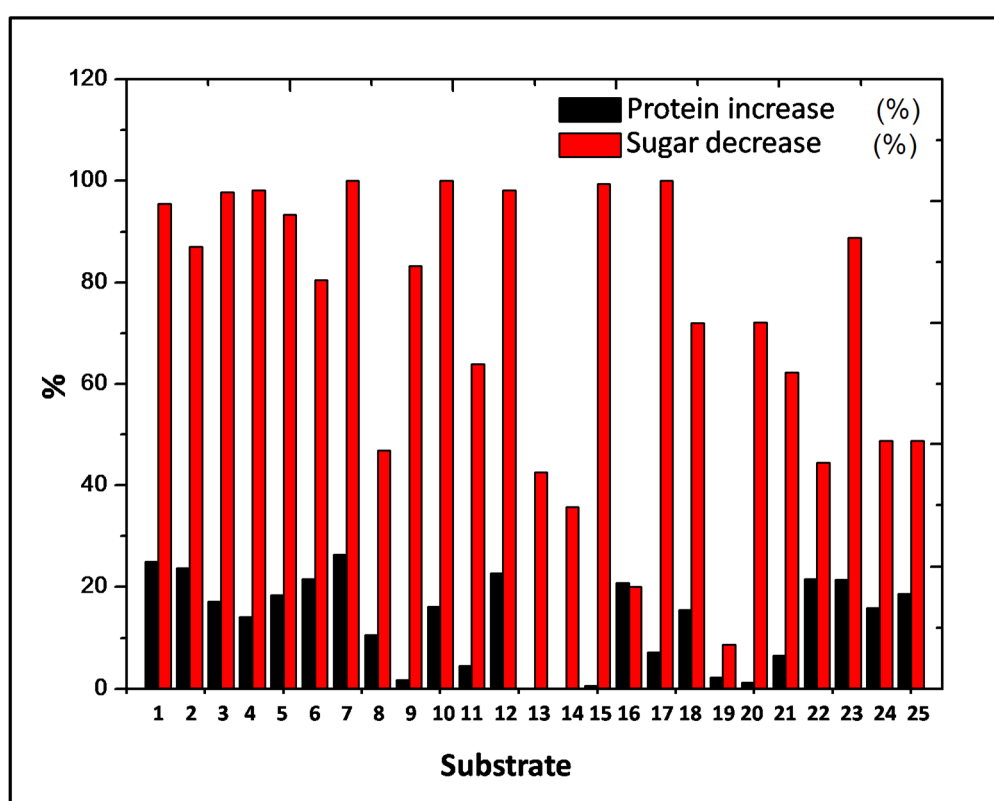


Figure 1. Sugar consumption and protein increase during *P. ostreatus* growth in the 25 mixed AISS substrates.

Subsequently, the effects of substrate pH (4.0, 5.5, and 7.0, at 25 °C) and temperature (15, 25, and 37 °C, at pH 4.0) on *P. ostreatus* growth on substrate S7 were studied. In this case, the biomass yield was expressed as g of mycelium/Kg of substrate, and the results are presented in Table 3. Higher mycelial yield was achieved at pH 4.0, while the lowest was at pH 7.0. This was expected as the optimal pH values for the growth of *Pleurotus* fungi range from 4.0 to 6.5 [31]. From Table 3, it can

also be seen that the fungus grows better at 25 °C. This is confirmed by other works that identified the optimal growth temperature of *P. ostreatus* to be about 22 °C [32–34].

For the scale-up experiments, the growth of *P. ostreatus* on the S7 substrate was then studied using three different bioreactor systems, as described in paragraph 2.2. After two days of incubation, the formation of mycelium appeared in all three systems, and at five days, it had covered the entire surface of the substrate. The incubation temperature was then reduced to 16 °C to facilitate the fruiting of the mushrooms, since the optimum temperature reported in the literature is 15–18 °C [32–34]. The incubation was continued for a further 35 days during which (every seven days) water was added to maintain stable substrate moisture. At the end of the 40 days of incubation, no appearance of fruit from the edible fungus was observed, but the mycelium, on the contrary, showed alterations in color and gas (probably methane) formation. Thus, further research is needed on the conditions (moisture, temperature, addition on trace elements, etc.) for mushroom fruiting bodies using the proposed mixed AISS.

3.3. Composition of the SSF Product and Autolyzate

The product of SSF of the mixed AISS (substrate S7) by *P. ostreatus* was analyzed for protein, total fat, ash, essential elements (Ca, Mg, Fe, Cu), and volatiles content (Table 4). The protein and fat levels were relatively high, higher than those that were reported in other works for fermented barley or dry citrus pulps [35–37], which makes the enriched material a potential nutritious food and feed supplement. The protein content was also higher when compared to that of the same AISS substrates that were fermented by SSF with *kefir* (23.6%), and slightly lower than those that were fermented by *S. cerevisiae* and *K. marxianus* (38.4% and 33.7%, respectively) [10]. The fat content (14.4%) was slightly higher than those of the *S. cerevisiae* and *kefir* fermented AISSs (12.9% and 12.3%, respectively), but lower than that of *K. marxianus* (25.5%) [10]. Regarding the minerals content, the *P. ostreatus* SSF product had similar amounts of Ca and Cu content with those of *S. cerevisiae*, *kefir*, and *K. marxianus* [10], but much higher Fe content when compared with these products (0.9–2.3 mg/Kg) [10]. Regarding Mg, the substrate that was fermented by *S. cerevisiae* contained the highest content of all compared products (771.1 mg/Kg) [10]. In general, it can be said that these essential minerals were found in significant amounts in the fermented AISS products.

Table 4. Composition of the product of S7 SSF by *P. ostreatus* and its autolyzate.

Characteristic		Content
SSF product	Moisture	76.0±2.3
	Dry matter	24.0±2.1
	Protein (%w/w)	30.3±1.9
	Ash (%w/w)	0.93 ± 0.1
	Fat (%w/w dry matter)	14.4±1.1
	Ca (mg/Kg)	91.1±8.3
	Mg (mg/Kg)	324.5±24.1
	Fe (mg/Kg)	17.8±2.1
	Cu (mg/Kg)	21.8±2.1
	Major volatile compounds (mg/Kg) (GC/FID)	
	Acetaldehyde	277.7±26.3
	Ethyl acetate	142.4±18.7
	Propan-1-ol	21.5±3.2
Isobutyl alcohol	31.9±4.3	
Amyl alcohols	7.5±1.3	
SSF product after autolysis	RNA (mg/100 g dry extract)	721.8±31.9

The composition of headspace volatile compounds that were identified by SPME GC/MS (totally 30 different compounds) in the same AISS as well as in their mixtures fermented by *S. cerevisiae*, *K. marxianus*, and *kefir*, has been reported in a previous study [10]. The volatile

compounds that contribute to the aroma of fermented products are produced during fermentation or derive from the raw materials. Some of the major volatile by-products that are produced by fermentation are acetaldehyde, ethyl acetate, propan-1-ol, isobutanol, and amyl alcohols (Table 4). These compounds were all found (by GC/FID analysis) in the SSF product of *P. ostreatus*, especially acetaldehyde, which was found at relatively high levels. Acetaldehyde has a pungent, etherial, fruity, or musty odor, depending on its concentration. Also, it may accumulate during mushroom growth at high temperatures due to anaerobic respiration and can have an inhibitory effect on mushroom growth (*Spawn-burning* syndrome) [38]. Therefore, inefficient oxygen supply may be the reason for the high acetaldehyde levels and the inability of the mushroom to grow fruiting bodies, and, in the future, further research will be needed focusing on the optimization of growth temperature and oxygen supply for scale-up of edible mushrooms production on these AISS.

Similar but fewer compounds were (positively) that were identified by SPME GC/MS analysis of the headspace of the *P. ostreatus* AISS fermentation product (Table 4). The principal compounds that are responsible for the aroma of raw and thermally processed mushrooms are aliphatic alcohols and ketones with eight carbon atoms [39]. However, only 1-octen-3-ol was positively identified. p-Xylene (1,4-dimethylbenzene) was possibly identified as a degradation product of aromatic amino acids, such as phenylalanine [40], as it usually found in other fermented and ripened products, such as cheese [41]. The presence of C6, C7, and C9 carbonylic compounds (2-hexanone, 3-heptanone, 2-heptanone, 5-nonanone) can be related to fatty acids oxidation [42]. Especially, methylketones, such as 2-hexanone and 2-heptanone, are very typical products of fungal species, biosynthesized by decarboxylation of β -keto-acids that are produced by β -oxidation of free fatty acids. 1,1,3,5-Tetramethyl-cyclohexane has also been previously identified in food as a result of microbial spoilage [42]. Finally, methyl pentanoate esters, such as isopropyl valerate, also occur naturally in fermented beverages, fruits, and vegetables [43,44].

The *P. ostreatus*SSF product of substrate S7 was autolyzed, freeze-dried, powdered, and analyzed for total RNA content. The texture of the powder was similar to that of commercial yeast extract. It contained about 722 mg total RNA/100 g of dry powder, which was similar to that reported for *K. marxianus* grown on whey [45]. RNA is a component of microbial extracts, which, after partial hydrolysis, provides oligonucleotides that can enhance the flavor and taste (*umami*) when used as additives in food [45].

4. Conclusions

The mixed AISS substrate for which the highest sugar consumption, protein increase, and *P. ostreatus* mycelium yield were observed, consisted of 20 mL molasses (diluted to 4° Baume density), 20 mL potato pulp, 5 mL whey, 5 mL orange pulp, 30 g BSG, and 5 g MSR (at 25 °C and substrate pH 4). In all of the experiments, orange pulp and BSG seemed to act as growth promoting ingredients. The produced fermented mycelium-enriched product was rich in protein, minerals (mainly Mg), and aroma volatile compounds, indicating the potential for use as protein-rich nutritious supplement for food, feed, or microbiology uses. High acetaldehyde levels were observed, which possibly inhibited the fruiting of the mushroom. Finally, the autolyzed dried product had high total ribonucleic acid content showing potential for use as natural food flavor enhancer. Future research will be needed focusing on optimization of growth temperature and oxygen supply for the scale-up of edible mushrooms production on these AISS, as well as the use of the autolyzed enriched substrates as food flavor enhancers and microbiology extracts.

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