Bacteriolyis by *Agaricus bisporus*

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*Agaricus bisporus*, the cultivated mushroom, was able to mineralize dead 14C-labelled *Bacillus subtilis* and utilize the cellular components as sole source of carbon and nitrogen for growth. Consistently higher levels of bacteriolytic activity were obtained when *A. bisporus* was grown on lower concentrations of bacteria, 1-5 mg ml<sup>-1</sup> as opposed to 3 mg ml<sup>-1</sup> basal liquid medium. *A. bisporus* also mineralized 14C-labelled bacteria in the presence of readily available alternative carbon and nitrogen sources such as glucose and ammonium sulphate. 14C-labelled bacteria were degraded to 14CO<sub>2</sub> more efficiently than [14C]cellulose. *A. bisporus* was able efficiently to mineralize biomass produced in situ (> 20% of total label released as 14CO<sub>2</sub>) where label was incorporated into the microbial biomass by composting rather than added as killed bacteria. This is the first time that quantitative data have been produced to show that the microbial biomass in wheat straw/animal manure compost might be significant in the nutritional strategy of basidiomycetes.

**Introduction**

The basic nutritional requirements *in vitro* of *Agaricus bisporus* (Lange) Imbach, the cultivated edible mushroom, are known (Wood & Fermor, 1985). Its commercial production medium, composted wheat straw, is prepared such that it is selective for mushroom mycelial growth. Successful mycelial colonization and subsequent degradation of this medium by *A. bisporus* is dependent on several ecophysiological factors (Wood & Smith, 1987), but these factors do not completely explain how selectivity can be described in chemical or biological terms (Straatsma et al., 1989).

The linear growth rate of *A. bisporus* in sterilized (autoclaved or irradiated) compost is half of that in unsterilized compost (Fermor et al., 1985). It is apparent, therefore, that micro-organisms or their metabolic products present in compost may be involved in stimulation of mushroom mycelial growth. Electron microscopy has revealed that during composting an amorphous coating builds up on straw in the compost mix (Atkey & Wood, 1983). This coating contains carbohydrate and phenolic fractions of microbial origin and contains abundant microbial cells, mycelium, spores and debris (Wain, 1981). Some 50% of this straw coating disappears within one month of the compost being inoculated with *A. bisporus* (Wain, 1981). These observations led to the hypothesis that the microbial biomass accumulated during composting may act as a concentrat-ed nutrient source for the mushroom mycelium as it colonizes the compost. *A. bisporus* has been shown to degrade and efficiently utilize the biomass of killed bacteria in both solid and liquid media (Fermor & Wood, 1981), and also to degrade and utilize fungal and actinomycete mycelium (Fermor & Grant, 1985). The types of lytic enzymes produced by *A. bisporus* which are responsible for this microbial degradation have been analysed (Fermor, 1983; Grant et al., 1984, 1986). They include β-N-acetylmuramidases and β-N-acetylglucosaminidases.

A range of actively growing bacteria stimulate mushroom mycelial extension on compost malt agar medium (Rainey, 1989). Similarly, *A. bisporus*, *Coprinus quadrijidus*, *Lepista nuda* and *Pleurotus ostreatus* are able to attack and digest live microcolonies of bacteria (Barron, 1988). In co-cultures these fungi show particular sensitivity to the presence of bacterial colonies and develop specialized directional hyphae terminating in haustorium-like, absorptive hyphae which penetrate the bacterial microcolonies (Barron, 1988). Mycelial growth of *A. bisporus* on sterilized compost is greatly stimulated by pre-incubating the compost with the thermophilic fungus *Scytalidium thermophilum*, and mushroom fruit-body yield was positively correlated with the *S. thermophilum* biomass (Straatsma et al., 1989). It has been shown that the most favourable temperature for compost degradation is 50–55 °C, whilst 40–45 °C is the optimum temperature for removal of ammonia from
compost, and that thermophilic fungi are most important in this process (Ross & Harris, 1983).

The total microbial biomass in a horse manure/wheat straw compost (after 14 d composting), estimated by both direct counts and biochemical methods, was 9.2 mg carbon (g dry wt compost)\(^{-1}\) (Sparling et al., 1982). This represented approximately 2% of the compost dry weight assuming that microbial biomass contains 50% (w/w) carbon. The studies reported here were designed to determine whether A. bisporus was able to degrade and utilize live or killed microbial biomass in compost in the presence of alternative carbon and nitrogen sources. This was examined by monitoring the biodegradation of radiolabelled bacterial biomass produced in vitro or by producing radiolabelled microbial biomass in vivo by the use of a bench-scale composting unit.

**Methods**

**Organisms.** Agaricus bisporus D621 was used. This is a heterokaryotic fertile strain of commercial origin obtained from the IHR (Littlehampton) collection of basidiomycete fungi. Bacillus subtilis 168 derivative S2861 (trivial name 261) spoIA69 spoG55 rif-2 (asporogenous) was obtained from the National Institute for Medical Research, London, UK. This sporless mutant was used to avoid the practical hazards associated with growing a spore-forming bacterium in a large-scale fermenter facility.

**Basic medium for growth of B. subtilis.** A. bisporus was grown in basal Spizizen Salts Medium (SSM; Piggot, 1975) which contained: (NH\(_4\))\(_2\)SO\(_4\), 2 g; K\(_2\)HPO\(_4\), 14 g; KH\(_2\)PO\(_4\), 6 g; trisodium citrate 2H\(_2\)O, 1 g; MgSO\(_4\).7H\(_2\)O, 0.2 g in 900 ml sterile distilled water. Bacto Casamino acids (1.0 g, Difco, Technical 0231-02) containing glucose and vitamin-free Casamino acids (1, 10, 50 or 100 mg; Difco) were added to the cooled medium to give a final concentration of 0.5% (w/v). A tritiated water (1.25 pCi ml\(^{-1}\)) and added to the cooled medium to give a final concentration of 0.95% (w/v).

**Preliminary experiment. Production of \(^{14}\)C-labelled bacteria.** A preliminary experiment was undertaken to determine which concentration of Casamino acids (a readily available nutrient source) gave the most efficient uptake of \(^{14}\)C-labelled protein hydrolysate, and a satisfactory yield of microbial biomass.

**Medium.** Each Erlenmeyer flask (100 ml) contained SSM (9 ml), D-glucose (1 ml, to give a final concentration in medium of 0.5% (w/v)) and vitamin-free Casamino acids (1, 10, 50 or 100 mg; Difco). There were two flasks for each of the four concentrations of Casamino acids. \([\text{U-}^{14}\text{C}]\) Protein hydrolysate (Amersham) was added to sterile distilled water (1.25 μCi ml\(^{-1}\)) and 1 ml of this diluted hydrolysate was added to each of the eight flasks.

**Counting \(^{14}\)C uptake by B. subtilis.** After shaking, a sample of the initial medium (1 ml) was removed from each flask and placed into PCS liquid scintillation fluid (10 ml, Amersham) to count the total radioactivity in each flask prior to inoculation with bacteria. The eight flasks were then inoculated with 0.2 ml B. subtilis (10\(^{10}\) c.f.u. ml\(^{-1}\)), incubated at 37°C and shaken at 150 r.p.m. Bacterial growth in these flasks was estimated by measuring the OD\(_{600}\) of B. subtilis growing under identical conditions in four control flasks containing the same media minus \([\text{U-}^{14}\text{C}]\) protein hydrolysate. The bacteria in flasks containing 10 or 5 μg Casamino acids \(^{14}\)C were harvested after 4.75 h and those in flasks containing 1 or 0.1 mg Casamino acids \(^{14}\)C after 6.25 h.

The following counts of radioactivity were made on samples of bacterial culture from replicate flasks containing each of the four concentrations of Casamino acids. (i) Total \(^{14}\)C present in bacteria plus medium. A sample (1 ml) was placed into 10 ml PCS counting fluid. (ii) \(^{14}\)C present in bacteria. Bacteria in liquid culture (2 ml) were added to ice-cold trichloroacetic acid (2 ml, 10% (w/v)), mixed rapidly and filtered (Whatman GF/C, 2.5 cm). The bacterial precipitate was washed thoroughly with sterile distilled water and vacuum dried for 16 h at 100°C and weighed. Dried bacteria and glass fibre filter disc were then added to 10 ml PCS counting fluid. (iii) The initial filtrates (1 ml) from the cultures were also added to 10 ml PCS counting fluid. From these counts it was then possible to calculate the percentage of the initial \(^{14}\)C added which had been taken up by the B. subtilis cells.

**Growth of A. bisporus on \(^{14}\)C-labelled killed bacteria as sole carbon and nitrogen source.** \(^{14}\)C-labelled B. subtilis cells were produced and added to unlabelled B. subtilis cells. These bacterial cells were then the sole carbon and nitrogen source in the medium used to grow A. bisporus.

**Production of \(^{14}\)C-labelled B. subtilis.** The bacteria were grown in five flasks (10 ml) each containing SSM (9 ml), D-glucose (5%, w/v, 1 ml) and Casamino acids (10 mg). \([\text{U-}^{14}\text{C}]\) Protein hydrolysate (0.2 ml, 10 μCi) was diluted in sterile distilled water (4.8 ml) and 1 ml was then added to each flask. Each flask was then inoculated with 0.2 ml B. subtilis (10\(^{8}\) c.f.u. ml\(^{-1}\)). The cultures were incubated for 6 h at 37°C and shaken at 150 r.p.m. The harvested cells (OD\(_{600}\) 0.603) were then killed by suspension in 10% (w/v) trichloroacetic acid (TCA).

**Production of unlabelled B. subtilis.** A modified basic medium for growth of B. subtilis was used containing 0.5% Casamino acids, as this gave the greatest biomass yield. The medium (8 l) was autoclaved at 121°C for 40 min in a fermenter. The medium was inoculated with 100 ml B. subtilis (10\(^{8}\) c.f.u. ml\(^{-1}\), grown on basic medium) and incubated with continuous aeration (10 l min\(^{-1}\) at 37°C for 12 h). The cells (OD\(_{600}\) 0.720) were harvested by centrifugation (MSE 18, 10000 r.p.m.). The B. subtilis cell paste was then suspended in 100 ml sterile distilled water to which was added 100 ml TCA (20% w/v). A second batch culture of B. subtilis was produced under identical conditions.

**Use of the bacterial cells in growth experiments.** The harvested cells were treated as follows: (i) \(^{14}\)C-labelled B. subtilis in 50 ml TCA (10% w/v) were added to unlabelled B. subtilis in 200 ml TCA (10% w/v); (ii) the mixture was sterilized at 100°C for 30 min, then (iii) centrifuged (12000 g, 15 min) to remove most of the TCA, and (iv) washed three times in copious distilled water; (v) cells were resuspended in sterile distilled water at 15 mg ml\(^{-1}\); (vi) a sample of the suspended cells was then digested (0.2 ml in 1 ml NCS tissue solubilizer (Amersham)) at 50°C for 12 h and radioactivity counted using a Beckman LC 7800 liquid scintillation counter, which corrects for quenching by external standards.

Into each of three flasks (100 ml) were dispensed 2 ml acid-killed \(^{14}\)C-labelled B. subtilis suspension (250000 p.d.m.) and 6.6 ml basal salts; these were autoclaved for 15 min at 121°C. Basal salts (Treschow, 1944) contained: KCl, 0.2 g; MgSO\(_4\).7H\(_2\)O, 0.2 g; CaCl\(_2\), 0.2 g; FeCl\(_3\), 6H\(_2\)O, 10 mg; and trace elements H\(_2\)BO\(_3\), 0.3 mg; CuSO\(_4\).5H\(_2\)O, 0.25 mg; MnCl\(_2\).4H\(_2\)O, 2 mg; Na\(_2\)MoO\(_4\).2H\(_2\)O, 0.4 mg; ZnSO\(_4\).7H\(_2\)O, 0.2 mg; CoCl\(_2\), 6H\(_2\)O, 0.7 mg; distilled water to 659 ml. Phosphate buffers Na\(_2\)HPO\(_4\).2H\(_2\)O (750 μl; stock solution 15 g l\(^{-1}\)) and KH\(_2\)PO\(_4\) (150 μl; stock solution 9 g l\(^{-1}\)) were added to the flasks after autoclaving separately. Vitamin solution (20 mg thiamin.HCl and 2 mg biotin in 100 ml distilled water) was sterilized by membrane filtration (Millipore 0.22 μm) and 10 μl was added to each flask which was then inoculated with 0.5 ml A. bisporus D621 mycelial suspension (Fermor & Wood, 1981). Each flask was then incubated at 25°C for 56 d, and \(^{14}\)CO\(_2\) evolved was monitored.
**Bacteriolysis by Agaricus bisporus**

**Growth of A. bisporus on '14C-labelled killed bacteria with an additional carbon and nitrogen source.** Ammonium sulphate was added to cultures together with glucose or cellulose. Experiments were again conducted with 10 ml cultures in 100 ml Erlenmeyer flasks. Twelve flasks contained the following medium: basal salts (6-6 ml), 14C-labelled B. subtilis suspension in sterile distilled water (1 ml, 140000 d.p.m.), and autoclaved at 121 °C for 15 min. Sterile phosphate buffers (0-9 ml) and vitamin solution (10 ml) were then added to each flask before inoculation with 0-5 ml A. bisporus D621 mycelial suspension in sterile distilled water. The flasks were divided into three groups of four and the following supplements added to each group. (1) Distilled water (1 ml). (2) D-glucose (18.75 mg in 0-5 ml) and ammonium sulphate (10-6 mg in 0-5 ml) sterilized separately by membrane filtration (Millipore 0-22 μm). The 14C-labelled bacterial suspension (1 ml) contained 15 mg dry wt bacteria. Assuming dry B. subtilis cells contain 50% by weight carbon and 15% by weight nitrogen (Jenkinson, 1981), then the 1 ml suspension contains 7-5 mg carbon and 2-25 mg nitrogen. Additional carbon and nitrogen supplements in these four flasks were calculated to give equal C and N to that of the bacteria already in the flask. (3) Distilled water (0-5 ml), cellulose (18-75 mg, Whatman CC41 powder) and ammonium sulphate (10-6 mg in 0-5 ml). All the flasks were incubated at 25 °C for 49 d and 14CO2 evolved was monitored.

**Growth of A. bisporus on 14C-labelled glucose and cellulose.** A similar set of 12 flasks containing basal salts (6-6 ml) was set up. After autoclaving at 121 °C for 15 min, to each was added sterile phosphate buffers (0-9 ml) and vitamin solution (10 ml) prior to inoculation with 0-5 ml A. bisporus D621 mycelial suspension. The flasks were divided into three groups of four replicates and to each group was added: (1) 14C-labelled bacterial suspension (1 ml, 120000 d.p.m.) and distilled water (1 ml), or (2) unlabelled B. subtilis (15 mg), distilled water (1 ml), ammonium sulphate (10-6 mg in 0-5 ml distilled water); D-glucose (18-75 mg in 0-5 ml distilled water) and D-[U-14C]glucose (18 μl, 40000 d.p.m., Amersham), or (3) unlabelled B. subtilis (15 mg) distilled water (1 ml), ammonium sulphate (10-6 mg in 0-5 ml distilled water), cellulose (18-75 mg, Whatman CC41 powder) and diluted aqueous suspension of 14C-cellulose [0-5 ml, 44000 d.p.m. (Cellulose-2-C14, ICN Pharmaceuticals, sp. act. 3.4 μCi mg-1)]. All cultures were incubated at 25 °C for 51 d, and 14CO2 evolved was monitored.

**Degradation of killed bacteria in compost.** Twelve 100 ml Erlenmeyer flasks were filled with 7 g (fresh wt; moisture content 70%, pH 7-0) of GCRI Formula 2 compost to each was added sterile phosphate buffers (0-9 ml) and vitamin solution (10 ml) prior to inoculation with 0-5 ml A. bisporus D621 mycelial suspension. The flasks were divided into three groups of four replicates and to each group was added: (1) 14C-labelled bacterial suspension (1 ml, 120000 d.p.m.) and distilled water (1 ml), or (2) unlabelled B. subtilis (15 mg), distilled water (1 ml), ammonium sulphate (10-6 mg in 0-5 ml distilled water); D-glucose (18-75 mg in 0-5 ml distilled water) and D-[U-14C]glucose (18 μl, 40000 d.p.m., Amersham), or (3) unlabelled B. subtilis (15 mg) distilled water (1 ml), ammonium sulphate (10-6 mg in 0-5 ml distilled water), cellulose (18-75 mg, Whatman CC41 powder) and diluted aqueous suspension of 14C-cellulose [0-5 ml, 44000 d.p.m. (Cellulose-2-C14, ICN Pharmaceuticals, sp. act. 3.4 μCi mg-1)]. All cultures were incubated at 25 °C for 51 d, and 14CO2 evolved was monitored.

**Radiaerospirometry.** Biodegradation of 14C-radiolabelled material by A. bisporus was determined by monitoring the formation of 14CO2. The apparatus was based upon that described by Kirk et al. (1978), with 10 ml cultures in 100 ml Erlenmeyer flasks. The flasks were fitted with butyl rubber stoppers through which 2 mm glass tubes were inserted. One of these tubes extended into the flask within 1 cm of the culture medium; outside the flask it was connected via rubber tubing to a sterilizing filter. The second glass tube extended 5-15 cm or less into the flask, and the outside was connected via rubber tubing to a 3 cm 180-degree Luer-type needle. Clamps on both the rubber tubing allowed the flasks to be closed. The following modifications to the original apparatus design of Kirk were made. (i) All tubing was Butyl-XX (Esco Rubber, UK), 1-6 mm wall x 5 mm bore. This was found to be less permeable to CO2 than natural or silicone rubber and withstood repeated autoclaving. (ii) 18 mm bacterial air vents (Gelman Sciences) were incorporated into both inlet and outlet lines of each flask to maintain sterility over long periods. (iii) Hoffman clamps were replaced by polypropylene quick release clamps (Gallenkamp, UK). Flasks were flushed every 3-4 d by opening the clamps and forcing air through the filters at 100-200 ml min-1 for 15 min. This was found to remove all the evolved 14CO2, which was trapped by placing the Luer-type needle on the end of the flask exit line in 10 ml of an ethanolamine-containing scintillation fluid (Kirk et al., 1975) in a 20 ml scintillation
vial. Studies showed that $^{14}$CO$_2$ trapping was >98% efficient. The flask closures fitted with flushing ports and the apparatus used to flush the A. bisporus culture head space and trap the evolved $^{14}$CO$_2$ have been illustrated by Roberts (1985). All radioactivity measurements were made in a Beckman LC 7800 liquid scintillation counter with quench correction by external standards.

Results

The Casamino acids concentration in basic salts medium was found to have a considerable effect on the efficiency of incorporation of $^{14}$C by B. subtilis from [U-$^{14}$C]protein hydrolysate (Table 1). It was decided to use 1 mg 0.1% (w/v) Casamino acids ml$^{-1}$ in the subsequent experiments as this concentration gave the most efficient uptake of the radiolabel, whilst producing adequate bacterial biomass.

Results from experiments to demonstrate bacteriolysis by A. bisporus were subjected to statistical analysis. Plots of cumulative percentage degradation against time showed sigmoid-shaped responses. In most instances the curves were asymmetric, with a rapid exponential phase followed by a slower deceleration phase prior to approaching a limiting value by about day 50.

Cumulative count data are not amenable to the usual methods of least squares (Brain & Butler, 1988), primarily because successive count values are not independent but are increasingly dependent on previous counts. There are two aspects to most of the data sets reported here: the proportion of labelled C that has degraded, and the nature of the response of that which was degraded. The former statistics can be estimated either as the proportion of labelled C degraded on the last day of sampling, or can be determined by fitting an empirical distribution function (in most instances the lognormal) to the data and estimating the limiting proportion (Brain & Butler, 1988). In examining the nature of the degradation response it is useful to consider the time to a specific level of degradation; thus useful summary statistics are the median (time to 50% degradation), and the inter-quartile range (time from 25% to 75% degradation). This latter statistic can be thought of as a sort of slope, defining the rate of degradation. These statistics obviously only apply to the proportion of labelled isotope that was degraded. Fig. 1 gives a simple visual interpretation.

By considering the data in uncumulated form, i.e. as a frequency table of counts between given time intervals, a mean time to degradation can also be calculated. The asymmetric nature of most of the sigmoid responses

<table>
<thead>
<tr>
<th>Casamino acids (mg l$^{-1}$)</th>
<th>10$^{-3}$ × Radioactivity in basic medium before inoculation (d.p.m.)</th>
<th>Harvested B. subtilis</th>
<th>Radioactivity [d.p.m. (μg dry wt bacteria)$^{-1}$]</th>
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<td>10</td>
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<td>475</td>
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<th>Inter-quartile range</th>
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<th>Limiting*</th>
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<tr>
<td></td>
<td></td>
<td></td>
<td>Lower</td>
<td>Upper</td>
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</tr>
<tr>
<td>1</td>
<td>26.5</td>
<td>24.7</td>
<td>20.6-31.6</td>
<td>11.0</td>
<td>48.7</td>
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<tr>
<td>2</td>
<td>29.3</td>
<td>29.1</td>
<td>23.6-34.8</td>
<td>11.2</td>
<td>51.7</td>
<td>51.8</td>
</tr>
<tr>
<td>3</td>
<td>24.0</td>
<td>21.7</td>
<td>17.4-29.3</td>
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<tr>
<td>Mean</td>
<td>26.6</td>
<td>25.2</td>
<td>20.5-31.9</td>
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<tr>
<td>SEM</td>
<td>1.53</td>
<td>2.15</td>
<td>1.79</td>
<td>0.27</td>
<td>4.45</td>
<td>4.47</td>
</tr>
</tbody>
</table>

SEM, standard error of the mean.

* Limiting percentage degradation estimated from fitted lognormal distribution.
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Cumulative count

\[ T \text{ (total labelled)} \]

Proportion degraded = \( D/T \)

\[ D \text{ (total degraded)} \]

\[ D \text{ (total degraded)} \]

\[ 100 \]

50

25

0

Median

Inter-quartile range

Time

Fig. 1. Determination of median and inter-quartile range from cumulative degradation response.

meant that means determined on logarithmically-transformed data were more compatible with the median; consequently such means have been used throughout.

In the experiments reported here, summary statistics were calculated for each replicate using the statistical program MLP (Ross, 1987), and these values were then analysed by analysis of variance. Mean values are presented for experimental treatments (where appropriate) together with standard errors of differences between means (SEDS).

A bisporus was able to mineralize dead \( ^{14}\text{C}-\text{labelled} \) bacteria and utilize their cellular components as a sole source of carbon and nitrogen for growth (Table 2). These results confirm the earlier findings of Fermor & Wood (1981) and Grant et al. (1984) that \( A. \) bisporus produces a set of enzymes capable of efficiently utilizing the complex of bacterial polymers. The \( ^{14}\text{C} \) label was partitioned at the end of the experiment as follows: \( ^{14}\text{CO}_2 \) evolved (45-8\% of total \( ^{14}\text{C} \) added), culture supernatant liquid (26-6\%), microbial mat consisting of mushroom mycelium and undegraded bacteria (27-6\%). Plots of cumulative percentage degradation against time showed a sigmoid-shaped response. In this experiment there was a symmetrical degradation response with the median and mean values quite close together (Table 2). By day 56 degradation had virtually ceased, with an average of 45-8\% of labelled carbon collected.

Consistently higher bacteriolytic activities were obtained when \( A. \) bisporus was grown on lower concentrations of bacteria. On media containing 1-5 mg bacteria ml\(^{-1}\) the mean cumulative degradation was 64-9\% after 56 d (Table 3), whereas on media containing 3-0 mg bacteria ml\(^{-1}\) the cumulative degradation was only 45-8\% over the same time period (Table 2). The most notable feature of the data from the experiment in Table 3 was the impact of one of the control samples: exclusion of this sample from the analysis causes most of the differences between treatments to be statistically significant. The sample was markedly different from the other three controls in that only 30-9\% of the labelled C had degraded by day 49 compared with a mean of 64-9\% for the others. We can offer no explanation as to why this marked difference occurred. Analysis of the data excluding this sample showed several significant fea-

Table 3. Degradation by \( A. \) bisporus of killed \( ^{14}\text{C}-\text{labelled} \) B. subtilis used as sole carbon and nitrogen source (control), with added glucose, or with added cellulose in Treshow's basal salts liquid medium

Values for time to degradation (presented in d) are means of four replicate samples for each treatment.

<table>
<thead>
<tr>
<th>Treatment</th>
<th>Mean</th>
<th>Median</th>
<th>Lower quartile</th>
<th>Inter-quartile range</th>
<th>Percentage degradation</th>
</tr>
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<tbody>
<tr>
<td>( ^{14}\text{C}-\text{Bacteria} ) (control)</td>
<td>16-5</td>
<td>16-5</td>
<td>12-2</td>
<td>11-6</td>
<td>56-4</td>
</tr>
<tr>
<td>(excluding 1 value)</td>
<td>17-6</td>
<td>17-7</td>
<td>13-4</td>
<td>11-8</td>
<td>64-9</td>
</tr>
<tr>
<td>( ^{14}\text{C}-\text{Bacteria} ) + glucose</td>
<td>17-8</td>
<td>18-6</td>
<td>13-6</td>
<td>13-3</td>
<td>50-8</td>
</tr>
<tr>
<td>( ^{14}\text{C}-\text{Bacteria} ) + cellulose</td>
<td>15-3</td>
<td>15-3</td>
<td>11-6</td>
<td>10-7</td>
<td>48-8</td>
</tr>
<tr>
<td>sde (9 d.f.)</td>
<td>1-12</td>
<td>1-37</td>
<td>1-05</td>
<td>1-40</td>
<td>8-46</td>
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<td>(excluding 1 value)</td>
<td>0-64</td>
<td>1-00</td>
<td>0-51</td>
<td>1-47</td>
<td>5-11</td>
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<td>Significance†</td>
<td>NS</td>
<td>( P&lt;0-01 )</td>
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<td>(excluding 1 value)</td>
<td>**</td>
<td>NS</td>
<td>NS</td>
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</table>

SED, standard error of the difference between two means.
† NS, Not significant; *, \( P<0-05 \); †*, \( P<0-01 \).
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Table 4. Degradation by A. bisporus of killed 14C-labelled bacteria, glucose and cellulose

Cultures were grown with killed 14C-labelled bacteria as sole carbon and nitrogen source (control), unlabelled killed bacteria with added [14C]glucose, or unlabelled killed bacteria with added [14C]cellulose, in basal salts liquid medium. Values for time to degradation (presented in d) are means of four replicate samples for each treatment.

<table>
<thead>
<tr>
<th>Treatment</th>
<th>Mean</th>
<th>Median</th>
<th>Lower quartile</th>
<th>Inter-quartile range</th>
<th>Percentage degradation</th>
</tr>
</thead>
<tbody>
<tr>
<td>Control</td>
<td>15.8</td>
<td>15.2</td>
<td>12.1</td>
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<tr>
<td>Glucose</td>
<td>13.0</td>
<td>12.5</td>
<td>10.1</td>
<td>6.6</td>
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</tr>
<tr>
<td>Cellulose</td>
<td>13.7</td>
<td>14.5</td>
<td>8.6</td>
<td>14.9</td>
<td>9.1</td>
</tr>
<tr>
<td>SED (9 d.f.)</td>
<td>0.75</td>
<td>0.76</td>
<td>0.62</td>
<td>0.61</td>
<td>4.98</td>
</tr>
<tr>
<td>Significance†</td>
<td>*</td>
<td>*</td>
<td>***</td>
<td>***</td>
<td>***</td>
</tr>
</tbody>
</table>

SED, standard error of the difference between two means.
† *, P<0.05; † † † †, P<0.001.

Table 5. Degradation of killed 14C-labelled bacteria added to unsterile compost (control), unsterile compost with A. bisporus, or sterile compost with A. bisporus

Values for time to degradation (presented in d) are means of four replicate samples for each treatment.

<table>
<thead>
<tr>
<th>Compost treatment</th>
<th>Inoculation with A. bisporus</th>
<th>Mean</th>
<th>Median</th>
<th>Lower quartile</th>
<th>Inter-quartile range</th>
<th>Percentage degradation</th>
</tr>
</thead>
<tbody>
<tr>
<td>Unsterile</td>
<td>−</td>
<td>4.7</td>
<td>4.5</td>
<td>2.4</td>
<td>6.0</td>
<td>27.7</td>
</tr>
<tr>
<td>Sterile</td>
<td>+</td>
<td>12.0</td>
<td>13.0</td>
<td>6.6</td>
<td>16.4</td>
<td>26.6</td>
</tr>
<tr>
<td>SED (9 d.f.)</td>
<td>0.91</td>
<td>1.31</td>
<td>0.57</td>
<td>1.38</td>
<td>2.02</td>
<td>1.88</td>
</tr>
<tr>
<td>Significance†</td>
<td>***</td>
<td>***</td>
<td>***</td>
<td>***</td>
<td>NS</td>
<td>NS</td>
</tr>
</tbody>
</table>

SED, standard error of the difference between two means.
† NS, Not significant; † † † †, P<0.001.

tures: total bacterial degradation in cultures containing additional cellulose was less than that for the control flasks containing bacteria alone, and it took place somewhat faster. The degradation of bacteria when additional glucose was incorporated was slower than that in the control cultures containing bacteria alone and it was estimated that some further degradation took place after day 49 with this treatment.

The proportion of labelled C degraded was much lower from the unlabelled bacteria plus 14C-cellulose treatment than from the control (14C-labelled bacteria) and unlabelled bacteria plus 14C-glucose-enriched treatments (Table 4). Degradation as determined by the limiting value was almost complete by day 51 when sampling was terminated. The rate of uptake of labelled glucose was faster than that for labelled bacteria in the control (a smaller lower quartile and a shorter inter-quartile range). The uptake of labelled cellulose was only 10%, so that the treatment summary for label uptake is of little consequence, though it should be noted that the inter-quartile range was greater than for the other two treatments. This suggests that the degradable carbon in the cellulose is less readily available than from the other two sources.

The micro-organisms present in compost can also degrade components of bacteria added to compost. Summary statistics of time to degradation of 14C-labelled bacteria in unsterile compost (control), in unsterile compost with A. bisporus, or in sterile compost with A. bisporus are presented in Table 5. By day 64 all treatments had effectively reached their limiting levels of degradation, which were not significantly different. On the unsterile compost half the degradable carbon had been released as CO₂ within about 6 d, but thereafter the rate slowed down more quickly when A. bisporus was present (note the longer inter-quartile range). The process of degradation was much slower for the sterile compost, though the same percentage degradation was achieved by day 64 (Table 5).

A. bisporus and the compost microflora were able to mineralize biomass produced in situ where label was incorporated into the microbial biomass by composting.
rather than added as killed bacteria. The effect of adding
_A. bisporus_ to sterile and unsterile labelled compost produced in this manner is shown in Table 6. The variability between samples without _A. bisporus_ was considerably less than that where the mushroom was present. In the latter case the percentage degradation was greater on day 49 for cultures containing _A. bisporus_ and the rate of degradation thereafter was greater, though not significantly so. Differences between sterile and unsterile composts were less marked: the difference in variability would suggest that differences between composts without _A. bisporus_ present are important.

### Discussion

This is the first study to produce quantitative data to show that _A. bisporus_ can mineralize (biodegrade) dead and live bacteria _in vivo_ and _in vitro_, validating previous evidence (Fermor & Wood, 1981; Grant _et al._, 1984; Barron, 1988; Rainey, 1989) of the bacteriolytic abilities of _A. bisporus_. The mineralization process is efficient, as evidenced by the release of >20% of total radiolabel in the compost as ^14^CO_2_, and confirms earlier suggestions that microbial biomass might be significant in the nutritional strategy of _A. bisporus_ (Sparling _et al._, 1982; Fermor, 1988). The total microbial biomass in commercial mushroom compost at the end of composting was estimated as 9.2 mg carbon (g dry wt compost)^-1_ (Sparling _et al._, 1982). This represents approximately 2% of the compost dry weight assuming that microbial biomass contains 50% (w/w) carbon. The weight of _A. bisporus_ mycelium in fully colonized compost has been estimated, using extracellular laccase activity as a growth marker, to be 50–125 mg (g compost)^-1_ (Wood, 1979). The mean ratio of mycelium dry weight to fruitbody dry weight was 1.78:1, giving a total _A. bisporus_ biomass of approximately 137 mg g^-1_. If _A. bisporus_ decomposes a high proportion of the compost microbial biomass, as indicated by the results reported here, assuming that 50% of the biomass weight was lost through respiration as CO_2_, then the maximum contribution of microbial biomass to the _A. bisporus_ biomass would be less than 10%. This means that _A. bisporus_ probably obtains the bulk of its carbon nutrition from the plant polymers within straw. However, the nitrogen content of the microbial biomass is high, and since micro-organisms will also concentrate minerals during composting this biomass could act as a concentrated source of nitrogen and minerals.

It is possible to manipulate microbial species diversity and populations in compost by both physical and chemical means; thus stimulation of _A. bisporus_ mycelium growth may be attainable by encouraging the growth of beneficial micro-organisms. It is not yet known whether there is any direct relationship between the quantity of microbial biomass in a compost and the quantity of mushroom mycelium that a compost will support. The lower mushroom yields obtained from substrates produced by accelerated composting techniques (Smith, 1983) might be attributable to the lower microbial biomass present or to less degradation of the lignocellulosics. An understanding of the nutritional relationship between white-rot basidiomycetes and other micro-organisms in solid substrates may help to answer the question why some basidiomycetes can successfully colonize uncomposted lignocellulosic substrates whilst others require lignocellulose to be considerably degraded. There have been no studies to determine whether _Lentinula, Pleurotus_ or _Volvariella_ (the other major cultivated fungi) can utilize microbial
biomass in wood, wheat straw or rice straw/cotton waste respectively. These three basidiomycetes can successfully colonize uncomposted materials. For example, *Lentinula* colonizes wood which has a C:N ratio in excess of 100:1. It is possible that the nitrogen supply for mycelial growth of these fungi comes from primary microbial colonization of the felled timber and the microorganisms present in this environment may include diazotrophs.

The bacteriolytic enzyme system produced by *A. bisporus* could have at least two roles. One is to act as a nutrient-releasing system to obtain carbon, nitrogen and minerals from deposits concentrated within the microbial biomass. The ability to efficiently release such nutrients from microbial biopolymers in a complex phenolic-rich environment may be one of the factors selecting for growth of *A. bisporus* mycelium. It would be of interest to determine the growth efficiency of a range of soil microorganisms to mineralize microbial biopolymers from such substrates.

Another possible role for the lytic and oxidative enzymes of *A. bisporus* could be to act as part of an antimicrobial system. *A. bisporus* mycelium growing in compost produces a large quantity of fungal biomass. This biomass would be an ideal nutritional source for competitive microorganisms in this habitat. However *A. bisporus* appears to suffer little microbial predation in this substrate.

These studies have shown that *A. bisporus* possesses an efficient bacteriolytic system in its commercial growth substrate, composted wheat straw. Further investigations are in progress to characterize the individual enzyme activities responsible for this system and to determine the regulation of their production.

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**References**


