COMPARATIVE GROWTH OF TRICHODERMA STRAINS IN DIFFERENT NUTRITIONAL SOURCES, USING BIOSCREEN C AUTOMATED SYSTEM

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ABSTRACT

Trichoderma is one of the fungi genera that produce important metabolites for industry. The growth of these organisms is a consequence of the nutritional sources used as also of the physical conditions employed to cultivate them. In this work, the automated Bioscreen C system was used to evaluate the influence of different nutritional sources on the growth of Trichoderma strains (T. hamatum, T. harzianum, T. viride, and T. longibrachiatum) isolated from the soil in the Juréia-Itatins Ecological Station (JIES), São Paulo State - Brazil. The cultures were grown in liquid culture media containing different carbon- (2%; w/v) and nitrogen (1%; w/v) sources at 28ºC, pH 6.5, and agitated at 150 rpm for 72 h. The results showed, as expected, that glucose is superior to sucrose as a growth-stimulating carbon source in the Trichoderma strains studied, while yeast extract and tryptone were good growth-stimulating nitrogen sources in the cultivation of T. hamatum and T. harzianum.

Key words: Bioscreen C system, growth, Trichoderma, tryptone, yeast extract.

INTRODUCTION

Fungi, in general, possess diverse abilities that form the basis for industrial research. Fungi are used in medical/pharmaceutical industries for the manufacture of antibiotics, such as penicillin, produced by Penicillium chrysogenum, of enzymes such as amylases, pectinases, and of cellulases produced by Aspergillus spp. and by Trichoderma spp., as well as in food processing (14,17).

Strains of Trichoderma can produce a great number of metabolites like antibiotics (10), degrade complex substrates such as cellulose (19), chitin, xylan, and lignin (6), and also possess proteolytic activity (25). Because of these properties, this genus has high biotechnological potential and is found in almost all ecosystems. Several studies have been carried out on Brazilian soil, such as in the Paranapiacaba Biological Reserve, SP (3). In order to improve knowledge of the biodiversity of soil microbiota, 112 fungi species were isolated and identified from the Juréia-Itatins Ecological Station (JIES), São Paulo, Brazil (31). These species are currently being studied with respect to their potential for producing macromolecules of interest to medicine and industry. The same authors determined that some of these strains are able to produce enzymes like xylanases and fatty acids including γ-linolenic acid (GLA), and observed that the species of Trichoderma probably plays an important role in soil nutrient recycling (20,28).

Like in the Bioscreen system, turbidimetric methods have been used for studies on microbial growth, providing fast results that are expressed in Optical Density (O.D.) units (21). This equipment has been used for many purposes in microbiology, including basic research (1), to study the effects of chemical compounds on microbial growth (17,19), for quality control (21), and in pollutant degradation tests (34).
Considering the importance of a knowledge of the factors that influence the growth of fungi and the biotechnological potential of the genus *Trichoderma*, four strains were chosen (*Trichoderma hamatum*, *Trichoderma harzianum*, *Trichoderma viride*, and *Trichoderma longibrachiatum*) and isolated from the Juréia-Itatins Ecological Station- SP, Brazil (JIES) soil (31), in order to compare their potential to grow in submerged cultures, in the presence of a carbon source (glucose or sucrose) and a nitrogen source (tryptone or yeast extract), by means of the Bioscreen C automated system.

**MATERIALS AND METHODS**

**Microorganisms**

Four strains were studied - *T. hamatum*, *T. harzianum*, *T. viride*, and *T. longibrachiatum* - that were isolated from the soil (0 - 15 cm deep) in the Banhado Grande area from JIES (LAT 24° 18' to 24° 37' S - LONG 47° 00' to 47° 31' W) in both natural- and secondary vegetation areas. The samples were collected, packaged into plastic bags, transported to the laboratory, and processed immediately to avoid storing them. These samples were sifted using 2.38 mm screens and homogenized. Isolation involved processing the soil samples by saline sterilization, followed by centrifugation, and a series of dilutions. Later, each sample was placed in Petri dishes (n = 3) followed by the respective culture medium (31).

These strains are kept in culture medium of MEA (29) (malt extract agar of composition (in w/v): malt extract 2%; bacteriological agar 2%; glucose 2%; peptone 0.1%; yeast extract 0.1% in Tween (T-80) 0.1% (v/v). The counting was done in a Neubauer chamber in NaCl, 0.85% (w/v).

**Cultures**

Strains were cultivated in a Bioscreen C system (Lab Systems Helsinki, Finland), in GY Medium, pH 6.5, with a single carbon source at 2% (w/v) (glucose or sucrose), and a single nitrogen source at 1% (w/v) (tryptone or yeast extract) (24). Accordingly, four different cultures were tested, namely Culture 1a (trypotone + glucose); Culture 1b (trypotone + sucrose); Culture 2a (yeast extract + glucose), and Culture 2b (yeast extract + sucrose). The working volume in the Bioscreen plate was 400 µL/well comprising 360 µL of each culture medium and 40 µL of each spore solution. Temperature was controlled at 28°C and O.D. at 540 nm. Absorbance values of the cell suspensions were automatically read at regular intervals of 2 h, over a 72 h period, and before each measurement, the cell cultures were automatically shaken for 60 seconds. All experiments were performed in quadruplicate. The control tubes contained only the culture medium.

**Data analysis**

1. **Comparative analysis of the strains’ growth**

The O.D. values from Bioscreen C were analyzed using spreadsheet software (Microsoft Excel 97) and the averages calculated of the quadruplicates for each type of culture medium. These averages were used to generate growth curves for each type of culture (O.D. 540 nm of the cultures versus incubation time). Figures present the data from the average of readings (from quadruplicates) taken from every fourth hour of cultivation.

Because the spore solution of each strain presented different colors, at time zero, the initial O.D. values were distinctive to each culture in relation to the control culture. This coloration difference (defined here as “specific color influence factor” for each strain) was calculated as follows:

**Color influence = O.D. of each culture (t0) - O.D. of the control (t0)**

Next, the influence factor specific to each strain was subtracted from all O.D values from 0 to 72 h of each culture, in order to express the real growth of the strains. The analysis of variance (Kruskal-Wallis test, BioEstat 4.0 software) was applied using the data obtained, and confirmed by Dunn *post hoc* test, which was used to verify the significance of the strain’s growth compared to the control and among the strains (5). The experimental results were separated into four classes based on the time taken for growth: I) 0 to 18 h, II) 18 to 36 h, III) 36 to 54 h, and IV) 54 to 72 h. The results were analyzed by triangulation of the statistical data with the differences of O.D. per period.

2. **Analyses of the growth of individual strains in different conditions of cultivation**

After excluding the color influence factor, the growth of each strain was calculated as the difference (Δ) between the average values of the O.D. 540 nm, final and initial, taking into account the fact that each average value of O.D. came from the replicates.

The relative growth of each strain, expressed in percentage, was obtained by comparing the performance of a single strain in every two different cultivation conditions, considering the smallest O.D. value as 100% and the highest O.D. value (greater than 100%) as the highest rate of relative growth among them.

**RESULTS AND DISCUSSION**

The quick determination of optimal growth of industrially interesting microorganisms, accompanied by the optimum production of economical value molecules, is very important for industry. The method used in this study is neither time consuming nor requires high costs, since it is no longer necessary to perform numerous tests like it is done by the traditional method of dilutions in series, while also enabling determination of optimal growth that is of industrial interest.
In optimizing the growth of *Trichoderma* sp., some parameters were reported for instance, the 72 h period required for cultivation (33). As for the temperature, studies on the activity of xylanases from mesophilic fungi showed that, the activity of these enzymes was optimized between 25ºC and 30ºC for submersc cultures (under constant agitation) and for stationary cultures with solid substrates (13).

Studies also verified the effect of different temperatures on the production of several enzymes, among them carboxymethylcellulases and xylanases, on strains of several species isolated from JIES - SP - Brazil soil, such as *Aspergillus* and *Trichoderma*, and found 28ºC to be the optimum temperature and which was the temperature chosen for the cultures prepared in this study (26).

Figs. 1 and 2 show the growth curves of the *Trichoderma* strains studied, obtained by their cultivation from 0 to 72 h and Table 1 records the results of the statistical analysis.

1. Comparative analysis of the growth of strains

In the cultures with tryptone and glucose, the growth among the strains of *T. hamatum*, *T. harzianum*, *T. viride* and *T. longibrachiatum* was significantly different in all the periods (H0-18h=33.36; H20-36h=40.21; H36-54h=40.90; H56-72h=41.83, and p<0.0001). With regard to the differences among the average scores per period (Table 1), it can be seen when individually analyzed against the control that in *T. hamatum* vs *T. viride* and in the strains of *T. hamatum*, *T. harzianum*, and *T. longibrachiatum*, all presented values of p<0.05 in most periods, and their growth rates were considered highly significant. Besides, *T. hamatum* differed not only from the control but also from *T. viride*, which did not happen with *T. harzianum* and *T. longibrachiatum*; whereas, growth rates of both *T. harzianum* and *T. longibrachiatum* differed only from the control, but not from each other. Thus, it can be concluded from these observations that *T. hamatum* presented the highest value of R and consequently higher growth in all periods (Table 2). This was followed by *T. harzianum* and *T. longibrachiatum* that exhibited similar growth rates, and lastly by *T. viride*, (Fig. 1).

In the cultures with tryptone and sucrose, the growth rates of the strains were also significantly different in all the periods (H0-18h=36.96; H20-36h=41.93; H36-54h=39.65; H56-72h=41.24; and p<0.0001). The differences among the average scores were not significant, especially between *T. hamatum* vs *T. viride*, *T. hamatum* vs control, *T. harzianum* vs *T. viride*, *T. harzianum* vs control, and *T. longibrachiatum* vs control. These findings can also be confirmed from the growth curves (Fig. 1B) in which *T. hamatum* and *T. harzianum* presented similar growth. Nevertheless, *T. longibrachiatum* exhibited a slightly smaller growth rate, while *T. hamatum* and *T. harzianum* showed the highest growth among the strains, followed by *T. longibrachiatum*, which is also confirmed by their respective values of R (Table 2).

It can be seen that *T. harzianum* showed a similar growth rate in the cultures containing tryptone with glucose or sucrose. The same was observed of *T. longibrachiatum*, but the growth rate of *T. hamatum* was higher in the medium containing glucose as carbon source than in the one with sucrose (Fig. 1).

Only few studies use tryptone as growth culture medium for fungi, as it is frequently used as nitrogen source for bacteria growth. Peptone is a similar molecule frequently used for growth of fungi such as *Mucor* (32), *Aspergillus* (23), and *Trichoderma* (35). Studies involving *Aspergillus flavus* growth report that tryptone is as effective as peptone (22), and is cheaper. Due to

![Figure 1](406.png)

**Figure 1.** Growth curves (O.D. 540 nm) of *T. hamatum*, *T. longibrachiatum*, *T. viride*, and *T. harzianum* in medium containing tryptone and glucose (a) or tryptone and sucrose (b) incubated at 28ºC from 0 until 72 h in an automated Bioscreen C system.
Table 1. Growth differences of the strains for culture periods in different culture media. The z values were obtained from variance analysis (Kruskal-Wallis test) and confirmed by Dunn post hoc test (BioEstat 4.0 software).

| Culture 1a (tryptone and glucose) | Culture 1b (tryptone and sucrose) | Culture 2a (yeast extract and glucose) | Culture 2b (yeast extract and sucrose) |
|-----------------------------------|-----------------------------------|--------------------------------------|--------------------------------------|
| 0-18h                             | 20-36h                            | 38-54h                               | 56-72h                               |
| T. hamatum x T. harzianum         | 2.010 2.082 1.561                 | 0.215 1.525 0.628                    | 0.969 2.737 2.261                    |
| T. hamatum x T. longibrachiatum   | 2.916 2.279 2.602 2.800 2.288 2.835 2.333 2.665 2.351 2.118 2.315 2.880 | 3.697 2.225 1.507 1.454             |
| T. hamatum x T. viride            | 5.133 4.361 4.361 4.361 3.508 4.361 3.894 4.119 5.016 4.343 3.822 4.334 2.862 4.092 2.961 2.907 |
| T. hamatum x controle             | 4.477 5.814 5.814 5.814 5.025 5.814 5.348 5.572 4.343 5.814 5.276 5.788 5.447 5.545 4.415 4.361 |
| T. harzianum x T. longibrachiatum | 0.906 0.197 0.843 1.238 2.073 1.310 1.705 1.696 0.386 0.144 1.884 1.481 1.166 0.897 2.853 2.907 |
| T. harzianum x T. viride          | 2.271 2.279 2.602 2.800 2.293 2.835 3.266 3.149 2.279 2.082 3.392 2.934 0.332 2.764 4.307 4.361 |
| T. harzianum x controle           | 2.468 3.733 4.056 4.253 4.809 4.289 4.720 4.603 1.606 3.553 4.845 4.388 2.916 4.217 5.761 5.814 |
| T. longibrachiatum x T. viride    | 2.216 2.082 1.759 1.561 1.220 1.525 1.561 1.454 2.665 2.225 1.507 1.454 0.834 1.866 1.454 1.454 |
| T. longibrachiatum x controle     | 1.561 3.535 3.212 3.015 2.737 2.979 3.015 2.907 1.992 3.697 2.961 2.907 1.750 3.320 2.907 2.907 |
| T. viride x controle              | 0.655 1.454 1.454 1.454 1.516 1.454 1.454 1.454 0.673 1.472 1.454 1.454 2.584 1.454 1.454 1.454 |

Critical z value for p is 2.807
z values in bold are significant (p<0.05)

Table 2. R-values of each strain obtained from variance analysis (Kruskal-Wallis test) confirmed by Dunn post hoc test (BioEstat 4.0 software).

| Culture 1a (tryptone and glucose) | Culture 1b (tryptone and sucrose) | Culture 2a (yeast extract and glucose) | Culture 2b (yeast extract and sucrose) |
|-----------------------------------|-----------------------------------|--------------------------------------|--------------------------------------|
| 0-18h                             | 20-36h                            | 38-54h                               | 56-72h                               |
| T. hamatum                        | 369 369 369 369 330 369 343 355.5 | 368 369 339 367.5                    | 369 354 291 288                      |
| T. harzianum                      | 257 253 271 282 318 284 308 301.5 | 215.5 243 315 289.5                  | 228 228 366 369                      |
| T. longibrachiatum                | 206.5 242 224 213 202.5 211 213 207 | 237 251 210 207                     | 163 230 207 207                     |
| T. viride                         | 83.5 126 126 126 134.5 126 126 126 | 88.5 127 126 126                    | 209.5 126 126 126                   |
the absence of studies related to the growth of fungi using tryptone, it was chosen as the nitrogen source in the present study.

In our study, differences in the growth rates among the strains could also be noticed ($H_0 - 18h = 30.63$; $H_{20} - 36h = 40.07$; $H_{36} - 54h = 39.90$; $H_{56} - 72h = 42.15$ and $p < 0.0001$) in the cultures containing yeast extract and glucose. The results of the average scores showed that only *T. hamatum* exhibited a significant difference in the growth in the period of 0 up to 18 h of cultivation. From the 38th hour of cultivation, the growth rate of the strains of *T. hamatum*, *T. harzianum*, and *T. longibrachiatum* increased (Table 1, Fig. 2), and a significant difference in growth could be seen from that period for *T. harzianum* vs *T. viride*, *T. harzianum* vs control, and *T. logibrachiatum* vs control (Table 1). Considering that the average scores of *T. hamatum* exhibited a significant rate in all the periods and also presented the highest value of R (Table 2), it can be assumed that this strain too presented the highest growth rate among those tested, followed by *T. harzianum* and *T. logibrachiatum*.

In cultures containing yeast extract and sucrose, different growth rates were also detected among the strains ($H_0 - 18h = 31.38$; $H_{20} - 36h = 38.97$; $H_{36} - 54h = 42.12$; $H_{56} - 72h = 42.42$ and $p < 0.0001$). The average scores of *T. hamatum* vs *T. viride* and the scores of *T. hamatum* when individually analyzed, against the control, were significantly different in all the periods (Table 1), while *T. harzianum* vs *T. longibrachiatum* and *T. harzianum* vs *T. viride* exhibited significant differences only after 38 h of cultivation. On comparing these data, it can be seen in the Fig. 2 that, from the 38th hour, the growth curve of *T. harzianum* surpassed that of *T. hamatum* and, consequently, the R-value of *T. harzianum* increased in that period too (Table 2).

### 2. Analyses of the growth of each individual strain in different cultivation conditions

On comparing the cultivation conditions tested here, it can be concluded that in the cultivation of *T. harzianum* there was an increase in the growth rate at approximately the 36th hour in all the conditions (Figs. 1, 2). From this period, the strain started to differ significantly from both *T. viride* (in the cultivation with yeast extract and glucose) and *T. longibrachiatum* (when cultivated in yeast extract and sucrose) (Table 1). Analyzing only the nitrogen sources in function of the carbon sources used, it was observed that the growth of *T. harzianum* was 30% higher with yeast extract plus glucose than with tryptone plus glucose (Figs. 1, 2). The same pattern was observed with regard to media with sucrose, where the growth with yeast extract was 65% higher than with tryptone (Figs. 1, 2). Thus, yeast extract appears to be the best nitrogen source for *T. harzianum*, compared to tryptone, apart from the carbon source used in this study. Studies involving xylanase production indicated that *T. harzianum* is a good producer of xylanases in media containing yeast extract (7). The comparative analysis of the carbon sources in function of the nitrogen sources showed that in the presence of tryptone, growth of *T. harzianum* in the cultures with glucose was 15% higher than that in sucrose (Fig. 1), while in presence of yeast extract the growth was 10% higher in the cultures with sucrose than in those with glucose (Fig. 2).

The data above show that yeast extract is better for growth of *T. harzianum*, especially when combined with sucrose, and differs in this respect from *T. viride* (Table 1).

As expected with all cultures, the growth of *T. hamatum* in glucose was much better than in sucrose, as glucose is a simpler metabolite than sucrose. Considering only the influence of sugar,
it can be seen that growth of *T. hamatum* was 40% higher in glucose than in sucrose in presence of tryptone as nitrogen source (Fig. 1). Likewise, the growth of *T. hamatum* was 45% higher in glucose than in sucrose in the medium with yeast extract (Fig. 2). These results indicate that for *T. hamatum* cultivation, glucose is better than sucrose as carbon source, irrespective of the nitrogen source used. Analyzing influence of the nitrogen source on the growth rate of *T. hamatum*, a slightly higher growth (20%) could be observed in yeast extract than in tryptone, when glucose was present in the medium (Figs. 1, 2). Almost the same result was detected when sucrose was used as carbon source, in which growth of *T. hamatum* was 15% higher than in yeast extract, indicating that in this strain yeast extract induces a higher growth rate (Figs. 1, 2).

*T. longibrachiatum* presented similar growth rates in all the cultivation conditions tested here (with small differences ranging from 10 to 15%) which can be observed in both Figures. According to the statistical data, this strain differed from the control only from the 20th hour of growth, and did not differ from *T. viride* in any period. This last strain did not show a growth pattern different from the control or exhibit a significant growth rate. This suggests that for these two strains, *T. longibrachiatum* and *T. viride*, further studies are necessary because the nutritional sources tested here may not be suitable for their cultivation.

Different carbon sources were also used in analysis of fungi and production of total lipids, as well as γ-linoleic acid, maintaining the nitrogen source (yeast extract) constant (26). The results showed a five times higher growth rate in the culture medium containing glucose, compared to that with sucrose.

Yeast extract has effectively been the best nitrogen source for growing filamentous fungi and is the best inducer for production of different macromolecules, a characteristic also observed, in recent studies, for thermophilic fungi (15). It was confirmed that significant mycelial growth resulted in *T. aureoviride*, *T. viride*, and *T. harzianum* strains when cultivated in media containing yeast extract and agricultural soil (16), this last being a nitrogen source considered very stimulating for production of enzymes such as xylanases and glucanases and many other metabolites (8), besides having clinical importance, causing infections, as reported for *T. viride* (18), *T. longibrachiatum* (16) and *T. harzianum* (12). Thus, the verification of nutritional parameters for the strains tested in our study is promising for optimizing their biotechnological- and industrial uses.

Some studies on microorganisms’ growth compared the Bioscreen automated system and the traditional cultivation method, and many patterns of similarity between them were found. One of these studies (34) confirmed that the Bioscreen automated system is a promising and reliable method for several uses in microbiology.

Considering that the microorganism cultivation in industries must be carried out on large scale for production of innumerable commercially interesting metabolites, our results indicate that the Bioscreen system provides a fast- and economical way to test and optimize growth conditions for further use in such large-scale production.

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**RESUMO**

*Crescimento de linhagens de Trichoderma em diferentes fontes nutricionais, empregando o sistema automatizado Bioscreen C.*

*Trichoderma* é um dos gêneros de fungos produtores de metabólitos de interesse industrial. O crescimento destes organismos é consequência das fontes nutricionais utilizadas, juntamente com as condições físicas de cultivo. Neste trabalho, o sistema automatizado Bioscreen C foi utilizado para avaliar a influência de diferentes fontes nutricionais sobre o crescimento de linhagens de *Trichoderma* (*T. hamatum*, *T. harzianum*, *T. viride* e *T. longibrachiatum*) isoladas do solo da Estação Ecológica da Juréia-Itatins (JIES), São Paulo - Brasil. Os cultivos foram feitos em meios líquidos de cultura contendo diferentes fontes de carbono (2%; w/v) e nitrogênio (1%; w/v) a 28°C, pH 6,5 e agitados a 150 rpm durante 72 h. Os resultados mostraram, conforme esperado, que a glicose é melhor do que a
sacarose como fonte de carbono induutora de crescimento das linhagens de *Trichoderma* testadas, enquanto que, o extrato de leveduras e a tripotona foram boas fontes de nitrogênio indutoras de crescimento para os cultivos de *T. hamatum* e *T. harzianum*.

**Palavras chaves**: Bioscreen C, crescimento, Trichoderma, tripotona, extrato de leveduras.

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