

# Sustainable Food Technology

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## Sustainability Spotlight Statement

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Ensuring global food security for a growing population requires sustainable, health-promoting protein sources beyond resource-intensive conventional livestock farming. Mycoprotein produced via submerged fermentation of edible fungi offers a promising alternative, with low carbon-footprint. This study advances the sustainable production of *Pleurotus ostreatus* mycoprotein by identifying key fermentation parameters that enhance mycelium biomass titer and protein content while controlling mycelial pellet morphology suitable for food applications. A significant advancement is the demonstration of oleic acid as a novel carbon source, highlighting a path towards valorizing lipid-rich feedstocks and reducing competition with food crops. This work directly supports UN SDG 2 (Zero Hunger) by enhancing mycoprotein availability and SDG 12 (Responsible Consumption and Production) by establishing efficient bioprocesses to support circular bioeconomy.



1 **Mushroom Mycelium as a Sustainable High-Protein Food Source: Effects of Submerged**  
2 **Fermentation Conditions on Mycoprotein Production and Mycelium Morphology**

New Article Online  
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7  
8 **Abstract**

9 Harnessing edible fungal mycelium as a source of health-promoting food presents a transformative  
10 pathway to enhancing global food security, and addressing the demand by world growing population in  
11 the 21<sup>st</sup> century. The mycoprotein from fungal mycelium also serves as a sustainable alternative to  
12 resource-intensive animal-sourced protein. This study investigated how submerged fermentation  
13 influences mycelium morphology and mycoprotein production by edible fungus *Pleurotus ostreatus*.  
14 Key parameters, including inoculum fragmentation, types of carbon sources, carbon-to-nitrogen (C/N)  
15 and carbon-to-phosphorous (C/P) ratios, and agitation rates, were systematically evaluated. Principal  
16 Component Analysis (PCA) was employed to identify the most influential parameters and elucidate  
17 their correlations with production metrics. Controlled inoculum fragmentation, inoculum density and  
18 agitation were found to be crucial for achieving uniformity of pelletized mycelium and improving  
19 productivity. Protein content and production increased with a decreasing C/N ratio, achieving a  
20 maximum of 39.7% (of dry biomass) and 3.89 g L<sup>-1</sup>, respectively, while not significantly influenced by  
21 the C/P ratio. Oleic acid, a plant-based fatty acid, was demonstrated for the first time as a sole non-  
22 sugar carbon source for *P. ostreatus* cultivation, achieving a biomass yield comparable to glucose.  
23 Maximum biomass production (12.9 g L<sup>-1</sup>) and productivity (1.61 g L<sup>-1</sup> d<sup>-1</sup>) were attained through  
24 desirable inoculum fragmentation, an inoculum density of 40 mg L<sup>-1</sup>, an agitation rate of 150 rpm, a  
25 carbon loading of 3.6 g L<sup>-1</sup>, a C/N ratio of 2.6 and a C/P ratio of 52.9. These findings provide valuable  
26 insights into establishing an efficient and sustainable biorefinery for mycelium-based foods.

27 **Keywords:** Fungal Mycelium; Mycoprotein; Submerged Fermentation; Pellet Morphology;  
28 Sustainable Biorefinery



## 35 Sustainability Spotlight Statement

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43 feedstocks and reducing competition with food crops. This work directly supports UN SDG 2 (Zero  
44 Hunger) by enhancing mycoprotein availability and SDG 12 (Responsible Consumption and  
45 Production) by establishing efficient bioprocesses to support circular bioeconomy.

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## 68 1. Introduction

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69 Securing a sustainable protein supply for a growing global population represents one of the  
70 most critical food security challenges.<sup>1</sup> A strategic transformation framework is needed towards  
71 building a more resilient and sustainable food supply to support global food security, as resource-  
72 intensive livestock farming cannot meet the accelerating demand alone.<sup>2</sup> Mycoprotein, referring to the  
73 whole-cell, protein-rich biomass derived from fungi, stands at the forefront of this transition.<sup>3</sup> Cultivated  
74 from edible mycelium, it offers not only a scalable, high-quality vegan protein, containing all essential  
75 amino acids, but also provides balanced nutrition and health-promoting benefits, such as a high content  
76 of  $\beta$ -glucan fiber known to improve gut health and help regulate blood cholesterol levels, along with  
77 other key metabolites like choline, an essential nutrient that supports liver function and brain health.<sup>3,4</sup>  
78 The edible fungus *Pleurotus ostreatus* is an especially promising microorganism for this purpose. Its  
79 fruiting body is well known as oyster mushroom, and considered nutritious owing to the significant  
80 content of carbohydrates (43.42 - 66.54 % biomass, dry basis (d.b.)), proteins (17.06 - 55.4 % biomass,  
81 d.b.), and dietary fibers (7.43 - 23.63 % biomass, d.b.).<sup>5-7</sup> *P. ostreatus* is also a good source of vitamins  
82 (e.g., A, C, E and B2), minerals (e.g., sodium, potassium, magnesium and calcium), and essential amino  
83 acids like lysine, phenylalanine, methionine, threonine, leucine and valine.<sup>5, 8, 9</sup> The species has been  
84 explored for the production of various bioactive compounds such as enzymes (e.g., laccase and  
85 cellulase)<sup>10, 11</sup> and exopolysaccharides (e.g.,  $\beta$ -glucans)<sup>7</sup>, which find diverse applications in the field of  
86 environment and agriculture, and food, beverage, health, pulp, textile and other industries<sup>12, 13</sup>.

87 Cultivation of fungal mycelium is a sustainable approach, offering profound environmental  
88 advantages over conventional livestock farming, including substantially lower land use, water  
89 consumption, and carbon emissions.<sup>14</sup> By growing mycelium in efficient, controlled fermentation  
90 systems, this technology can directly meet the growing market demand for vegan proteins and novel  
91 food ingredients, decoupling food production from large-scale agriculture. Moreover, fungi can be  
92 successfully cultivated on waste substrates, such as pomelo peel powder, solid digestate from anaerobic  
93 digester, wheat straw, tree leaves, apple peels and banana peels.<sup>15-17</sup> Solid-state fermentation has been  
94 widely prevalent for fungal cultivation mainly attributed to the lower input of energy and water and  
95 cost-effectiveness in terms of potential to propagate on cheaper substrates and valorize agro-industrial  
96 wastes.<sup>18</sup> However, the cultivation method is constrained by the requirement of large footprint,  
97 inconvenient separation of mycelium biomass that is adhered to the fermented solid substrate, and  
98 challenges to maintain homogeneity of pH, temperature and moisture throughout fermentation, which  
99 may potentially lead to batch-to-batch variability in the characteristics of the biomass produced.<sup>19</sup>  
100 Submerged fermentation on the contrary is advantageous in terms of much smaller footprint, shorter  
101 cultivation period, capability of regulating and maintaining uniform culture conditions throughout  
102 fermentation, and easiness of scaling-up.<sup>20</sup> Furthermore, many fungi species, including *P. ostreatus*,  
103 form spherical mycelium pellets in submerged fermentation, which facilitates efficient harvesting of



104 the mycelial biomass from the culture medium for subsequent downstream applications.<sup>21, 22</sup> While  
105 solid-state fermentation of *P. ostreatus* has been extensively investigated for the traditional cultivation  
106 of its fruiting bodies, there are comparatively few studies on its submerged cultivation.<sup>19, 23</sup> Submerged  
107 fermentation of *P. ostreatus* is advantageous for large-scale mycoprotein production owing to its  
108 significantly shorter production cycles (days vs. weeks), higher volumetric productivity, and the ability  
109 to maintain precise control over mycelium morphology and biomass composition suitable for  
110 developing food ingredients. Moreover, submerged fermentation has been an emerging area for  
111 developing mycelium-based foods, such as cultured caviar and mycoboba from *Aspergillus awamori*  
112 (manufactured by Optimized Foods), meat analogue for Harbin red sausage by *Fusarium venenatum*<sup>24</sup>,  
113 oncom from *Neurospora intermedia*<sup>25</sup>, and dairy analogue for yogurt from *Fusarium* strain *flavolapis*  
114 (manufactured by Nature's Fynd), suggesting the need of more comprehensive research.

115 The type and composition of the substrate play a crucial role in determining the production of  
116 biomass as well as metabolites thereof. For instance, biomass production in submerged fermentation of  
117 *Monascus purpureus* was reported to be significantly influenced by the choice of carbon (*e.g.*, glucose,  
118 sucrose, xylose and citric acid) and nitrogen (*e.g.*, yeast leavening, NH<sub>4</sub>Cl, peptone and soybean meal)  
119 sources.<sup>21</sup> Similarly, the availability of glucose and yeast extract, and supplementation of olive oil was  
120 observed to have critical impact on protein production in *A. oryzae* and *N. intermedia*.<sup>26</sup> Investigating  
121 such non-sugar carbon sources (*e.g.*, fats/oil) is critical for developing more sustainable and cost-  
122 effective bioprocesses, as lipids and fatty acids can often be derived from low-cost agri-food industry  
123 byproducts, thereby avoiding competition with food crops. However, these lipids are typically used  
124 only as supplements rather than as the primary substrate; to the best of our knowledge, no studies have  
125 yet investigated the use of a long-chain fatty acid as the *sole* carbon source for submerged fungal  
126 cultivation. Furthermore, like nitrogen forms a key part of proteins, phosphorus forms a critical portion  
127 of phosphorylated biomolecules, *e.g.*, adenosine triphosphate (ATP), adenosine diphosphate (ADP) and  
128 adenosine monophosphate (AMP), and the reductant nicotinamide adenine dinucleotide phosphate  
129 (NADPH), which are essential molecules involved in energy transfer in the cells.<sup>27</sup> Only a few studies  
130 have explored the effect of phosphorus on production of mycelium biomass and protein, with existing  
131 research being limited to other products like manganese-dependent peroxidase, glucosamine and lipid  
132 in fungi species, *e.g.*, *Phanerochaete chrysosporium*, *Mucor indicus* and *Umbelopsis vinacea*.<sup>27-29</sup>  
133 Beyond specific nutrients, the stoichiometric relationship between carbon and nitrogen (C/N ratio) and  
134 carbon and phosphorus (C/P ratio) are critical determinants with potential influence on cell growth and  
135 metabolic pathways. Precise regulation of these ratios is therefore indispensable for targeted modulation  
136 of metabolic fluxes, enabling the preferential allocation of cellular resources towards biomass  
137 accumulation or protein biosynthesis.<sup>30</sup> However, in contrast to well-studied platforms like bacteria and  
138 microalgae, these stoichiometric relationships remain critically under-investigated in fungal submerged  
139 cultivation. Specifically, while the C/N ratio has received limited attention, the effect of the C/P ratio



140 on biomass and protein synthesis remains entirely unexplored. Moreover, other parameters, including  
141 inoculum properties (e.g., inoculum density) and agitation speed have been identified as influential  
142 process conditions for mycelial growth, in addition to pellet morphology in submerged fermentation of  
143 certain species like *Ganoderma curtisii* and *Aspergillus awamori*.<sup>31,32</sup> Based on the current state-of-the-  
144 art for submerged fungal fermentation, a comprehensive investigation is still required to evaluate the  
145 influence of inoculum properties, different sugar and non-sugar carbon sources, and key stoichiometric  
146 ratios (C/N and C/P) on the interplay between mycelial morphology, biomass accumulation, and protein  
147 synthesis to ultimately develop an economically viable and environmentally sustainable process for  
148 global food security.

149 Therefore, to address the research gap, this study aimed at developing a robust submerged  
150 fermentation process for *Pleurotus ostreatus* to maximize the co-production of mycelial biomass and  
151 protein. The central hypothesis was that the systematic modulation of physical factors to control  
152 mycelial morphology, combined with the regulation of chemical factors to tailor nutrient availability  
153 and metabolic flux, would substantially enhance both biomass titer and protein content. The objectives  
154 of this research were to: (1) evaluate the effects of key inoculum and fermentation parameters, including  
155 inoculum fragmentation level, inoculum density, and agitation rate, on mycelial pellet morphology; (2)  
156 systematically investigate the influence of media formulation factors, including both sugar and non-  
157 sugar (such as long-chain fatty acid) as sole carbon sources and the stoichiometric C/N and C/P ratios,  
158 on the production of mycelium biomass and protein; and (3) to elucidate the complex correlations  
159 between various submerged fermentation parameters and the resulting production metrics, thereby  
160 identifying the most favorable conditions for sustainable mycoprotein production.

## 161 2. Materials and methods

### 162 2.1. Organism and culture conditions

#### 163 2.1.1. Reagents and chemicals

164 All the chemicals were of analytical grade. Glucose and fructose were procured from Ward's  
165 Science (Rochester, NY, USA). Lactose, xylose and lauric acid were procured from Thermo Fisher  
166 Scientific (Waltham, MA, USA). Magnesium sulfate and monopotassium phosphate were purchased  
167 from Ambeed Inc. (Buffalo Grove, IL, USA). Yeast extract and potato dextrose agar were purchased  
168 from Hardy Diagnostics (Santa Maria, CA, USA). Oleic acid was purchased from TCI (Portland, OR,  
169 USA). 3,5-Dinitrosalicylic acid was procured from Alfa Aesar (Ward Hill, MA, USA).  
170 Chloramphenicol was purchased from G Biosciences (Saint Louis, MO, USA). All the reagents were  
171 prepared in de-ionized water.

#### 172 2.1.2. Genome sequencing and species identification



173 As the source of fungal mycelium, the Oyster mushroom fruiting bodies were sourced from a  
174 local store (Ithaca, NY, USA). A 2 mm x 2mm piece of tissue was aseptically removed from the  
175 mushroom flesh in the middle of the stem butt-end and then cultured on potato dextrose agar (PDA) for  
176 approximately 10 days.<sup>33</sup> The Genomic DNA of the strain was extracted following a previous method  
177 <sup>34</sup> with minor modification: a small fragment of mycelium was collected from a mature plate and placed  
178 into a Polymerase Chain Reaction (PCR) tube with 50  $\mu$ L of nuclease-free water. The mixture was  
179 briefly vortexed to ensure homogeneity. The tube was then incubated in a thermocycler (BIO-RAD,  
180 Model T100 Thermal Cycler, Hercules, CA, USA) for 15 minutes at 95 °C to facilitate cell lysis and  
181 DNA release through heat shock.<sup>34</sup> Following incubation, the sample was immediately cooled in an ice-  
182 water bath for 1 minute to stabilize the released genomic DNA. The resulting extract was used directly  
183 as a template for PCR amplification without further mechanical or chemical treatment. PCR was carried  
184 out using ITS 5 (Forward, GGA AGT AAA AGT CGT AAC AAGG, position 1737 - 1758) and ITS 4  
185 (Reverse, TCC TCC GCT TAT TGA TAT GC, position 2390 - 2409) primers (Supplementary  
186 information, Fig. S1). The amplification was performed in a 25  $\mu$ L total reaction volume containing  
187 12.5  $\mu$ L of 2X Master Mix, 0.5  $\mu$ L of Forward and Reverse Primer each, 0.15  $\mu$ L GoTaqFlexi Taq  
188 Polymerase, 9.85  $\mu$ L of nuclease-free water and 1.5  $\mu$ L of the extracted DNA. The PCR tubes were  
189 then placed in a thermocycler and run using the appropriate conditions: initial denaturation at 95°C for  
190 5 min; 35 cycles of denaturation at 95°C for 1 minute, annealing at 56°C for 1 minute and extension at  
191 72 °C for 1 min, and a final extension at 72 °C for 10 min. A 3  $\mu$ L aliquot of the resulted sample was  
192 then electrophoresed on 1.5% (w/v) agarose gel. The gel was loaded with a 100bp DNA ladder  
193 (Promega Bench Top DNA Ladder) as molecular weight markers for comparison. The gel box  
194 (Instaview Blue LED Transilluminator, Model, E1201-BLT) (Accuris Instruments, Edison, NJ, USA)  
195 was run at 120 Volts for 30 minutes until the ladder bands were appropriately separated. The gel was  
196 visualized using the gel box orange filter. Based on the primers used, we observed the sample bands  
197 appearing on the agarose gel as expected (Supplementary information, Fig. S2). The PCR sample was  
198 then cleaned using ExoSAP and subjected to sanger sequencing by Cornell Biotechnology Resource  
199 Center (BRC) (Ithaca, NY, USA). The ITS sequences were then compared with other ITS sequences in  
200 the GenBank database using the native nucleotide Basic Local Alignment Tool (BLAST) function  
201 accessible through the National Center for Biotechnology Information (NCBI) (Supplementary  
202 information, Fig. S3 and S4), based on which the species was identified as *Pleurotus ostreatus*.

### 203 2.1.3. Media, inoculum preparation and culture conditions

204 The fungal mycelia were cultivated on agar plates in basal media containing (g L<sup>-1</sup>): glucose  
205 (30.0), yeast extract (5.0), KH<sub>2</sub>PO<sub>4</sub> (1.0), MgSO<sub>4</sub> (0.244) and agar (20.0), as previously described by  
206 Shen et al. <sup>35</sup> with minor modifications. The agar plates were incubated at 30 °C under static conditions  
207 for 6–7 days until the mycelia colonized approximately 90% of the agar surface area, as determined by  
208 the radial expansion of the mycelial front from the center of the plate. To ensure the collection of



209 actively growing biomass, plates were harvested before the mycelium reached the edges of the petri  
210 dish.<sup>35</sup> The plates were maintained at 4 °C for further use. Inoculum preparation was performed in 300-  
211 mL shake flasks containing 150 mL of liquid basal media (without agar). The liquid media was  
212 inoculated with 3-4 units of 8-10 mm mycelial discs cut out from the pre-grown agar plates using sterile  
213 forceps, and the seed flasks were incubated at 30 °C and 150 rpm for 6 days in a shaker incubator  
214 (MaxQ™ 6000, Thermo Scientific, Waltham, MA, USA). The pH of the media was adjusted to 6.0 by  
215 adding 2 M HCl or 2 M NaOH. The medium was sterilized by autoclaving at 121 °C for 15 min. The  
216 basal medium was selected based on preliminary trials (data not shown), where it supported faster  
217 mycelial growth compared to potato dextrose agar/broth to reach comparable biomass levels for  
218 inoculation.

219 For inoculation of the experimental flasks, 80 mL of the seed culture was transferred into a  
220 sterile 100-mL beaker containing majority of smaller pellets of comparable sizes. The pellets were  
221 fragmented through homogenization by a mechanical homogenizer (T25 digital Ultra Turrax, IKA,  
222 Staufen, Germany) at 5000 rpm for 40 seconds. This specific fragmentation level was selected based  
223 on the study described later in the Section 2.2. The resulting inoculum granulometry exhibited a mean  
224 fragment size of 924.5 µm. For all subsequent experiments, 1 mL of this homogenized pellet suspension  
225 was used as inoculum, corresponding to an inoculum density of ~20 mg L<sup>-1</sup> biomass (dry basis (d.b.))  
226 in 150 mL of liquid media. The experimental flasks comprised of the same basal media as was used for  
227 that of inoculum generation (unless otherwise specified), except that it was supplemented with 0.1 g L<sup>-1</sup>  
228 chloramphenicol to prevent microbial contamination in the submerged fermentation of fungi.<sup>35</sup>

## 229 2.2. Evaluation of the effect of inoculum fragmentation through homogenization

230 *Pleurotus ostreatus* was investigated for the production of biomass and morphology of pellets  
231 under varying levels of inoculum fragmentation comprising of different homogenization speeds and  
232 durations. Pre-grown seed culture was subjected to four types of fragmentation treatments following a  
233 method previously reported by Shen et al.<sup>35</sup>, with modifications based on our preliminary experimental  
234 trials (data not shown) on *P. ostreatus*: F1 (pellets homogenized at 3000 rpm for 5 seconds), F2  
235 (homogenized at 5000 rpm for 10 seconds), F3 (homogenized at 5000 rpm for 40 seconds), and control  
236 (no homogenization). The pellet fragmentation was achieved using a mechanical homogenizer (T25  
237 digital Ultra Turrax, IKA, Staufen, Germany). Inoculation was performed using the method as  
238 mentioned earlier in Section 2.1. The cultures were incubated for 10 days under continuous shaking of  
239 150 rpm at 30 °C.

## 240 2.3. Submerged fermentation with variable media composition

### 241 2.3.1. Evaluation of the effect of different carbon sources and carbon loading



242 Five carbon sources including glucose, lactose, fructose, xylose and oleic acid were studied as  
243 the sole carbon substrate for submerged fermentation of *P. ostreatus*. Three carbon loadings to  
244 cultivation media were investigated including 0.88, 3.60 and 6.33 g L<sup>-1</sup>, corresponding to glucose  
245 concentrations of 2.19, 9.0 and 15.82 g L<sup>-1</sup>, lactose concentrations of 2.08, 8.55 and 15.03 g L<sup>-1</sup>, fructose  
246 concentrations of 2.19, 9.0 and 15.82 g L<sup>-1</sup>, xylose concentrations of 2.19, 9.0 and 15.82 g L<sup>-1</sup>, and oleic  
247 acid concentrations of 1.14, 4.71 and 8.28 g L<sup>-1</sup>, respectively. Yeast extract was used as nitrogen source  
248 for all assays at a fixed concentration (5 g L<sup>-1</sup>). The cultures with different carbon loadings (0.88, 3.60  
249 and 6.33 g L<sup>-1</sup>) were incubated under constant agitation of 150 rpm at 30 °C until the exhaustion of  
250 carbon source (*e.g.*, total sugars or fatty acid) in the respective media, *i.e.*, on Day 5, Day 8 and Day 12.

### 251 2.3.2. Evaluation of the effect of different C/N and C/P ratios

252 *P. ostreatus* was cultured in media containing different concentrations of nitrogen, including  
253 0.06, 0.39, 0.72, 1.05 and 1.38 g L<sup>-1</sup>, which were prepared by adding yeast extract at equivalent  
254 concentrations (0.5, 3.5, 6.5, 9.5 and 12.5 g L<sup>-1</sup>, respectively). The experiment was performed using the  
255 basal liquid media containing 9 g L<sup>-1</sup> glucose (and 1 g L<sup>-1</sup> KH<sub>2</sub>PO<sub>4</sub>), which corresponded to equivalent  
256 C/N ratios of 65.5, 9.4, 5.0, 3.4 and 2.6, respectively.

257 Further, the organism was cultured under varying concentrations of phosphorus (0, 0.07, 0.52,  
258 0.98 and 1.43 g L<sup>-1</sup>) in the media, which correspond to KH<sub>2</sub>PO<sub>4</sub> concentrations of 0, 0.3, 2.3, 4.3 and  
259 6.3 g L<sup>-1</sup>. The study was carried out in media containing 9 g L<sup>-1</sup> glucose (and 5 g L<sup>-1</sup> yeast extract),  
260 which corresponded to equivalent C/P ratios of 52.9, 6.9, 3.7 and 2.5, respectively.

261 The cultures were incubated for 8 days under constant agitation of 150 rpm at 30 °C.

### 262 2.4. Evaluation of the effect of different inoculum densities and agitation rates

263 The organism was investigated in submerged cultivation at varying inoculum densities of 20  
264 and 40 mg L<sup>-1</sup> biomass (d.b.), under different agitation rates of 75, 150 and 255 rpm at 30 °C. The  
265 cultures were grown for 8 days in media supplemented with glucose at a carbon loading of 3.6 g L<sup>-1</sup>,  
266 yeast extract at a nitrogen loading of 1.38 g L<sup>-1</sup> and KH<sub>2</sub>PO<sub>4</sub> at a phosphorous loading of 0.07 g L<sup>-1</sup>,  
267 which were selected based on previous studies carried out in Section 2.3.

### 268 2.5. Analytical methods

#### 269 2.5.1. Determination of the size distribution of inoculum fragments and pellets

270 The size distribution of inoculum fragments was determined using Mastersizer particle size  
271 analyzer (MS-2000, Malvern Instruments Limited, Westborough, MA, USA) by gradually adding 30  
272 mL sample after different fragmentation treatments. The instrument mathematically interprets light  
273 scattering pattern by the fragments and fits the measured data to estimate a volume-based (% v/v)  
274 particle size distribution in the range of 10 – 3000 µm, which is obtained as a bell-shaped plot.



275 For determination of pellet sizes after submerged fermentation, 10 mL of homogeneously mixed  
 276 culture was poured into a petri dish, and mycelial pellets were spread out across the dish to remove any  
 277 clumps and aggregates, and visualize discrete units of pellets. Images of the petri dish were captured  
 278 inside a photo light box and the sizes of individual pellets were estimated using the software, ImageJ,  
 279 developed by the National Institutes of Health (Bethesda, MD, USA).

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### 280 2.5.2. Estimation of biomass, residual sugar and fatty acid concentrations

281 For sampling of biomass, the experimental flask was mixed thoroughly by gentle shaking to  
 282 avoid sedimentation of mycelium pellets, and 10 mL of the culture was collected by pouring into a  
 283 sterile measuring cylinder. The mycelium biomass was separated from the liquid medium using vacuum  
 284 filtration through a pre-weighed Whatman filter paper (United Scientific, Libertyville, IL, USA),  
 285 assisted by a vacuum pump (Rocker 400, Rocker Scientific, Kaohsiung City, Taiwan). The biomass  
 286 residue was washed thoroughly using de-ionized water (150 mL) for two cycles and subjected again to  
 287 vacuum filtration to remove residual water and any surface-adhered medium components. The filtered  
 288 and washed samples were dried at 50 °C in a convective oven (VWR Oven F Air 3.65CF, VWR  
 289 International, Radnor, PA, USA) until it attained a constant weight with change of no more than  
 290  $\pm 0.5\%$ .<sup>21</sup> Moisture content of mycelium from different assays was observed to be within a range of 90.0  
 291 to 96.3 % (Supplementary information, Table S5). The concentration of biomass in the fermentation  
 292 media was expressed as dry weight (DW) of biomass per liter of media ( $g L^{-1}$ ).

293 Biomass productivity was calculated as per Eq. (1):

$$294 \text{Biomass productivity } (g L^{-1} d^{-1}) = \frac{\text{Biomass concentration } (g L^{-1})}{\text{Fermentation period } (d)} \quad (1)$$

295 The initial media and cell-free spent media were collected and analyzed for the content of  
 296 residual reducing sugar through DNS (3,5-dinitrosalicylic acid) assay as described earlier by Miller<sup>36</sup>,  
 297 and measuring the absorbance using UV-Vis Spectrophotometer (UV5Bio, Mettler Toledo, Columbus,  
 298 OH, USA) at 540 nm. Similarly, the cell-free supernatant was analyzed for residual concentration of  
 299 fatty acid spectrophotometrically using commercial free fatty acid assay kit (Sigma-Aldrich, St. Louis,  
 300 MO, USA) and determining the absorbance at 570 nm.

301 Biomass yield with respect to substrate ( $Y_{x/s}$ ) was calculated as the ratio of biomass  
 302 concentration ( $g L^{-1}$ ) at a given time point and the corresponding concentration of carbon source  
 303 consumed, as per Eq. (2):

$$304 \text{Biomass yield } (g g^{-1}) = \frac{\text{Biomass concentration } (g L^{-1})}{\text{Concentration of carbon source consumed } (g L^{-1})} \quad (2)$$

### 305 2.5.3. Estimation of intracellular proteins



306 Harvested mycelium biomass initially separated from liquid media through vacuum filtration  
 307 was subjected to lyophilization using a freeze-dryer with cycles of 12 h freezing and 12 h drying (EW-  
 308 03336-44, Harvest Right, Salt Lake City, UT, USA). To evaluate the nutritional quality of the harvested  
 309 mycelium biomass, the intracellular protein content was analyzed as previously described by Bakratsas  
 310 et al.<sup>23</sup> with minor modifications. 50 mg of freeze-dried biomass was resuspended in 5 mL of phosphate  
 311 buffer (pH 7.0) and subjected to ultrasonication (Q125, QSonica, Newtown, CT, USA) at 8 kHz (40%  
 312 intensity) in a pulse mode (15 s ON, 3 s OFF) for 6 min. The sample was placed in ice water bath during  
 313 ultrasonication to eliminate the risk of protein degradation owing to excess heat generating from the  
 314 sonication process. Intracellular contents were obtained in the supernatant by centrifuging (Centrifuge  
 315 5810 R, Eppendorf, Hamburg, Germany) the sample at 5000 rpm for 15 min. Protein content was  
 316 estimated through Bradford Assay (Pierce Bradford Protein Assay Kit, Thermo Scientific, Waltham,  
 317 MA, USA), using bovine serum albumin (BSA) as standard. In brief, 1.5 mL Bradford reagent was  
 318 mixed with 30  $\mu$ L of cell-free supernatant and the mixture was incubated for 10 minutes, followed by  
 319 the measurement of absorbance at 595 nm using a spectrophotometer. The absorbance value was  
 320 converted into mass of proteins using the correlation based on calibration curve ( $y = 1.0404 x$ ;  $R^2$   
 321  $= 0.9923$ ):

$$322 \quad \text{Concentration of protein in the sample (mg mL}^{-1}\text{)} = \frac{\text{Absorbance (595 nm)}}{1.0404} \quad (3)$$

323 Protein content in the biomass (d.b.), protein concentration in the fermentation media,  
 324 productivity and protein yield were calculated as per Eq. (4), (5), (6) and (7):

$$325 \quad \text{Protein content (\%)} = \frac{\text{Mass of protein (g)}}{\text{Mass of biomass (d.b.) (g)}} \times 100\% \quad (4)$$

$$326 \quad \text{Protein concentration (g L}^{-1}\text{)} = \text{Protein content (\%)} \times \text{Biomass concentration (g L}^{-1}\text{)} \quad (5)$$

$$327 \quad \text{Protein productivity (g L}^{-1}\text{d}^{-1}\text{)} = \frac{\text{Protein concentration (g L}^{-1}\text{)}}{\text{Fermentation period (d)}} \quad (6)$$

$$328 \quad \text{Protein yield (g g}^{-1}\text{)} = \frac{\text{Protein concentration (g L}^{-1}\text{)}}{\text{Concentration of carbon source consumed (g L}^{-1}\text{)}} \quad (7)$$

## 329 2.6. Statistical analysis

330 Experiments evaluating carbon source types and carbon loading for glucose and lactose were  
 331 performed in triplicate; all other experiments were conducted in duplicate (Supplementary information,  
 332 Table S3). Results were expressed as the mean  $\pm$  standard deviation of the replicates. Statistical  
 333 significance of the difference between the means was estimated through one-way or two-way analysis  
 334 of variance (ANOVA) with a post-hoc Tukey test using the Origin Pro software (OriginLab,  
 335 Northampton, MA, USA) at a significance level ( $\alpha$ ) of 0.05. Principal Component Analysis and Pearson  
 336 Correlation were performed using Origin Pro software.



### 337 3. Results and discussion

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#### 338 3.1. Effect of inoculum fragmentation

339 The size of mycelium fragments in the inoculum critically influences the development of new  
340 pellets and biomass accumulation.<sup>31</sup> To this end, inoculum was subjected to varying levels of  
341 fragmentation to obtain mycelium fragments of various sizes, viz., control: no homogenization; F1:  
342 3000 rpm, 5 seconds; F2: 5000 rpm, 10 seconds; and F3: 5000 rpm, 40 seconds. The mean sizes of  
343 mycelium fragments resulting from different treatments including control, F1, F2 and F3 were 1110.25,  
344 1089.13, 1036.37 and 924.5  $\mu\text{m}$ , respectively. Fig. 1A shows the size distribution of mycelium  
345 fragments in the inoculum. After fragmentation, the fraction of mycelium fragments (% v/v of the  
346 inoculum) having sizes in the range of 1000 – 2500  $\mu\text{m}$  were 53.9, 51.9 and 46.3, respectively, in  
347 control, F1 and F2 treatments. Whereas, F3, which was subjected to the highest degree of fragmentation  
348 resulted in a major fraction (64.5%, v/v) of the fragments in a size a range less than 1000  $\mu\text{m}$ , while  
349 only 35.5% exhibited sizes more than 1000  $\mu\text{m}$ . This fragmentation had a direct effect on the  
350 development of new pellets, which were harvested after 10 days. The pellets sizes were observed to  
351 decrease with increasing degree of fragmentation, resulting in mean diameters of 10.9, 6.41, 3.95 and  
352 2.26 mm, respectively, from control, F1, F2 and F3 treatments (Fig. 1B). Furthermore, the pellet size  
353 distribution became less scattered and more even with increasing fragmentation level. The control and  
354 lower fragmentation levels (F1 and F2) resulted in lower pellet count, which did not differ significantly  
355 among the treatments ( $p > 0.05$ ); whereas, F3 exhibited substantially higher pellet count (count: 2400)  
356 relative to other treatments ( $p < 0.05$ ) (Fig. 1C). As observed from Fig. 5A, while control, F1 and F2  
357 resulted in larger-sized mycelium pellets with spiky and rougher surfaces, F3 generated more compact  
358 pellets with relatively smoother surfaces and more homogenous size distribution. The underdeveloped  
359 pellets were observed as numerous fine and short thread-like structures dispersed all over the medium,  
360 and were excluded from size and count estimations. To the best knowledge of the authors, this is the  
361 first study demonstrating the effect of inoculum fragmentation level on resulting pellet sizes and number  
362 in submerged cultivation of *P. ostreatus*.

363 Moreover, effect of inoculum fragmentation levels was investigated on different growth  
364 parameters, including biomass production, productivity and yield. Lower biomass titer and productivity  
365 of 1.8 g L<sup>-1</sup> and 0.18 g L<sup>-1</sup>d<sup>-1</sup>, respectively, were observed for control, whereas, relatively higher titers  
366 (6.65 – 12.8 g L<sup>-1</sup>) and productivities (0.66 – 1.28 g L<sup>-1</sup>d<sup>-1</sup>) were observed for F1, F2 and F3 (Fig. 1D),  
367 although the values did not vary significantly among the different fragmentation treatments ( $p > 0.05$ ).  
368 Biomass yields across different treatments varied in the range of 0.66 – 1.57 g g<sup>-1</sup> (Fig. 1E). These  
369 observations are in close alignment with the study reported for *Inonotus hispidus* culture, where, the  
370 biomass production was observed to increase with increment in homogenization rate and time from  
371 6000 to 26000 rpm and 1 to 3 min, respectively, resulting in maximum biomass of 10.29 g L<sup>-1</sup>.<sup>35</sup>  
372 Moreover, Shen et al. <sup>35</sup> reported that lower shear rate and shorter shear time yielded loose mycelial

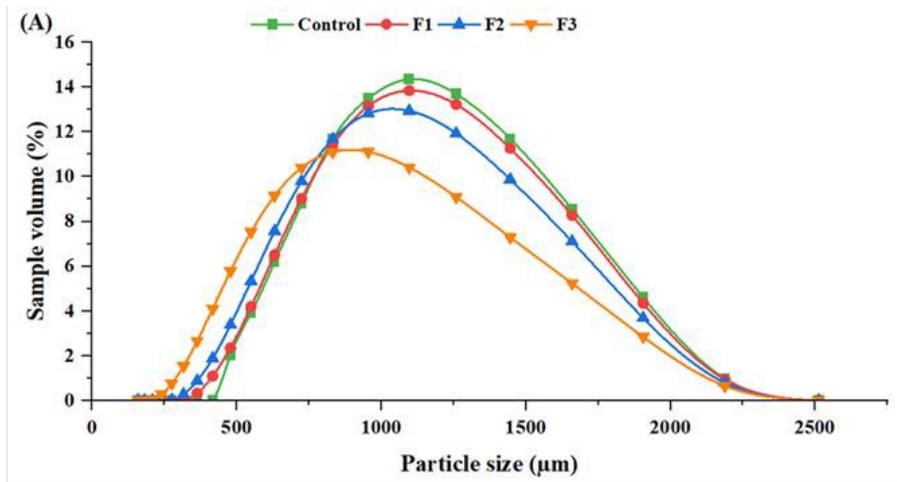


373 clumps and heterogeneous pellets, similar to our control; whereas, high shear rate and longer shear time  
374 produced well-dispersed evenly distributed mycelia, analogous to our F3 treatment. The dispersed  
375 morphology seems essential for creating higher surface area per unit volume of fragmented mycelium  
376 which can potentially enhance the uptake of nutrients and oxygen from the liquid medium, consequently  
377 leading to improved growth. Furthermore, the enhanced biomass titer and productivity under different  
378 fragmentation treatments (F1, F2 and F3) may also be attributed to availability of a greater number of  
379 densely-branched hyphal tips per unit length of an inoculum fragment, promoting higher germination  
380 capacity.<sup>35</sup> Another study also reported a similar pattern for biomass production in *Aspergillus niger*  
381 involving dispersion of inoculum.<sup>37</sup> Implementation of pellet recycling strategy revealed that  
382 fragmentation of mycelium pellets before cultivation could enhance biomass production from 27.0 g L<sup>-1</sup>  
383 (1<sup>st</sup> batch) to 32.1 g L<sup>-1</sup> (8<sup>th</sup> batch); whereas, direct inoculation with pellets (without fragmentation)  
384 resulted in decline in biomass production from 28.2 g L<sup>-1</sup> (1<sup>st</sup> batch) to 9.2 g L<sup>-1</sup> (6<sup>th</sup> batch), thus,  
385 demonstrating the significance of inoculum fragmentation.<sup>37</sup>

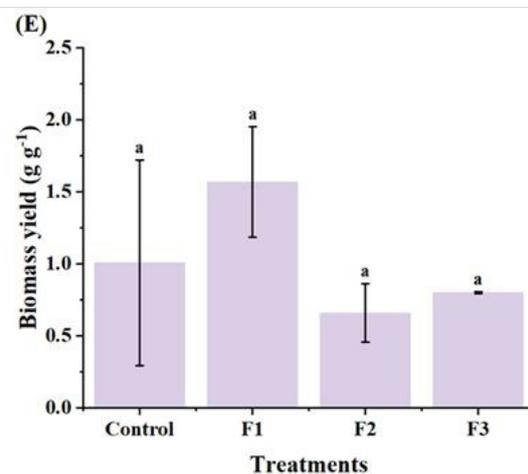
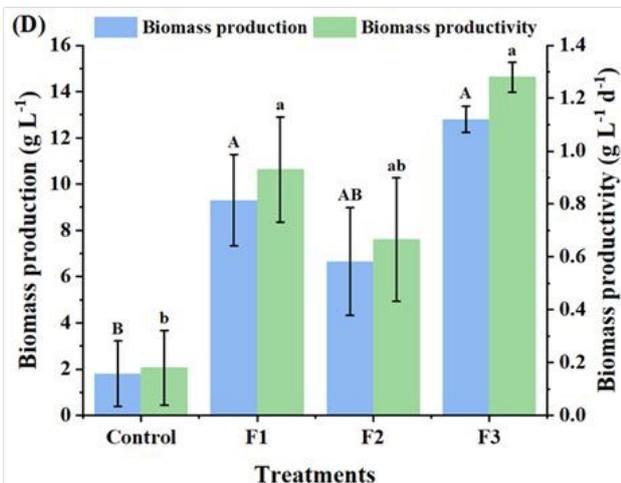
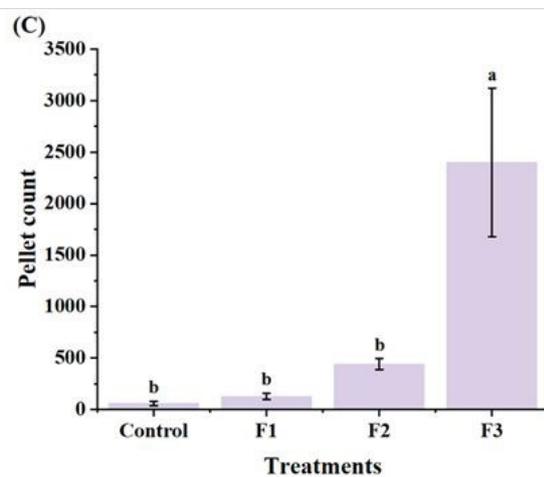
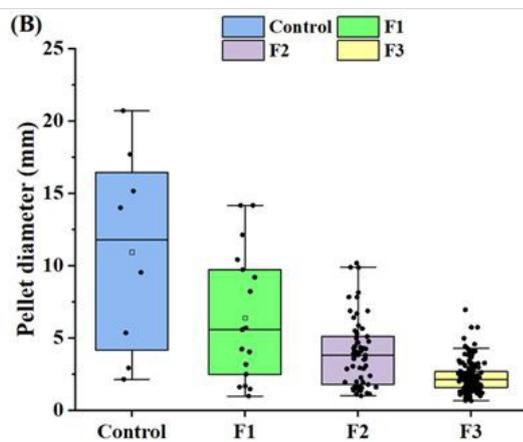
386 In the present study, since, no significant differences were observed between the treatments F1,  
387 F2 and F3 in terms of biomass titer, productivity and yield (Fig. 1D and E), therefore F3 was selected  
388 as the inoculum fragmentation level for further experiments, owing to significantly higher biomass  
389 production and productivity (compared to control), as well as compact and relatively uniform pellet  
390 sizes (Fig. 5A).

391





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392

393 **Fig. 1.** Effect of different fragmentation levels on (A) size distribution of mycelium fragments in  
 394 inoculum, (B) diameter of mycelium pellets in the culture on Day-10 (Quartile summary of the Box  
 395 and Whisker plot is provided in Supplementary information, Table S1), (C) number of pellets in the  
 396 culture, (D) biomass production and productivity, and (E) biomass yield. Control: no homogenization;



397 F1: 3000 rpm, 5 s; F2: 5000 rpm, 10 s; and F3: 5000 rpm, 40 s. \* Groups that share same letters are  
 398 considered statistically insignificant. View Article Online  
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### 399 3.2. Effect of media composition on submerged fermentation of *P. ostreatus*

#### 400 3.2.1. Effect of carbon source and loading on mycelium biomass

401 *P. ostreatus* was cultivated with different sugar and non-sugar carbon sources including  
 402 monosaccharides (glucose, fructose and xylose), disaccharide (lactose), and long-chain fatty acid (oleic  
 403 acid) at varying carbon loadings of 0.88, 3.60 and 6.33 g L<sup>-1</sup>, which were harvested on Day 5, Day 8  
 404 and Day 12, respectively. Among all sole carbon sources, the highest mycelium biomass production  
 405 and productivity were obtained using glucose. Biomass production was observed to improve with  
 406 increment in carbon loading of glucose from 0.88 – 6.33 g L<sup>-1</sup> (equivalent glucose concentrations: 2.19  
 407 – 15.82 g L<sup>-1</sup>), resulting in a maximum titer of 11.2 g L<sup>-1</sup> (Fig. 2A). However, highest biomass  
 408 productivity (1.09 g L<sup>-1</sup>d<sup>-1</sup>) was obtained at carbon loading of 3.6 g L<sup>-1</sup> (Fig. 2B). Bakratsas et al.<sup>23</sup>  
 409 reported similar patterns in *P. ostreatus* biomass production which was observed to increase from 6 to  
 410 approx. 14 g L<sup>-1</sup> with the increase in glucose concentration from 5 to 20 g L<sup>-1</sup>. It was interesting to  
 411 observe that oleic acid, as a non-sugar sole carbon source, achieved a higher biomass yield than majority  
 412 of sugar-based carbon sources in this study. A biomass production of 8.33 g L<sup>-1</sup> was obtained from oleic  
 413 acid at a carbon loading of 6.33 g L<sup>-1</sup>; and its biomass productivity was 0.69 g L<sup>-1</sup> d<sup>-1</sup> at the same carbon  
 414 loading which was not significantly different from the productivity attained at 3.6 g L<sup>-1</sup> carbon loading.  
 415 Furthermore, oleic acid exhibited higher biomass yield in the range of 1.0 – 1.27 g g<sup>-1</sup> at different carbon  
 416 loadings, and was observed to be comparable with glucose at higher loadings (3.6 – 6.33 g L<sup>-1</sup>) (Fig.  
 417 2C).

418 Filamentous fungi are known to assimilate fatty acids from fats and oils within endogenous  
 419 organelles like peroxisome mediated by  $\beta$ -oxidation and glyoxylate pathways resulting in the  
 420 production of carbohydrates, energy and acetyl-CoA.<sup>38</sup> Oleic acid is an 18-carbon unsaturated fatty acid  
 421 with 76.5% (w/w) carbon content in a molecule; hence, this higher energy density possibly accounts for  
 422 the higher biomass production owing to higher energy accumulation in *P. ostreatus* in the present study.  
 423 Oleic acid is the major fatty acid comprising of 55–83% of olive oil.<sup>39</sup> Nazir et al.<sup>26</sup> demonstrated the  
 424 use of olive oil (40 g L<sup>-1</sup>) as sole carbon source to grow *Aspergillus oryzae* in presence of yeast extract,  
 425 attaining a biomass titer of 13.21 g L<sup>-1</sup>, which was relatively higher than the biomass production (5.63  
 426 g L<sup>-1</sup>) obtained using 20 g L<sup>-1</sup> glucose as sole carbon source. Moreover, researchers have demonstrated  
 427 the use of olive oil as a medium supplement to enhance biomass production. For instance, Rosales-  
 428 López et al.<sup>31</sup> reported that supplementation of 1% v/v olive oil to the culture medium could increase  
 429 biomass production by 32% and 240% relative to optimized basal medium (containing glucose) and  
 430 commercially available potato dextrose broth, respectively, in submerged fermentation of *Ganoderma*  
 431 *curtisii*. However, to the best knowledge of the authors, there are no other studies demonstrating the



432 cultivation of filamentous fungi utilizing a long-chain fatty acid as the sole carbon source. This finding  
433 is particularly significant from a sustainability standpoint, as it demonstrates the potential to valorize  
434 plant-based or lipid-rich waste streams, such as used cooking oils or agricultural byproducts, which do  
435 not compete with the arable land and food crops required for sugar-based feedstocks like glucose and  
436 fructose. In the present study, an attempt was also made to utilize lauric acid as carbon source under  
437 different carbon loadings (0.88 – 6.33 g L<sup>-1</sup>). However, no growth was observed, primarily owing to the  
438 higher melting point (~43 °C), leading to a clear phase separation, and flaking out of the added lauric  
439 acid as a solid layer on top of the aqueous media under specified culture conditions (30 °C and 150  
440 rpm). Whereas, oleic acid, could form an oil-in-water emulsion<sup>40</sup> under the given conditions. Although,  
441 initial addition of oleic acid led to formation of a distinct layer on the surface of the aqueous medium;  
442 however, this phase separation gradually diminished during the cultivation period at 150 rpm. While no  
443 external emulsifying agents were utilized, the dispersion of the fatty acid was likely facilitated by the  
444 continuous mechanical agitation in conjunction with the biological activity of the fungus. *P. ostreatus*  
445 is known to secrete extracellular lipases<sup>41</sup> and polysaccharides with surfactant-like properties (e.g.,  $\beta$ -  
446 glucan)<sup>42</sup>, which can reduce interfacial tension and promote the emulsification of lipids in submerged  
447 culture. This biological stabilization of the oleic acid likely enhanced its bioavailability to the inoculum  
448 fragments for uptake and growth.

449 The biomass production, productivity and yield were observed to be comparable between other  
450 sugars like lactose, fructose and xylose at the highest carbon loading of 6.33 g L<sup>-1</sup>, with values  
451 significantly lower ( $p < 0.05$ ) than that obtained with glucose and oleic acid (Fig. 2A, 2B and 2C).  
452 However, substantially low biomass production and productivity were observed with xylose at 3.6 g L<sup>-1</sup>  
453 carbon loading. These results are in alignment with other studies which have identified glucose as a  
454 superior carbon source that provided higher cell biomass and product titers, among other sugars. For  
455 instance, higher production of biomass and exopolysaccharides (EPS) have been reported in *P.*  
456 *ostreatus* and other filamentous fungi, e.g., *Inonotus hispidus*, using glucose as carbon source, in  
457 contrast to other substrates, including lactose and fructose, at the same concentration.<sup>19, 35</sup> Li et al.<sup>43</sup>  
458 justified the slower growth rate of *Neurospora crassa* on xylose through the gene expression profile of  
459 sugar transporters; deciphering that out of 39 putative sugar transporter genes in the genome of *N.*  
460 *crassa*, 15 and 6 genes were expressed respectively when D-xylose and D-glucose were used as carbon  
461 sources. This suggests that glucose, being the preferred and most efficiently metabolized sugar, is  
462 handled by a small number of specialized and effective transporters. However, xylose triggers a broader,  
463 less-specialized response, upregulating a larger suite of transporters as part of a wider metabolic  
464 adjustment to a less-desirable carbon source.

### 465 3.2.2. Effect of carbon source and loading on mycelium protein



466 The protein content (% biomass, d.b.) and protein production from submerged fermentation of  
467 *P. ostreatus* were observed to increase with higher carbon loadings from 0.88 to 6.33 g L<sup>-1</sup>, and attained  
468 respective maximum values with glucose (31.1% and 3.47 g L<sup>-1</sup>), followed by oleic acid (22% and 1.83  
469 g L<sup>-1</sup>) (Fig. 2D and 2E). However, protein content and production were found to be relatively lower  
470 with other carbon sources like lactose, fructose and xylose, which could be attributed to corresponding  
471 lower biomass yield and production from these sugars, as compared to glucose and oleic acid. Hamza  
472 et al.<sup>19</sup> reported similar relationship between biomass and protein production in *Pleurotus ostreatus*,  
473 where protein production exhibited exponential increase from Day 6 to Day 10, with exponential growth  
474 of biomass following Day 5 of cultivation. Furthermore, Bakratsas et al.<sup>23</sup> demonstrated significantly  
475 higher protein content and production in *P. ostreatus* biomass cultivated using glucose (29.0 ± 2.0%  
476 and 3.32 ± 0.20 g L<sup>-1</sup>) as compared to lactose, fructose and xylose on Day 8 of cultivation, which is well  
477 in accordance with the findings of the present study. The protein content and production from oleic acid  
478 were 7.99 – 21.98% and 0.12 – 1.83 g L<sup>-1</sup>, respectively, under carbon loadings from 0.88 to 6.33 g L<sup>-1</sup>,  
479 which were significantly higher than majority of sugar substrates, but lower than glucose. This could  
480 be attributed to possible metabolic switch in the cells, resulting in the diversion of energy from fatty  
481 acid towards accumulation of fat in the cells, as a priority over synthesis of proteins. This is corroborated  
482 by Nazir et al.<sup>26</sup>, wherein, *A. oryzae* biomass accumulated 14% protein and 34% fat from 40 g L<sup>-1</sup> olive  
483 oil, while it produced 48% protein and 3% fat from 20 g L<sup>-1</sup> glucose. Similarly, protein productivity and  
484 protein yield from glucose were observed to attain maximum values of 0.29 g L<sup>-1</sup> d<sup>-1</sup> and 0.30 g g<sup>-1</sup>,  
485 respectively, which were significantly higher than that from oleic acid (0.15 g L<sup>-1</sup> d<sup>-1</sup> and 0.22 g g<sup>-1</sup>)  
486 (Fig. 2F and 2G) and other sugars.

487 Overall, the superior performance of protein production by *P. ostreatus* in presence of glucose,  
488 followed by oleic acid, and relatively lower performance in presence of other sources, e.g., lactose,  
489 fructose and xylose, could be attributed poor efficiency of assimilation of these sugars into the cells as  
490 compared to glucose and oleic acid. As the maximum biomass productivity and protein yield were  
491 observed with glucose at carbon loading of 3.6 g L<sup>-1</sup> (culture period: 8 days), these set of conditions  
492 were selected for performing further experiments, keeping in view the industrial viability of the process.  
493 Fig. 5B shows the morphology of the pellets grown using different carbon sources at loading of 3.6 g  
494 L<sup>-1</sup>. Glucose and lactose were observed to produce several round pellets with relatively uniform size  
495 distribution, as compared to fructose and xylose which yielded majority of pellets with rough and spiky  
496 surfaces and a non-uniform size distribution. Pellets generated from oleic acid-containing medium had  
497 interesting morphological features which were prominently distinguishable from the ones yielded from  
498 sugar substrates. For instance, oleic acid yielded a greater number of compact pellets with even  
499 smoother surfaces and a more homogenous size distribution as compared to the rest of the carbon  
500 sources, including glucose and lactose. The enhanced pellet morphology (compact, smooth, uniform)  
501 observed with oleic acid suggests a potential role of lipids in modulating mycelial aggregation and

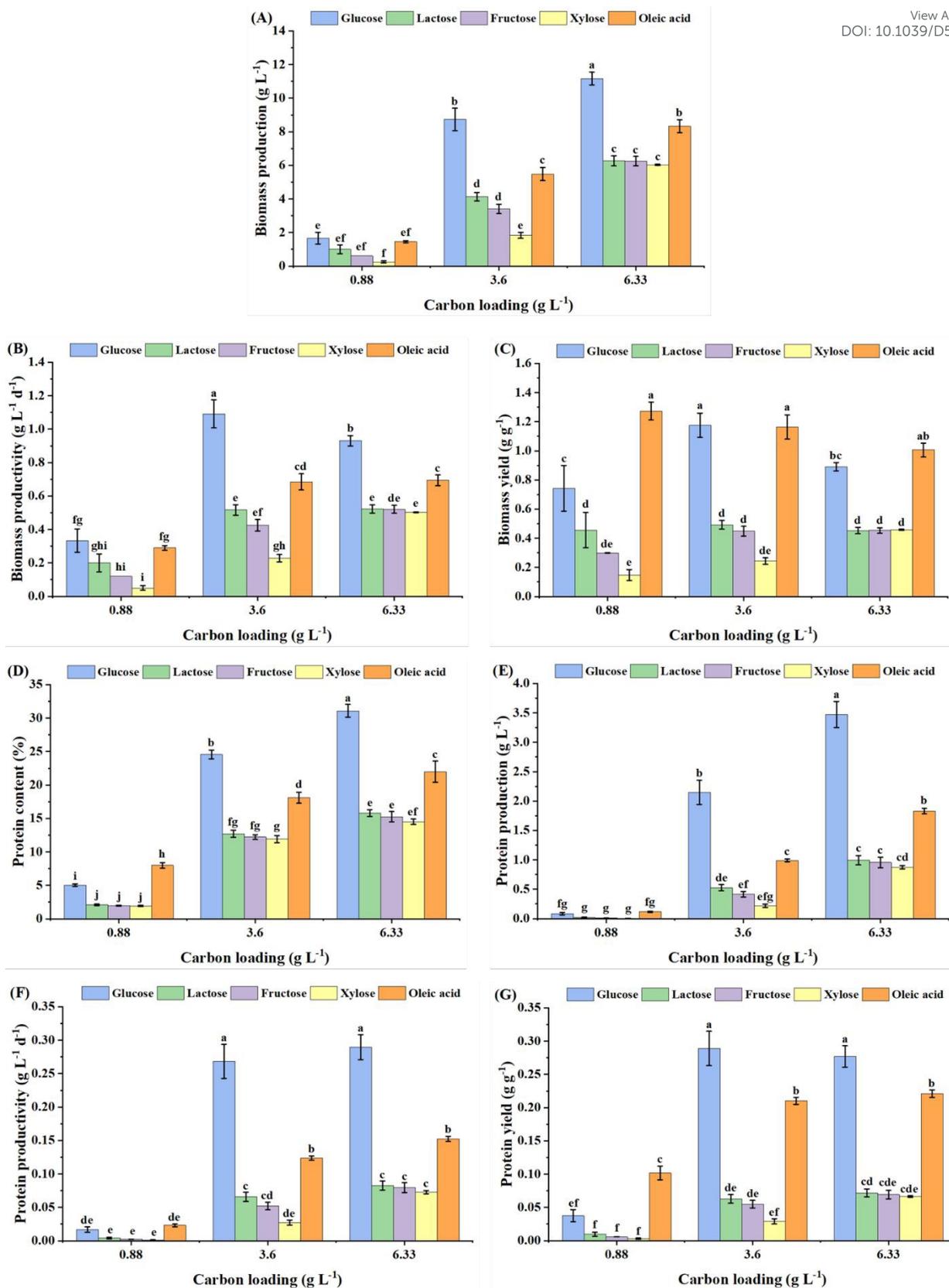


502 morphology. This aligns with Cheng et al. <sup>44</sup>, who improved *Pleurotus* sp. pellet compaction and  
503 circularity using olive oil (containing oleic acid) combined with the surfactant Tween 80 (a synthetic  
504 source of oleic acid). Cheng et al. <sup>44</sup> speculated that Tween 80 might assist in olive oil emulsification,  
505 implying the lipid phase influences pellet formation. Observations from our study indicate oleic acid  
506 alone may promote similar characteristics, possibly by altering hyphal surface interactions due to its  
507 hydrophobicity, thus mimicking surfactant effects. Alternatively, this could be attributed to the potential  
508 of *P. ostreatus* to secrete extracellular lipases <sup>41</sup> and polysaccharides with surfactant-like properties  
509 (e.g.,  $\beta$ -glucan) <sup>42</sup>, facilitating the emulsification of oleic acid. Although, the exact mechanism warrants  
510 further investigation, these findings underscore the significant impact of lipid-based carbon sources on  
511 fungal pellet morphology.

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513

514 **Fig. 2.** Effect of different carbon loadings and carbon types on (A) biomass production, (B) biomass  
 515 productivity, (C) biomass yield, (D) protein content, (E) protein production, (F) protein productivity



516 and (G) protein yield in *Pleurotus ostreatus*. \* Groups that share same letters are considered  
517 statistically insignificant. View Article Online  
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### 518 3.2.3. Effect of C/N and C/P ratios on mycelium biomass

519 *P. ostreatus* was cultivated under different C/N ratios of 65.5, 9.4, 5.0, 3.4 and 2.6, which were  
520 achieved by varying the nitrogen loading in the media to 0.06, 0.39, 0.72, 1.05 and 1.38 g L<sup>-1</sup>,  
521 respectively, at a fixed carbon loading. Biomass production and productivity were observed to increase  
522 significantly with increment in nitrogen loading and decrement in C/N ratio (Fig. 3A), attaining highest  
523 values of 9.8 g L<sup>-1</sup> and 1.23 g L<sup>-1</sup> d<sup>-1</sup>, respectively, at a N-loading of 1.38 g L<sup>-1</sup> (C/N=2.6). These trends  
524 are in alignment with the findings of Bakratsas et al.<sup>23</sup>, where increasing production of *P. ostreatus*  
525 biomass was achieved with increase in the concentration of yeast extract from 5 to 15 g L<sup>-1</sup>, which can  
526 be indirectly attributed to increase in nitrogen loading. Biomass yield was remarkably low (0.4 g g<sup>-1</sup>) at  
527 the lowest nitrogen loading (highest C/N ratio, *i.e.*, 65.5), however, comparable yields in the range of  
528 0.88 – 1.09 g g<sup>-1</sup> were obtained with further increment in nitrogen loading from 0.39 to 1.38 g L<sup>-1</sup> (Fig.  
529 3B). Cao et al.<sup>32</sup> reported similar observations in *Aspergillus awamori*, wherein, a high C/N ratio of  
530 153 (control) resulted in the lowest production (~5 g L<sup>-1</sup>) and yield (~0.5 g g<sup>-1</sup>) of biomass, however, a  
531 reduction in C/N ratio to 15 led to substantial increment in the values (14.00 g L<sup>-1</sup> and 0.98 g g<sup>-1</sup>),  
532 relative to the control.

533 The improved growth metrics achieved at lower C/N ratios could be attributed to higher  
534 nitrogen concentration in the media, which is critical for the endogenous production of nucleotides,  
535 amino acids and other supporting cofactors. Increased availability of easily assimilated organic nitrogen  
536 sources (like those in yeast extract) directly fuels pathways like the glutamine synthetase cycle, boosting  
537 the production of amino acids needed for protein synthesis and overall cell structure.<sup>45</sup> Simultaneously,  
538 ample nitrogen ensures the production of purine and pyrimidine bases for nucleotide synthesis,  
539 supporting DNA replication and RNA production required for cell division and growth.<sup>30</sup> Furthermore,  
540 many essential enzymatic cofactors (*e.g.*, nicotinamide adenine dinucleotide and flavin adenine  
541 dinucleotide) also contain nitrogen atoms. With a preferable carbon substrate such as glucose providing  
542 energy and carbon skeletons, abundant nitrogen allows these precursors to be readily assimilated into  
543 biomass, resulting in higher biomass titers. Conversely, a high C/N ratio leads to faster nitrogen  
544 depletion, and subsequent nitrogen starvation causing earlier transition of the fungi into stationary  
545 phase, and consequent reduction in biomass growth and yield.<sup>32</sup>

546 In the next step, *P. ostreatus* was evaluated under varying C/P ratios of 52.9, 6.9, 3.7 and 2.5,  
547 which were obtained by supplementing different loadings of phosphorus to the media, *viz.*, 0.07, 0.52,  
548 0.98 and 1.43 g L<sup>-1</sup> (corresponding to KH<sub>2</sub>PO<sub>4</sub> concentrations of 0.3, 2.3, 4.3 and 6.3 g L<sup>-1</sup>), in addition  
549 to the control unit without phosphorus (0 g L<sup>-1</sup>). The highest biomass production (4 g L<sup>-1</sup>) and  
550 productivity (0.5 g L<sup>-1</sup> d<sup>-1</sup>) were achieved at the highest C/P ratio of 52.9 (corresponding to a



551 phosphorous loading of 0.07 g L<sup>-1</sup>), which were significantly higher than the control ( $p < 0.05$ ) but  
 552 were comparable with the performance at lower C/P ratios or higher phosphorous loadings (Fig. 3D).  
 553 Biomass yield was obtained in the range of 0.9 – 1.04 g L<sup>-1</sup> d<sup>-1</sup>, clearly demonstrating that variation of  
 554 phosphorous loading did not impose any impact on the yield (Fig. 3E). Liang et al.<sup>28</sup> reported a similar  
 555 pattern in *Phanerochaete chrysosporium* involving increase in biomass titer from 0.002 to 2.006 g L<sup>-1</sup>  
 556 with initial increase in KH<sub>2</sub>PO<sub>4</sub> concentration from 0 to 0.05 g L<sup>-1</sup>, however, further increment in  
 557 KH<sub>2</sub>PO<sub>4</sub> loading (0.5 – 4.0 g L<sup>-1</sup>), led to reduced growth rate producing only 2.397 – 2.678 g L<sup>-1</sup> of  
 558 biomass, which was not significantly different from the biomass titer attained at lower KH<sub>2</sub>PO<sub>4</sub> loading  
 559 of 0.05 g L<sup>-1</sup>. Hence, it can be inferred that although availability of phosphorus is critical for boosting  
 560 the growth of fungi, excess availability of phosphorus beyond a certain threshold does not impose  
 561 substantial impact on biomass production.

#### 562 3.2.4. Effect of C/N and C/P ratios on mycelium protein

563 Similarly, the increase in nitrogen loading and corresponding reduction in C/N ratio (from 65.5  
 564 to 2.6) led to significant increase in protein content from 1.84% to 39.7%, and equivalent protein  
 565 production from 0.01 to 3.89 g L<sup>-1</sup> (Fig. 3C). There is lack of studies delving into the effect of C/N ratio  
 566 on protein production in submerged fermentation of filamentous fungi. However, the effect is well-  
 567 studied in other proteinaceous cell factories. For instance, microalgae *Chlorella vulgaris* exhibited  
 568 similar pattern in enhancement of protein content from 27.61% to 61.56% with the decrease in C/N  
 569 ratio from 32:1 to 12:1, which has been primarily attributed to higher availability of nitrogen to the  
 570 intracellular protein synthesis machinery at lower C/N ratio.<sup>46</sup> Likewise, as the C/N ratio declined from  
 571 50 to 24, single-cell protein production was observed to increase from 3.7 to 8.4 g L<sup>-1</sup> in red yeast  
 572 *Sporidiobolus pararoseus*.<sup>47</sup> A lower C/N ratio, associated with a higher nitrogen availability stimulates  
 573 the cells to build more proteins primarily to support cell growth<sup>47</sup>, which possibly explains the reason  
 574 for higher protein content associated with higher biomass production in the present study. Moreover,  
 575 the highest protein content attained in present study (39.7%) aligns well with the range (42.8 – 44.0%)  
 576 reported for *Pleurotus ostreatus* cultivated using synthetic media as well waste substrates, e.g., wine  
 577 lees and fibre sludge-derived cellulosic hydrolysate.<sup>10,23</sup>

578 The maximum protein content (24.0%) and protein production (0.96 g L<sup>-1</sup>), achieved at a  
 579 phosphorous loading of 0.07 g L<sup>-1</sup> (equivalent C/P ratio of 52.9), exhibited increments of 10.2% and  
 580 52.1%, respectively, relative to the control without phosphorus (Fig. 3F). However, further increase in  
 581 phosphorous loading did not cause any significant changes in the protein content and production.  
 582 Similar observations were reported in *Mucor indicus*, where variation in KH<sub>2</sub>PO<sub>4</sub> concentration between  
 583 0 and 7.5 g L<sup>-1</sup> did not cause any significant changes in the protein content which was secreted within  
 584 a range of 505 to 520 g kg<sup>-1</sup> biomass under different conditions.<sup>29</sup> In the present study, the plateauing of  
 585 biomass production at higher phosphorous loadings beyond 0.07 g L<sup>-1</sup>, could be attributed to feedback

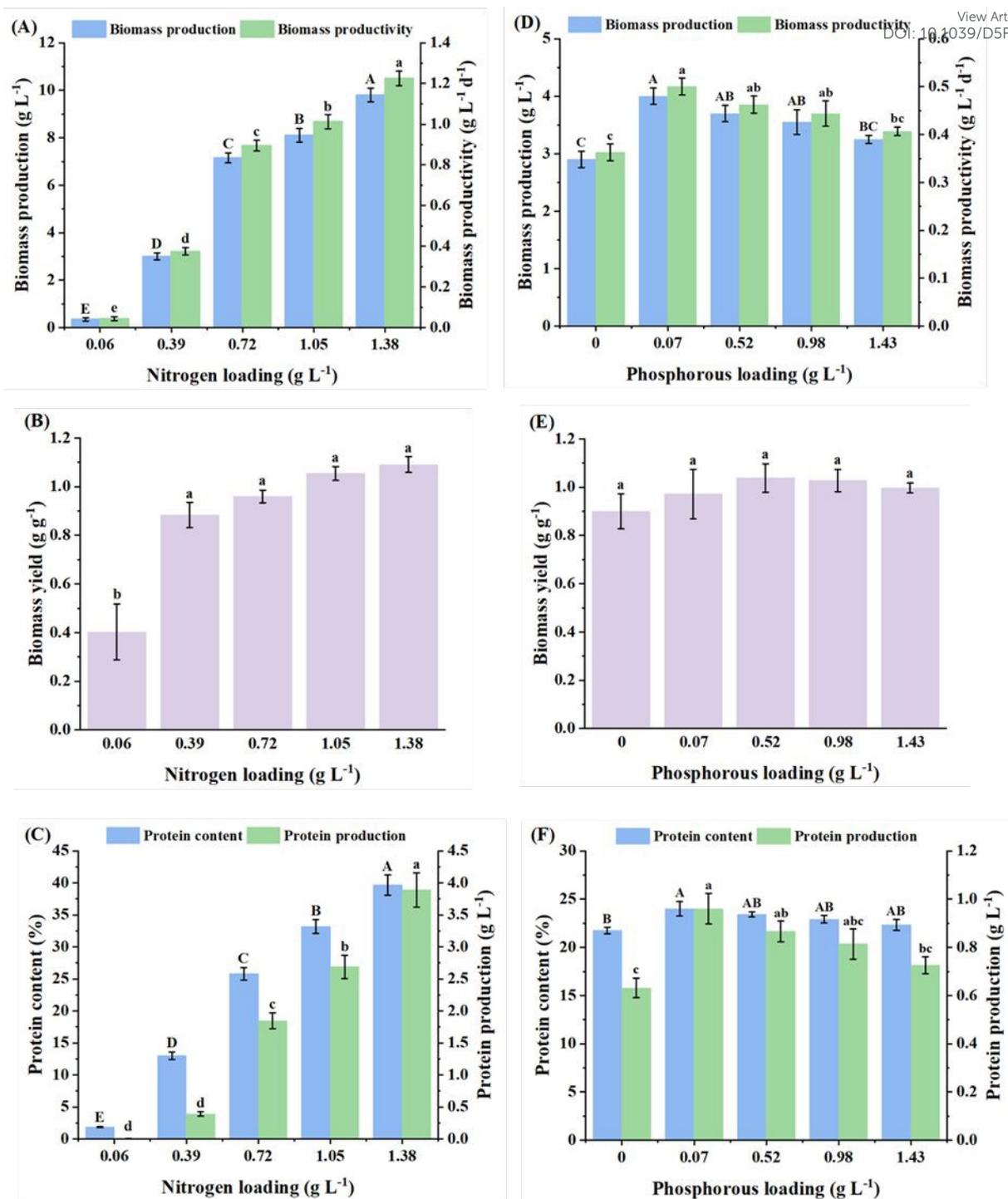


586 inhibition, implying that at 0.07 g L<sup>-1</sup>, the fungal cells were able to fill their internal phosphorus  
587 reservoirs (*e.g.*, ATP, polyphosphate granules), thereby, signaling the transporters to slow down or stop  
588 importing more phosphorus.<sup>48</sup> Beyond this concentration, even with abundant external phosphorus, the  
589 cell's regulatory systems might stop taking it up, thus plateauing the growth. Furthermore, since the  
590 accumulation of protein is associated with growth as previously reported for other species, *e.g.*,  
591 *Chlorella pyrenoidosa*<sup>49</sup>, likewise in the present study, saturation in protein content and production  
592 were observed with plateauing of biomass production beyond 0.07 g L<sup>-1</sup> phosphorous loading.

593 To the best knowledge of the authors, this is the first study demonstrating the effect of C/P ratio  
594 on the production of biomass and protein in submerged fermentation of filamentous fungi. However,  
595 the effects of C/P ratio observed in the present study are comparable with the outcomes of other studies  
596 concerning non-fungi-based cell systems. For instance, variation of C/P ratio between 282 to 474 did  
597 not cause a significant change in biomass production, showing saturation within a range of 48.1 – 48.9  
598 g L<sup>-1</sup> in the yeast *Sporidiobolus pararoseus*.<sup>47</sup> Based on the present findings, nitrogen loading of 1.38  
599 g L<sup>-1</sup> and phosphorous loading of 0.07 g L<sup>-1</sup>, corresponding to C/N and C/P ratios of 2.6 and 52.9,  
600 respectively, were selected for further experiments. Variation of C/N and C/P ratio did not impose a  
601 significant impact on the morphology of the mycelium pellets in terms of size distribution pattern and  
602 external appearance of the surface (data not shown). In general, these pellets exhibited the same  
603 morphological attributes as those cultured using glucose at a carbon loading of 3.6 g L<sup>-1</sup> (Fig. 5B (i)).  
604 This implies that pellet morphology remains consistent under fixed sources of carbon and nitrogen,  
605 unless other fermentation parameters are altered (*e.g.*, inoculum fragmentation, inoculum density and  
606 agitation).

607





608

609 **Fig. 3.** Effect of varying nitrogen loadings on (A) biomass production and productivity, (B) biomass610 yield, and (C) protein content and protein production in *Pleurotus ostreatus*. Different nitrogen611 loadings (0.06 – 1.38 g L<sup>-1</sup>) correspond to equivalent C/N ratios of 65.5, 9.4, 5.0, 3.4 and 2.6,

612 respectively. Effect of varying phosphorous loadings on (D) biomass production and productivity, (E)

613 biomass yield, and (F) protein content and protein production in *Pleurotus ostreatus*. Different614 phosphorous loadings (0.07 – 1.43 g L<sup>-1</sup>) correspond to equivalent C/P ratios of 52.9, 6.9, 3.7 and 2.5,

615 respectively. \* Groups that share same letters are considered statistically insignificant.



### 616 3.3. Effect of inoculum density and agitation rate

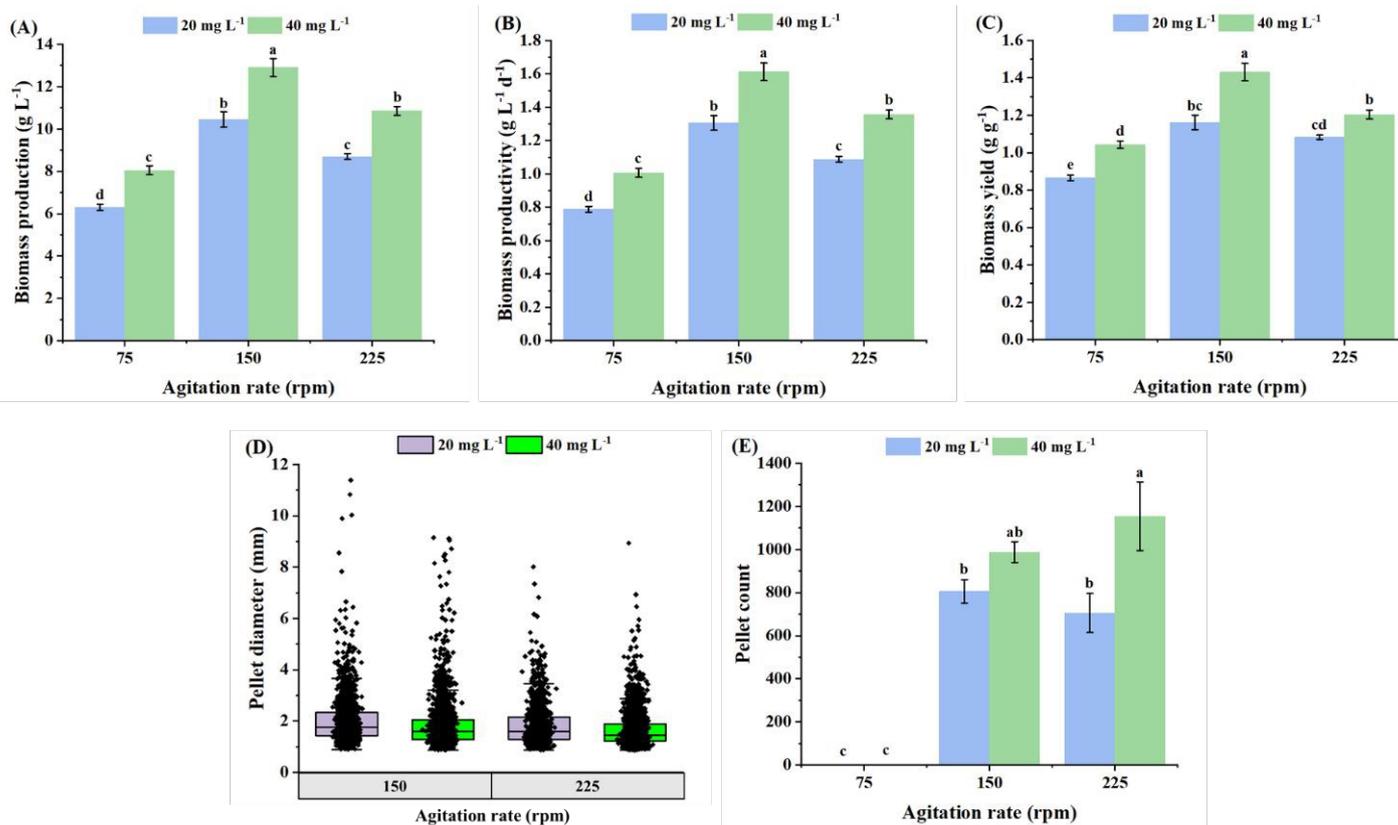
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617 Inoculum density and agitation rate were observed to have significant effects on growth as well  
618 as the morphology of pellets in terms of size and number. Biomass production, productivity and yield  
619 were observed to increase with increase in inoculum density from 20 to 40 mg L<sup>-1</sup> under different  
620 agitation rates (Fig. 4A, 4B and 4C). Although these growth parameters exhibited significant  
621 improvement with the increase in agitation rate from 75 to 150 rpm, further increment in agitation to  
622 225 rpm, resulted in substantial decline in biomass growth in terms of titer, productivity and yield (Fig.  
623 4A, 4B and 4C). Cao et al.<sup>32</sup> reported similar observations in *Aspergillus awamori* culture where  
624 maximum biomass concentration was attained at 150 rpm, however, the production was observed to  
625 decline under both lower (50 rpm) as well as higher (250 rpm) agitation rates. The decline in biomass  
626 growth at higher agitation rate has been attributed to the fragmentation of pellets induced by high  
627 hydrodynamic shear. The broken pellets subsequently attempt to restructure into new pellets; the  
628 underlying process consumes energy, thereby diverting it away from active biomass formation and  
629 consequent decline in growth.<sup>32</sup> In the present study, highest biomass production (12.9 g L<sup>-1</sup>),  
630 productivity (1.61 g L<sup>-1</sup> d<sup>-1</sup>) and yield (1.43 g g<sup>-1</sup>) were attained using an inoculum density of 40 mg L<sup>-1</sup>  
631 at 150 rpm; which were 23.4, 23.4 and 23.1% higher than the respective values at lower inoculum  
632 density (20 mg L<sup>-1</sup>) under the same agitation rate. These observations are in alignment with the study  
633 reported by Shen et al.<sup>35</sup>, where biomass production was observed to increase with the increment in  
634 inoculum density (from 8 to 40 mg L<sup>-1</sup>), resulting in a maximum titer of 10.22 g L<sup>-1</sup> at 40 mg L<sup>-1</sup>  
635 inoculum concentration in *Inonotus hispidus* fermentation.

636 The lowest agitation rate of 75 rpm resulted in single, large-sized mycelium pellets with average  
637 sizes of 63.8 mm and 72.7 mm at inoculum densities of 20 and 40 mg L<sup>-1</sup>, respectively (Fig. 5C). Fig.  
638 5C shows the morphology of the pellets harvested on Day 8. The pellet sizes reduced and the pellet  
639 count increased significantly with the increase in agitation to 150 rpm. Whereas, further increment in  
640 agitation rate (from 150 to 225 rpm) did not impose a significant difference on the size nor the number  
641 of the pellets under specific inoculum densities (Fig. 4D, 4E and 5C). The average pellet sizes obtained  
642 at 150 rpm were 2.02 mm (20 mg L<sup>-1</sup>), 1.85 mm (40 mg L<sup>-1</sup>), and at 225 rpm were 1.83 mm (20 mg L<sup>-1</sup>)  
643 and 1.66 mm (40 mg L<sup>-1</sup>), as shown in the Box and Whisker plot in Fig. 4D. A few other studies have  
644 reported similar observations in terms of reduction in pellet size with increase in agitation rate. For  
645 instance, average pellet diameter of *Neurospora intermedia* was observed to reduce from 6.54 ± 0.62  
646 mm to 1.92 ± 0.33 mm, with the increase in agitation from 100 to 150 rpm.<sup>22</sup> Although inoculum density  
647 did not have a remarkable effect on the size of the pellets at higher agitation rates (150 – 225 rpm), the  
648 pellet number was observed to increase significantly ( $p < 0.05$ ) with the increase in the inoculum density  
649 at the highest agitation rate of 225 rpm, *i.e.*, 705 pellets at 20 mg L<sup>-1</sup> to 1154 pellets at 40 mg L<sup>-1</sup> (Fig.  
650 4D and 4E).



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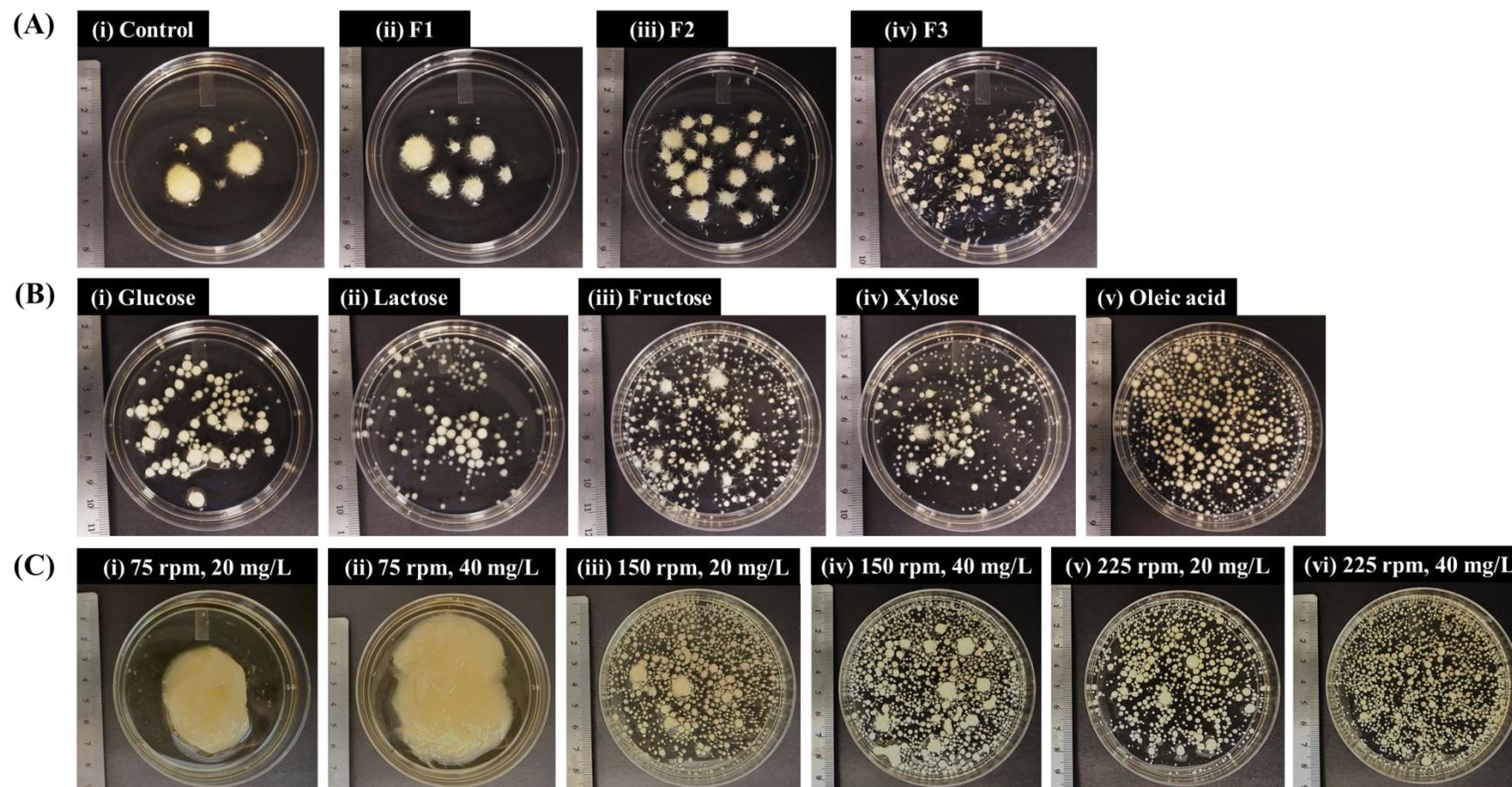
652

653 **Fig. 4.** Effect of different agitation rates and inoculum densities on (A) biomass production, (B)  
 654 biomass productivity, (C) biomass yield, and (D) diameter of mycelium pellets on Day 8 (Quartile  
 655 summary of the Box and Whisker plot is provided in Supplementary information, Table S2). Only a  
 656 single pellet was observed at 75 rpm with a mean diameter of 63.8 mm and 72.7 mm at inoculum  
 657 densities of 20 and 40 mg L<sup>-1</sup>, respectively, and therefore, has not been included in the Box and  
 658 Whisker plot, and (E) number of pellets in 10 mL culture. Inoculum densities are on dry weight basis.

659 \* Groups that share same letters are considered statistically insignificant.

660





661

662 **Fig. 5.** Morphology of mycelium pellets under different (A) fragmentation levels. Control: no homogenization; F1: 3000 rpm, 5 s; F2: 5000 rpm, 10 s; and F3:663 5000 rpm, 40 s; (B) carbon sources at 3.6 g L<sup>-1</sup> carbon loading harvested on Day 8; and (C) inoculum densities and agitation rates.

664 From a technological perspective, controlling *P. ostreatus* pellet morphology via inoculum  
665 fragmentation, inoculum density, agitation, and media formulation can profoundly impact industrial  
666 downstream processing. In the present study, compact, uniform pellets, achieved using fragmentation  
667 level F3, inoculum density of 20 mg L<sup>-1</sup>, and agitation rate of 150 rpm, are significantly more  
668 advantageous for industrial-scale operations than large, loose, and heterogeneous mycelial clumps. The  
669 production of compact, uniform pellets is further supported by media components like glucose and oleic  
670 acid, which facilitate rounder pellet formation. In terms of primary recovery, such uniform pellet  
671 characteristics can facilitate more predictable filtration and dewatering rates. The compact nature of  
672 these pellets, combined with their smoother surface topography, reduces the entrapment of interstitial  
673 water within the fungal matrix, which can enhance drying efficiency and reduce the energy footprint of  
674 moisture removal. Beyond harvesting, compact pellets provide a higher biomass density (as observed  
675 in Sections 3.1 and 3.3), that can improve the efficiency of mechanical cell disruption, ensuring a more  
676 consistent release of intracellular proteins. Furthermore, a standardized size distribution is critical for  
677 texturization processes like extrusion, where uniform raw material ensures consistent rheological  
678 behavior and heat transfer.<sup>50</sup> This morphological control, governed by both physical and chemical  
679 parameters, is therefore essential for ensuring consistent quality and streamlining the transition from  
680 fungal biomass to a structured food ingredient.

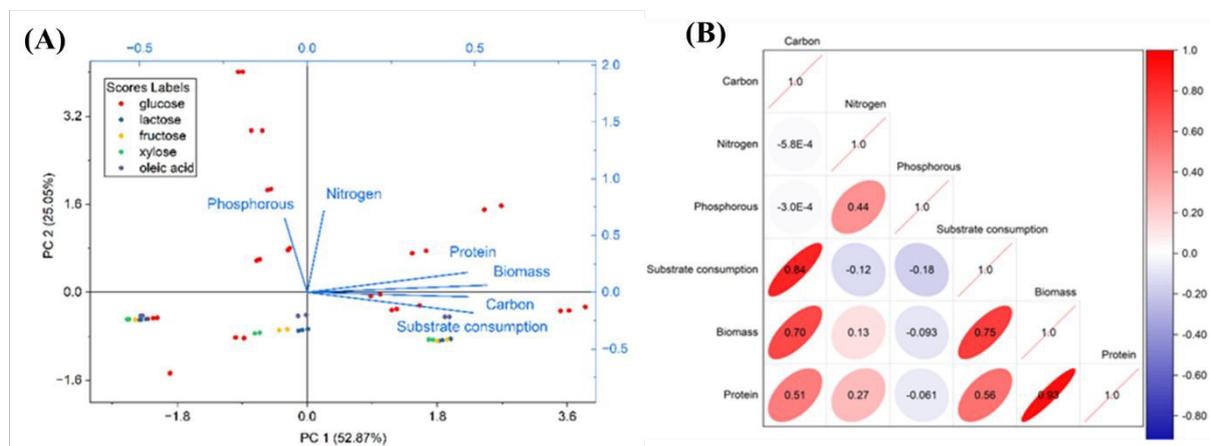
### 681 3.4. Principal Component Analysis

682 Principal Component Analysis (PCA) was conducted to study how submerged fermentation  
683 parameters influence biomass and protein production, and substrate consumption. Essentially it reduces  
684 the dimension of variables from the primary data into principal components and visualizes how much  
685 variance of each experimental factor was weighted. It was found that two major principal components,  
686 PC1 and PC2, describe 77.92% of the total variation; and a third principal component, PC3, provides  
687 an additional 12.7% of the variation (as shown in Supplementary information, Fig. S5). The loading  
688 plot (Fig. 6A) demonstrates the relative weight of each variable in PC1 and PC2, where all factors  
689 except for phosphorous loading contribute positively to PC1, and carbon loading and substrate  
690 consumption contribute negatively to PC2. The dots in score plot represent the coordinates of individual  
691 data points in the new principal component space, with different carbon sources being distinguished by  
692 color. The results suggest that the majority of observations with glucose as carbon source contribute  
693 positively to PC2, while observations with other carbon sources including lactose, fructose, xylose and  
694 oleic acid contribute negatively to PC2. Pearson Correlation Analysis (Fig. 6B) was performed to  
695 understand the correlations among all variables. The analysis suggests that among three key medium  
696 constituents including carbon, nitrogen and phosphorous loadings, carbon loading had a significantly  
697 positive correlation with production of both biomass and protein, and substrate consumption.  
698 Meanwhile, nitrogen loading only had a strong positive correlation with protein production, rather than



699 biomass. Phosphorous loading did not have any significant correlation with production or substrate  
 700 consumption. These results are in good agreement with our previous findings.

701



702  
 703 **Fig. 6.** (A) Biplot of Principal Component Analysis; and (B) Pearson Correlation Matrix on the  
 704 medium composition, carbon substrate consumption, biomass and protein production from submerged  
 705 fermentation of *P. ostreatus*

#### 706 4. Conclusions

707  
 708 The study herein developed a robust framework for enhancing mycelium-based protein  
 709 production from *Pleurotus ostreatus* in submerged fermentation. The results highlight that precise  
 710 control of mycelial morphology can be achieved under specific inoculum fragmentation, agitation and  
 711 inoculation density. The media composition influences protein production, where a low C/N ratio (2.6)  
 712 and a high C/P ratio (52.9) provided the highest protein content of mycelium (39.7% of biomass, d.b.).  
 713 By integrating these findings with appropriate agitation and inoculum density, the process achieved a  
 714 maximum biomass titer of 12.9 g L<sup>-1</sup>, with a productivity of 1.61 g L<sup>-1</sup> d<sup>-1</sup>. Furthermore, statistical  
 715 analysis via PCA confirmed that carbon loading strongly influenced both biomass and protein  
 716 production, whereas nitrogen loading was a significant driver primarily for protein synthesis, and  
 717 phosphorous loading showed no significant correlation with either metric. This outlines a potential  
 718 strategy for leveraging carbon and nitrogen-rich waste streams to optimize sustainable co-production  
 719 of mycelium biomass and protein at industrial-scale. Moreover, this work establishes oleic acid as a  
 720 viable non-sugar carbon source, demonstrating the potential to improve the economic feasibility of  
 721 large-scale production by valorizing lipid-rich waste streams (e.g., used cooking oils or agricultural  
 722 byproducts), thus avoiding competition with arable land and food crops required for conventional sugar  
 723 feedstocks. This research advances the development of mycelium as a sustainable, food-grade vegan



724 protein, offering a promising alternative to conventional animal-sourced protein. However, industrial  
 725 scale submerged fungal cultivation could be constrained by high broth viscosity, limited oxygen  
 726 transfer, and pellet clumping that can obstruct/adhere to bioreactor components like baffles, leading to  
 727 process inconsistencies. Future research must address these scale-up dynamics by managing rheological  
 728 behavior and maintaining morphological stability in large-scale bioreactors. Concurrently, further  
 729 exploration can be directed towards deciphering the molecular-level mechanisms governing the variable  
 730 performance of the species in response to media formulation.

731

### 732 **Author contributions**

733 **Krishna Kalyani Sahoo:** Investigation, Visualization, Formal analysis, Writing - Original Draft,  
 734 Writing - Review & Editing. **Bruno Meireles Xavier:** Resources, Supervision. **Mohammad Afiq**  
 735 **Hafiy Mohammad Taufiq:** Investigation, Writing - Original Draft. **Ke Wang:** Conceptualization,  
 736 Supervision, Funding acquisition, Formal analysis, Writing - Original Draft, Writing - Review &  
 737 Editing.

### 738 **Conflicts of interest**

739 There are no conflicts to declare.

### 740 **Data availability**

741 The data supporting this article have been included as part of the Supplementary Information.

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749

### **References**

- 750 1. K. Smith, A. W. Watson, M. Lonnie, W. M. Peeters, D. Ooninx, N. Tsoutsoura, G. Simon-  
 751 Miquel, K. Szepe, N. Cochetel, A. G. Pearson, O. C. Witard, A. M. Salter, M. Bennett and B.  
 752 M. Corfe, *European Journal of Nutrition*, 2024, **63**, 1425-1433.
- 753 2. U.S. Department of Agriculture, [https://www.usda.gov/about-usda/news/press-](https://www.usda.gov/about-usda/news/press-releases/2022/06/01/usda-announces-framework-shoring-food-supply-chain-and-transforming-food-system-be-fairer-more)  
 754 [releases/2022/06/01/usda-announces-framework-shoring-food-supply-chain-and-](https://www.usda.gov/about-usda/news/press-releases/2022/06/01/usda-announces-framework-shoring-food-supply-chain-and-transforming-food-system-be-fairer-more)  
 755 [transforming-food-system-be-fairer-more](https://www.usda.gov/about-usda/news/press-releases/2022/06/01/usda-announces-framework-shoring-food-supply-chain-and-transforming-food-system-be-fairer-more), (accessed October 2025).
- 756 3. E. J. Derbyshire and J. Delange, *Frontiers in Sustainable Food Systems*, 2021, **5**.
- 757 4. K. K. Sahoo, S. Hao, J. M. I. Aquino and K. Wang, *Trends in Food Science & Technology*,  
 758 2026, DOI: 10.1016/j.tifs.2026.105614.



- 759 5. M. E. Effiong, C. P. Umeokwochi, I. S. Afolabi and S. N. Chinedu, *Frontiers in Nutrition*, 2024, **10**. View Article Online  
DOI: 10.1039/D3FB00752F
- 760
- 761 6. A. Irshad, A. Tahir, S. Sharif, A. Khalid, S. Ali, A. Naz, H. Sadia, A. Ameen and D. P. Fulzele,
- 762 *BioMed Research International*, 2023, **2023**.
- 763 7. G. Bakratsas, C. Tsoumanis, H. Stamatis and P. Katapodis, *Fermentation*, 2024, **10**.
- 764 8. F. Ahmad Zakil, L. H. Xuan, N. Zaman, N. I. Alan, N. A. A. Salahutheen, M. S. M. Sueb and
- 765 R. Isha, *Bioresource Technology Reports*, 2022, **17**.
- 766 9. D. Tagkouli, A. Kaliora, G. Bekiaris, G. Koutrotsios, M. Christea, G. I. Zervakis and N.
- 767 Kalogeropoulos, *Molecules*, 2020, **25**.
- 768 10. G. Bakratsas, K. Antoniadis, P. E. Athanasiou, P. Katapodis and H. Stamatis, *Biomass*, 2023,
- 769 **4**, 1-22.
- 770 11. O. Datsomor, Q. Yan, L. Opoku-Mensah, G. Zhao and L. Miao, *Fermentation*, 2022, **8**.
- 771 12. R. Rajput, B. Singh and M. Bashir Mir, in *Beta-Glucan: Sources, Properties and Applications*,
- 772 2025, DOI: 10.1007/978-3-031-95788-8\_11, ch. Chapter 11, pp. 189-205.
- 773 13. H. El-Gendi, A. K. Saleh, R. Badierah, E. M. Redwan, Y. A. El-Maradny and E. M. El-
- 774 Fakharany, *Journal of Fungi*, 2021, **8**.
- 775 14. F. Humpenöder, B. L. Bodirsky, I. Weindl, H. Lotze-Campen, T. Linder and A. Popp, *Nature*,
- 776 2022, **605**, 90-96.
- 777 15. K. Majumder, B. Paul and R. Sundas, *Journal of Genetic Engineering and Biotechnology*, 2020,
- 778 **18**.
- 779 16. A. Musatti, E. Ficara, C. Mapelli, C. Sambusiti and M. Rollini, *Journal of Environmental*
- 780 *Management*, 2017, **199**, 1-6.
- 781 17. V. Elisashvili, M. Penninckx, E. Kachlishvili, N. Tsiklauri, E. Metreveli, T. Kharziani and G.
- 782 Kvesitadze, *Bioresource Technology*, 2008, **99**, 457-462.
- 783 18. M. M. Pascual, L. T. Herbert, M. Campos, V. Jurski, J. C. Paineofilú and C. M. Luquet,
- 784 *Innovative Food Science & Emerging Technologies*, 2025, **102**.
- 785 19. A. Hamza, M. P. Shankar, U. S. Chowdary, S. Ghanekar, S. Sahoo, C. V. Krishnaiah and D. S.
- 786 Kumar, *Food and Humanity*, 2024, **2**.
- 787 20. G. M. Mascarin, P. S. Golo, C. de Souza Ribeiro-Silva, E. R. Muniz, A. de Oliveira Franco, N.
- 788 N. Kobori and É. K. K. Fernandes, *Applied Microbiology and Biotechnology*, 2024, **108**.
- 789 21. X. Zhang, H. Liu, M. Zhang, W. Chen and C. Wang, *Journal of Fungi*, 2023, **9**.
- 790 22. R. B. Nair, P. R. Lennartsson and M. J. Taherzadeh, *AMB Express*, 2016, **6**.
- 791 23. G. Bakratsas, A. Polydera, O. Nilson, L. Kossatz, C. Xiros, P. Katapodis and H. Stamatis,
- 792 *Sustainable Food Technology*, 2023, **1**, 377-389.
- 793 24. X.-L. Li, X.-N. Qi, J.-C. Deng, P. Jiang, S.-Y. Wang, X.-L. Xue, Q.-H. Wang and X. Ren,
- 794 *Foods*, 2025, **14**.
- 795 25. V. Maini Rekdal, J. M. Villalobos-Escobedo, N. Rodriguez-Valeron, M. Olaizola Garcia, D.
- 796 Prado Vásquez, A. Rosales, P. M. Sørensen, E. E. K. Baidoo, A. Calheiros de Carvalho, R.
- 797 Riley, A. Lipzen, G. He, M. Yan, S. Haridas, C. Daum, Y. Yoshinaga, V. Ng, I. V. Grigoriev,
- 798 R. Munk, C. H. Wijaya, L. Nuraida, I. Damayanti, P. Cruz-Morales and J. D. Keasling, *Nature*
- 799 *Microbiology*, 2024, **9**, 2666-2683.
- 800 26. M. T. Nazir, A. M. Soufiani, J. A. Ferreira, T. Sar and M. J. Taherzadeh, *Journal of Chemical*
- 801 *Technology & Biotechnology*, 2022, **97**, 2626-2635.
- 802 27. S. Dzurendova, B. Zimmermann, V. Tafintseva, A. Kohler, D. Ekeberg and V. Shapaval,
- 803 *Applied Microbiology and Biotechnology*, 2020, **104**, 8065-8076.
- 804 28. H. Liang, D.-W. Gao and Y.-G. Zeng, *Bioresource Technology*, 2012, **107**, 535-538.
- 805 29. M. Mohammadi, A. Zamani and K. Karimi, *Applied Biochemistry and Biotechnology*, 2013,
- 806 **171**, 1465-1472.
- 807 30. Z. Wang, S. Li, H. Pan, Y. Li, X. Wang, H. Zhou and J. Shan, *Journal of Microbiological*
- 808 *Methods*, 2025, **238**.
- 809 31. C. Rosales-López, A. Vargas-López, M. Monge-Artavia and M. Rojas-Chaves,
- 810 *Microorganisms*, 2022, **10**.
- 811 32. L. Cao, H. M. El Mashad, Z. Pan and R. Zhang, *Food and Bioprocess Technology*, 2025, **18**,
- 812 8735-8750.
- 813 33. G. E. Zharare, S. M. Kabanda and J. Z. Poku, *Scientia Horticulturae*, 2010, **125**, 95-102.



- 814 34. H. Jeon, H. Son and K. Min, *Bio Protoc*, 2023, **13**, e4889.
- 815 35. K. Shen, Y. Liu, L. Liu, A. W. Khan, N. Normakhamatov and Z. Wang, *Applied Biochemistry and Biotechnology*, 2024, **197**, 1534-1555.
- 816
- 817 36. G. L. Miller, *Analytical chemistry*, 1959, **31**, 426-428.
- 818 37. B. Wang, J. Chen, H. Li, F. Sun, Y. Li and G. Shi, *Bioprocess and Biosystems Engineering*, 2016, **40**, 45-53.
- 819
- 820 38. C. Falter and S. Reumann, *Molecular Plant Pathology*, 2022, **23**, 781-794.
- 821 39. M. L. Hernández, M. D. Sicardo, A. Belaj and J. M. Martínez-Rivas, *Frontiers in Plant Science*, 2021, **12**.
- 822
- 823 40. J. Shen, Y. Wang, P. Fan, L. Jiang, W. Zhuang, Y. Han and H. Zhang, *Colloids and Surfaces A: Physicochemical and Engineering Aspects*, 2019, **568**, 66-74.
- 824
- 825 41. S. Nakagame, H. Minagawa and N. Motegi, *Applied Biochemistry and Biotechnology*, 2022, **195**, 1085-1095.
- 826
- 827 42. F. Gallotti, C. Turchiuli and V. Lavelli, *Journal of Food Process Engineering*, 2021, **45**.
- 828 43. J. Li, L. Lin, H. Li, C. Tian and Y. Ma, *Biotechnology for Biofuels*, 2014, **7**.
- 829 44. C.-Y. Cheng, Y.-S. Wang, Z.-L. Wang and S. Bibi, *Foods*, 2023, **12**.
- 830 45. M. C. B. Tecson, C. Geluz, Y. Cruz and E. R. Greene, *Biochemistry*, 2025, **64**, 547-554.
- 831 46. Y. Cai, L. Zhai, X. Fang, K. Wu, Y. Liu, X. Cui, Y. Wang, Z. Yu, R. Ruan, T. Liu and Q. Zhang, *Biotechnology for Biofuels and Bioproducts*, 2022, **15**.
- 832
- 833 47. H. Wang, B. Hu, J. Liu, H. Qian, J. Xu and W. Zhang, *Bioprocess and Biosystems Engineering*, 2020, **43**, 1403-1414.
- 834
- 835 48. M. Takado, T. Komamura, T. Nishimura, I. Ohkubo, K. Ohuchi, T. Matsumoto and K. Takeda, *Journal of Biological Chemistry*, 2023, **299**.
- 836
- 837 49. D. Xing, X. Li, Y. Wang, S. Deng, C. Jin, Y. Zhao and L. Guo, *Journal of Water Process Engineering*, 2023, **51**.
- 838
- 839 50. T. A. L. Do, J. M. Hargreaves, B. Wolf, J. Hort and J. R. Mitchell, *Journal of Food Science*, 2007, **72**.
- 840
- 841

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**Data availability**

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The data supporting this article have been included as part of the Supplementary Information.

