

Bioreactor design and implementation strategies for the cultivation of filamentous fungi and the production of fungal metabolites: from traditional methods to engineered systems

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The production of fungal metabolites and conidia at an industrial scale requires an adequate yield at relatively low cost. To this end, many factors are examined and the design of the bioreactor to be used for the selected product takes a predominant place in the analysis. One approach to addressing the issue is to integrate the scaling-up procedure according to the biological characteristics of the microorganism considered, *i.e.* in our case filamentous fungi. Indeed, the scaling-up procedure is considered as one of the major bottlenecks in fermentation technology, mainly due to the near impossibility of reproducing the ideal conditions obtained in small reactors designed for research purposes when transposing them to a much larger production scale. The present review seeks to make the point regarding the bioreactor design and its implementation for cultivation of filamentous fungi and the production of fungal metabolites according to different developmental stages of fungi of industrial interest. Solid-state (semi-solid), submerged, fermentation and biofilm reactors are analyzed. The different bioreactor designs used for these three processes are also described at the technological level.

Keywords. Bioreactors, fungi, secondary metabolites, conidia, solid state fermentation.

Conception de bioréacteurs et mise en œuvre de stratégies pour la culture de champignons filamenteux et la production de métabolites d'origine fongique : des méthodes traditionnelles aux technologies actuelles

La production de métabolites d'origine fongique et de conidies à l'échelle industrielle requiert un procédé adéquat à cout relativement bas. À cet effet, beaucoup de facteurs sont analysés et la configuration du bioréacteur à utiliser pour le produit retenu occupe une place prépondérante dans l'analyse. Une approche à suivre pour résoudre le problème relatif à la production de métabolites et de conidies par les champignons consiste à maîtriser la montée en échelle du procédé de fermentation avec le respect de caractéristiques biologiques de micro-organismes, tels que les champignons filamenteux pour le cas qui concerne notre étude. En effet, cette montée en échelle est considérée comme l'un des principaux points d'achoppement en technologie de fermentation : ceci en raison de la quasi impossibilité de reproduire exactement les mêmes conditions idéales obtenues en utilisant les petits réacteurs de recherche au moment de les transposer à la production à grande échelle. La présente revue vise à faire le point sur la conception de bioréacteurs et leur mise en œuvre pour la culture de champignons filamenteux et la production de métabolites en fonction des différents stades de développement de champignons d'intérêt industriel. Les cultures solide (semi-solide), submergée et en biofilm ont été prises en compte dans les différentes études. Les différents types de bioréacteurs utilisés pour les trois procédés sont aussi décrits du point de vue technologique.

Mots-clés. Bioréacteur, champignon, métabolite secondaire, conidie, fermentation à l'état solide.

1. INTRODUCTION

Three fungal genera are widely used for biotechnological applications: *Aspergillus* sp., *Penicillium* sp., and to a lesser extent, *Trichoderma* sp. Filamentous fungi have the capability to produce extremely large quantities of homologous proteins, including several enzymes that can be used as industrial biocatalysts. Several volatile metabolites (MVOC) can also be produced. However, the production of these metabolites relies on the culture mode at both the quantitative and qualitative levels. Indeed, these fungi are able to grow in different forms: from free mycelium and pellet in liquid phase, to pellet and mycelium to conidia at a solid-air interface. Based on this, different bioreactor strategies have been developed in order to promote a given development stage. In this context, three modes of cultivations can be considered:

- submerged fermentation where two kinds of morphologies are most often depicted, *i.e.* free mycelium (with different degrees of ramification) and pellets;
- semi-solid fermentation which is reflected by the development of mycelium on the surface and at the interior of a solid substrate in the absence of free water;
- biofilm formation, represented by the development of mycelium on an inert support, such as stainless steel, glass or Teflon material integrated in an adapted reactor.

Among these three strategies, submerged fermentation is the most widely used mode of fermentation. However, the last two strategies lead to the enhancement of the development of aerial hyphae with the production of conidia on conidiophore position of filaments depending on the development stage of the microorganism. They will be able to generate fungal products of biotechnological interest, since the excretion capacity is increased when fungal biomass is attached on a given support. Barrios-Gonzalez (2012) reported that the solid-state fermentation is considered as an alternative culture method that has gained researchers attention over the past 20 years, and credibility among many industrial corporations. The present review is aimed at understanding the influence of the cultural conditions on the developmental stage of the fungi, as well as on the excretion of secondary metabolites of industrial interest, which highlight the orientation for bioreactor design needed. It is focused on bioreactor, fermentation, strategies for secondary metabolites enhancement and some applications cultivation of *Trichoderma*, *Penicillium* and *Aspergillus*. In the conclusion, a new range of biofilm reactors will be proposed for the improvement of the production of fungal metabolites.

2. TRADITIONAL METHODS AND ENGINEERED SYSTEMS APPLIED FOR THE CULTIVATION OF FILAMENTOUS FUNGI IN ORDER TO PRODUCE THE METABOLITES AND CONIDIA

2.1. Basic bioreactor design: from submerged fermentation to biofilm reactor

A bioreactor can be defined as mechanically stirred vessel in which organisms are cultivated in a controlled manner and/or materials are converted or transformed through specific reactions. Indeed, traditional methods for solid state fermentation include tray, drum and packed bed bioreactors with problems regarding the control of different parameters; while, for submerged fermentation they include continuous stirred-bioreactors, continuous flow stirred-tank reactors, plug-flow reactor, and fluidized-bed reactors. For biofilm processes, the material supporting the microorganism can be used in reactors applied for submerged fermentation and adapted for used in order to cultivate immobilized cells or biofilms with respect to the production of value-added molecules (Muffler et al., 2014).

2.2. Factors affecting bioreactor design

Many factors contribute to the design of a bioreactor. Durand (2003) reported that compared to submerged fermentation, the solid media used in solid-state fermentation contain less water but an important gas phase existed between the particles. This feature is of great importance because of the poor thermal conductivity of air compared to water. Another point is the wide variety of matrices used in solid-state fermentation, which vary in terms of composition, size, mechanical resistance, porosity and water-holding capacity. All these factors can affect the reactor design and the strategy to control the key parameters. Indeed, with submerged fermentation, it can be considered an approximation that all the media are made up essentially of water. In this environment, temperature and pH regulations are trivial and do not pose problems during the scaling-up of a process.

With submerged fermentation, the difficulty mainly encountered is related to limitations at the level of oxygen transfer capacity, which depends upon the shape and size of the reactor and the agitation/aeration system used. To characterize this transfer, a parameter, $K_L a$ (oxygen transfer coefficient), has been defined. It can be considered as a “similarity invariant”, which means that its value expresses the capacity of the equipment to transfer oxygen independent of the volume of the reactor and, as such, constitutes an important parameter used in scale-up studies in submerged fermentation.

With solid-state fermentation, besides oxygen transfer, which can be a limiting factor for some designs, the problems are more complex and affect the control of three important parameters, *i.e.* temperature, pH and water content of the solid medium. There are also other factors affecting the bioreactor design, namely:

- the morphology of the fungus (the presence or not of a septum in the hyphae) and, related to this, its resistance to mechanical agitation,
- the necessity or not of having a sterile process.

Concerning fungal morphology, as reported by Papagianni (2004), filamentous fungi are morphologically complex microorganisms, exhibiting different structural forms throughout their life cycles. The basic vegetative structure of growth consists of tubular filaments known as hyphae that originate from the germination of a single reproductive conidium or a piece of mycelium. As the hyphae continue to grow, they branch frequently and repeatedly to form a mass of hyphal filaments referred to as a mycelium. When grown in submerged culture, these fungi exhibit different morphological forms, ranging from dispersed mycelial filaments to densely interwoven mycelial masses referred to as pellets, whereas conidiation, as the end point of the fungi's developmental cycle, is rarely achieved during submerged cultivation, partially due to relatively good nutrient availability and partly to the physical nature of the hyphal wall. Different metabolites are produced by these different forms in submerged culture, so the successful production of fungal metabolites requires detailed knowledge of the growth characteristics and physiology of the fungus considered. Not only does the production of different metabolites require different physiological conditions, but each fungal species is also unique in its anatomical, morphological and physiological development. Thus, for fungal fermentation, the precise physiological conditions and correct stage of development must be established to achieve the maximal product formation. In other words, controlling the form of these microorganisms is a real issue that needs great attention in order to make optimal use of their potential production capacities during their development according to the structure of bioreactor designed.

2.3. Solid-state fermentation processes

Many examples of solid-state fermentation have been reported by researchers, some of which are depicted in a figure presented by Durand (2003) and others where they represented tray bioreactors, rotating drum bioreactors, packed bed bioreactors and others more sophisticated. Regarding examples of submerged and biofilm fermentations, the different developmental

stages observed depend on the processes used to obtain the final product desired at the end of the process. Some examples are outlined in **figures 1A, 2A and 3A**. The figures confirmed that the biomass that can be accumulated on the support was higher in biofilm reactor (**Figure 3B**) than its equivalent in submerged reactor, pointing out a possible way of bioprocess intensification (**Figure 1A**). The same conclusion has been outlined by Seye et al. (2014) concerning immobilisation of microorganisms on support used in fungal biofilm reactor for production of conidia using *Aspergillus clavatus*. The choice of a bioreactor for a given fermentation is based on the production or productivity yields and the regulation of the system required for the filamentous fungi's growth. Regarding bioreactors for solid-state fermentation at the pilot and industrial levels, a few categories can be distinguished based on the aeration and mixing strategies. These are: forced or unforced aeration; static, pulsed mixed, or continuously mixed.

Among cultivation systems, while solid-state fermentation exhibits a higher productivity of secondary metabolites, Kumar et al. (1987) as well as Hölker et al. (2004) compared submerged with liquid-surface fermentations and pointed to some inherent disadvantages, such as poor heat dissipation. Slow diffusion rates of nutrients, products, water, and oxygen in packed beds have also been reported (Viccini et al., 2001; Bellon-Maurel et al., 2003; Durand, 2003; Mitchell et al., 2003; Mo et al., 2004). Muffler et al. (2014) pointed out that the relevance of the emerging field of biofilm research in biotechnology becomes particularly obvious if one considers the number of publications per year for the terms “biofilm” and “biotechnology” collected by the Web of Knowledge (Thomson Reuters) in June 2013. Even though only a small fraction of the number of papers is directly focused on productive biofilms, the illustration shows the prosperous field of this research area.

2.4. Strategies for enhancing the production of fungal metabolites

By considering the fungal biology found when biomass is attached onto a support, some researchers tried to develop a process by which secondary metabolites could be enhanced according to the needs of the microorganisms. Nevertheless, generally much higher yields were obtained in solid-state fermentation. They used, for example, submerged cultures in flasks and/or a stirrer tank (Gaspar et al., 1997; Gaspar et al., 1998; Vavilova et al., 2003; Seyis et al., 2005; Ahamed et al., 2008; Meshram et al., 2008), or solid-state fermentation (Bakri et al., 2003; Sandhya et al., 2005; Assamoi et al., 2008a; Assamoi et al., 2008b; Mitchell et al., 2010) to produce the compounds under study.

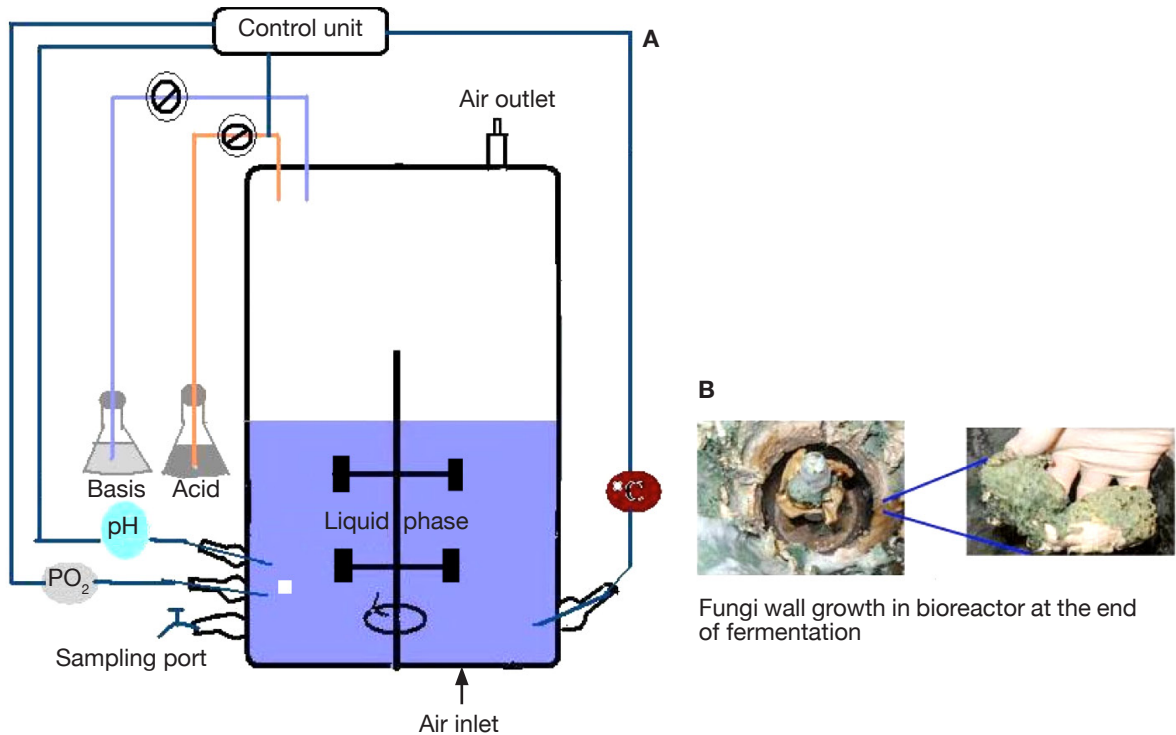


Figure 1. Scheme of a bioreactor used in submerged fermentation and illustration of wall growth fungi — *Schéma d'un bioréacteur utilisé en culture submergée et illustration de la croissance du micro-organisme sur les parois de la cuve.*

A: classical stirred tank bioreactor in which two or three identical turbines were mounted on the central shaft — *bioréacteur agité classique équipé d'un axe comprenant deux à trois mobiles d'agitation identiques*; **B:** wall growth of fungi in submerged bioreactor — *biomasse sur les parois en culture submergée.*

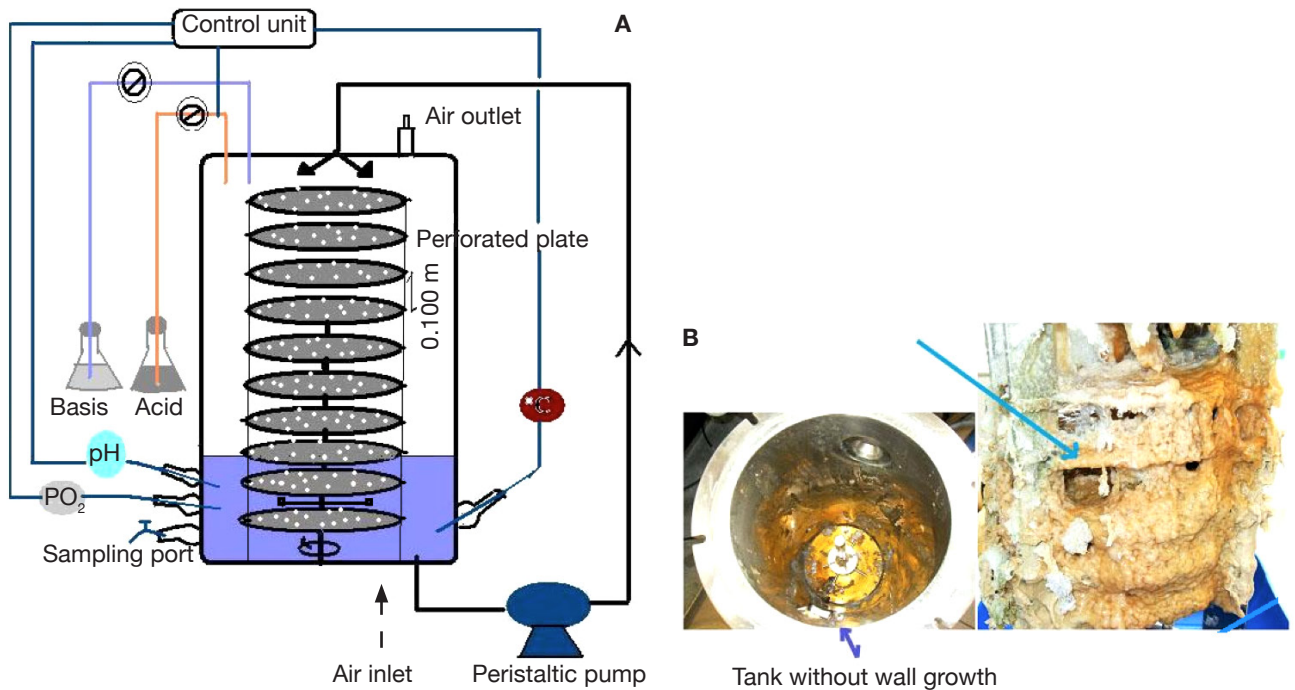


Figure 2. Scheme of a bioreactor used in biofilm fermentation and illustration of the biomass accumulated on the support used — *Schéma d'un bioréacteur utilisé en culture à biofilm et la biomasse accumulée sur les plateaux perforés utilisés.*

A: perforated plate bioreactor adapted from a mechanically stirred tank in which ten perforated stainless steel plates were mounted at a regular distance and a flowing medium system adopted — *bioréacteur à plateaux perforés avec système d'aspersion*; **B:** biomass accumulated on the plates used in a perforated plate bioreactor — *illustration de la biomasse accumulée sur les plateaux perforés utilisés.*

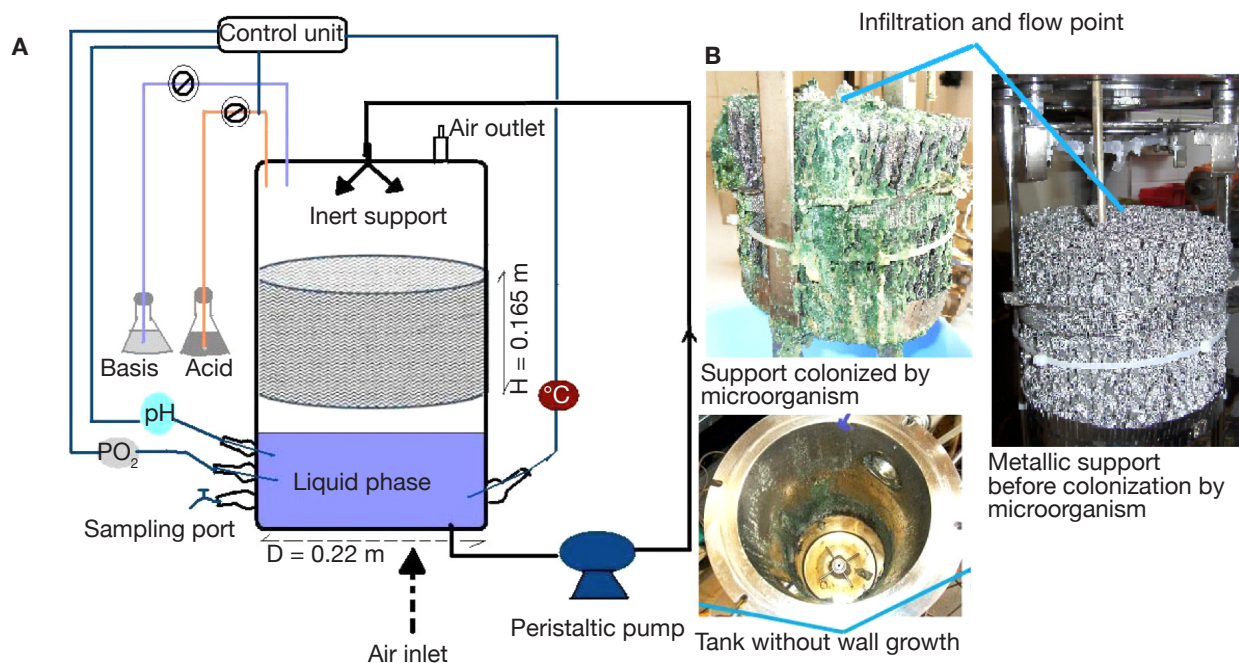


Figure 3. Scheme of a bioreactor used in biofilm fermentation and illustration of the biomass accumulated on support bioreactors with biofilm fermentation of *Trichoderma harzianum* — Schéma d'un bioreacteur utilisé en culture à biofilm avec support inerte et illustration de la biomasse formée en culture submergée et biofilm.

A: Biofilm bioreactor with inert support adapted form a mechanically stirred tank to which an inert support can be added and medium circulated in system during fermentation — bioréacteur à biofilm avec système de circulation du milieu aspergé sur le garnissage métallique structuré; **B:** biomass accumulated on an inert support and an empty one — incrustation de biomasse dans le garnissage métallique structuré et un autre sans dépôt.

Hölker et al. (2004) reported that solid-state fermentation is currently the best method for obtaining fungal conidia from aerial hyphae. The properties of conidia produced by solid-state fermentation differ distinctly from those obtained by submerged fermentation. Fungal conidia, used as biocontrol agents against fungal plant pathogens, are preferentially produced in solid-state fermentation; the conidia obtained manifest a higher quality. They are more resistant to desiccation and more stable in the dry state. To obtain a high number of conidia, a combination of submerged fermentation (for biomass production in the first step) and solid-state fermentation (for subsequent conidia production) proved to be successful. However, a direct comparison between solid-state fermentation and submerged fermentation is very difficult due to the different consistencies of the fungal cultures used in the two technologies. In this regard, biofilm cultures should represent a good objective due to the fact that aerial growth with a flowing medium, which seems to be similar to their natural development, can result in continuous high density production of secondary metabolites and conidia.

3. APPLICATIONS INVOLVING *TRICHODERMA*, *PENICILLIUM* AND *ASPERGILLUS*

According to their capability of producing different compounds, the three genera, *Trichoderma*, *Penicillium* and *Aspergillus* are used in many industrial applications. These mycofactories can be used for two classes of applications *i.e.* either the production of secondary metabolites, or the production of conidia for biocontrol applications.

3.1. Cultivation of *Trichoderma* for the production of fungal metabolites

The filamentous fungus *Trichoderma* has been cultivated using submerged, solid-state and biofilm fermentation in order to produce secondary metabolites as well as conidia. The molecules which have been taken into account were cyclosporin, 6-pentyl- α -pyrone, cellulase, hemicellulase and amylase.

Vinale et al. (2008) showed that MVOCs of the fungus *Trichoderma* act antibiologically against pathogenic plant moulds and can confer plant growth

promoting effects as well as systemic resistance to plants, thus rendering plants less vulnerable to fungal pathogens (Harman et al., 2004).

A number of studies have been conducted to evaluate the effect of nutritional factors (Yong et al., 1986; Serrano-Carreón et al., 1992) on solid and liquid fermentation methods (Kalyani et al., 2000; Sarhy-Bagnon et al., 2000), on 6-pentyl- α -pyrone production. Particular attention has been accorded to enzymes with high xylanolytic activity produced by *Trichoderma* spp. (Wong et al., 1992). Various fermentation systems have been used with *Trichoderma harzianum* Rifai, *Trichoderma viride* Persoon and *Trichoderma koningii* Oudemans to produce these compounds, such as submerged (Cutler et al., 1986; Simon et al., 1988; Evidente et al., 2003), aqueous two-liquid-phase (Rito-Palomares et al., 2000; Rito-Palomares et al., 2001), organic-aqueous two-liquid-phase (Serrano-Carreón et al., 2002; Rocha-Valadez et al., 2006), liquid-surface (Kalyani et al., 2000), and solid-state cultivation systems (Cooney et al., 1997a; Cooney et al., 1997b).

Shinobu et al. (2009) developed an extractive fermentation system with characteristics of both a liquid surface immobilization (LSI) system and a liquid-liquid interface bioreactor (L-L IBR). The system is tentatively named by the authors an “extractive liquid surface immobilization” (Ext-LSI) system. In the Ext-LSI system, the fungus-microsphere mat produces hydrophobic secondary metabolites using nutrients contained in a liquid medium. It is expected that the hydrophobic metabolites are spontaneously extracted from the fungal cells into an organic phase where they accumulate to high concentrations because feedback inhibition and cellular toxicity of the metabolites may be effectively alleviated. The reported maximum accumulations of 6-pentyl- α -pyrone (6PP) in submerged, liquid surface, and solid-state fermentations were 474 mg·l⁻¹ (Serrano-Carreón et al., 2004), 455 mg·l⁻¹ (Kalyani et al., 2000) and 3 g·kg⁻¹ (de Araujo et al., 2002), respectively. Shinobu et al. (2009) also tried to apply the Ext-LSI system with a fungus to produce an aromatic coconut compound. Following optimization of the conditions to produce this volatile, the accumulation of the metabolite in the organic phase reached 7.1 g·l⁻¹ during a four week cultivation. These examples proved that the production of secondary metabolites can be enhanced by using solid-state fermentation.

Looking at the relationship between secondary metabolites and conidiation, Calvo et al. (2002) suggested three kinds of relations that can be considered:

- metabolites that activate conidiation (for example, the linoleic acid derived compounds produced by *Aspergillus nidulans*);
- pigments required for conidiation structures (for example, melanins required for the formation or

integrity of both sexual and asexual conidia and overwintering bodies);

- toxic metabolites secreted by growing colonies at the approximate time of conidiation (for example, the biosynthesis of some deleterious natural products, such as mycotoxins).

Table 1 summarizes some examples of secondary metabolites produced by microorganisms with commercial importance, like antibiotics, organic acids and enzymes.

The term “secondary metabolite” includes a heterogeneous group of chemically distinct natural compounds possibly related to survival functions for the producing organism such as competition (against other micro- and macroorganisms), symbiosis, metal transport, differentiation, etc. (Demain et al., 2000). Included in this group are antibiotics, which are natural products that can inhibit microbial growth. Antibiotic production is often well correlated with biocontrol ability, and the application of purified antibiotics has been found to show positive effects on the host pathogen similar to those obtained using the corresponding living microorganism. Ghisalberti et al. (1990) demonstrated that the biocontrol efficacy of *Trichoderma harzianum* Rifai isolates against *Gaeumannomyces graminis* var. ‘*tritici*’ is related to the production of pyrone like antibiotics.

Trichoderma spp. produces a plethora of secondary metabolites with biological activity (Ghisalberti et al., 1991; Sivasithamparam et al., 1998). The production of secondary metabolites by *Trichoderma* spp. is strain dependent and includes antifungal substances belonging to a variety of chemical classes that were classified by Ghisalberti et al. (1991) into three categories:

- volatile antibiotics, for example 6-pentyl- α -pyrone and most of the isocyanide derivatives;
- water-soluble compounds, like heptelidic acid or koningic acid;
- peptaibols, which are linear oligopeptides of 12–22 amino acids rich in α -aminoisobutyric acid, N-acetylated at the N-terminus and containing an amino alcohol (Pheol or Trpol) at the C-terminus (Le Doan et al., 1986; Rebuffat et al., 1989).

The results obtained from experiments conducted by some researchers suggest that microorganisms produce secondary metabolites under different bioprocesses (**Table 2**). Both mutant and wild strains were used. In some cases, a mutant gave a higher production than the wild strain. The particular bioprocess used also had an impact on enhancing the production.

Polizeli et al. (2005) reported that in either submerged or solid-state fermentation could be used to generate xylanase production by microorganisms.

Table 1. Some biocompounds (antibiotics, organic acids and enzymes) produced by filamentous fungi appreciated for their commercial value — *Quelques composés (antibiotiques, acides organiques et enzymes) produits par les champignons filamenteux appréciés pour leur importance commerciale.*

N°	Fungal biocompounds	Filamentous fungi in used
A Antibiotic		
Fungal strains		
1	Cyclosporins	<i>Trichoderma polysporum</i> Gams
2	Echinocandin B	<i>Aspergillus nidulans</i> Winter, <i>Aspergillus rugulosus</i> Thom and Raper.
3	Griseofulvin	<i>Penicillium aurantiogriseum</i> Dierckx, <i>Penicillium griseofulvum</i> Dierckx, <i>Penicillium italicum</i> Wehmer
4	Patulin	<i>Penicillium urticae</i> Bainier
5	Penicillins	<i>Aspergillus flavus</i> Link, <i>Aspergillus giganteus</i> Wehmer, <i>Aspergillus nidulans</i> Winter, <i>Aspergillus niger</i> van Tieghem, <i>Aspergillus oryzae</i> (Ahlburg) Cohn, <i>Aspergillus parasiticus</i> Spaere, <i>Aspergillus</i> sp., <i>Penicillium baculatum</i> Westling, <i>Penicillium chrysogenum</i> Thom, <i>Penicillium turbatum</i> Westling
B Organic acid		
Fungal strains		
6	Citric acid	<i>Aspergillus citricus</i> (Wehmer) Mosseray, <i>Aspergillus clavatus</i> Desm., <i>Aspergillus niger</i> van Tieghem, <i>Aspergillus phoenicis</i> Wisconsin, <i>Penicillium decumbens</i> Thom, <i>Penicillium isariiforme</i> Stolk and Meyer, <i>Trichoderma viride</i> Persoon
7	Gluconic acid	<i>Aspergillus carbonarius</i> (Bainier) Thom, <i>Aspergillus niger</i> van Tieghem, <i>Aspergillus oryzae</i> (Ahlburg) Cohn, <i>Aspergillus wentii</i> Wehmer, <i>Penicillium chrysogenum</i> Thom, <i>Penicillium luteum</i> Zukal, <i>Penicillium simplicissimum</i> (Oudemans) Thom
8	Itaconic acid	<i>Aspergillus itaconicus</i> Kinoshita, <i>Aspergillus terreus</i> Thom
9	Kojic acid	<i>Aspergillus candidus</i> Link, <i>Aspergillus flavus</i> Link, <i>Aspergillus oryzae</i> (Ahlburg) Cohn, <i>Aspergillus parasiticus</i> Spaere, <i>Aspergillus tamarii</i> Kita, <i>Penicillium jensenii</i> Zalessky
10	L-Malic acid	<i>Aspergillus atroviolaceus</i> Mosseray, <i>Aspergillus citricus</i> (Wehmer) Mosseray, <i>Aspergillus flavus</i> Link, <i>Aspergillus niger</i> van Tieghem, <i>Aspergillus ochraceus</i> Wilhelm, <i>Aspergillus oryzae</i> (Ahlburg) Cohn, <i>Aspergillus wentii</i> Wehmer
11	D-Araboascorbic acid	<i>Penicillium chrysogenum</i> Thom
12	Erythorbic acid	<i>Penicillium griseoroseum</i> Dierckx
C Enzyme		
Fungal strains		
13	α -Amylase	<i>Aspergillus niger</i> van Tieghem, <i>Aspergillus oryzae</i> (Ahlburg) Cohn, <i>Aspergillus awamori</i> Nakazawa, <i>Trichoderma viride</i> Persoon
14	Amyloglucosidase	<i>Aspergillus niger</i> van Tieghem, <i>Aspergillus oryzae</i> (Ahlburg) Cohn, <i>Aspergillus awamori</i> Nakazawa, <i>Aspergillus phoenicis</i> Wisconsin
15	Catalase	<i>Aspergillus niger</i> van Tieghem, <i>Penicillium vitale</i> Pidoplichko and Bilai
16	Cellulase	<i>Aspergillus niger</i> van Tieghem, <i>Aspergillus soyae</i> Sakagawa and K. Yamada, <i>Aspergillus terreus</i> Tohm, <i>Penicillium citricum</i> Thom, <i>Penicillium funiculosum</i> Thom, <i>Trichoderma longibrachiatum</i> Rifai, <i>Trichoderma reesei</i> Rifai, <i>Trichoderma viride</i> Persoon
17	Dextranase	<i>Aspergillus carneus</i> (van Tieghem) Blockwitz, <i>Penicillium funiculosum</i> Thom, <i>Penicillium lilacinum</i> Thom, <i>Penicillium pinophilum</i> Hedgecock
18	α -Galactosidase	<i>Aspergillus awamori</i> Nakazawa, <i>Aspergillus niger</i> van Tieghem, <i>Aspergillus oryzae</i> (Ahlburg) Cohn, <i>Penicillium dupontii</i> Griffon and Maubl.
19	β -Galactosidase	<i>Aspergillus awamori</i> Nakazawa, <i>Aspergillus niger</i> van Tieghem, <i>Aspergillus nidulans</i> Winter, <i>Aspergillus oryzae</i> (Ahlburg) Cohn, <i>Penicillium funiculosum</i> Thom
20	β -Glucanase	<i>Aspergillus niger</i> van Tieghem, <i>Aspergillus oryzae</i> (Ahlburg) Cohn, <i>Trichoderma viride</i> Persoon, <i>Trichoderma reesei</i> Rifai
21	Glucoamylase	<i>Aspergillus niger</i> van Tieghem, <i>Aspergillus awamori</i> Nakazawa, <i>Aspergillus oryzae</i> (Ahlburg) Cohn

Table 1 (continued). Some biocompounds (antibiotics, organic acids and enzymes) produced by filamentous fungi appreciated for their commercial value — *Quelques composés (antibiotiques, acides organiques et enzymes) produits par les champignons filamenteux appréciés pour leur importance commerciale.*

N°	Fungal biocompounds	Filamentous fungi in used
C	Enzyme	Fungal strains
22	Glucose aerohydrogenase	<i>Aspergillus niger</i> van Tieghem
23	Glucose oxidase	<i>Aspergillus niger</i> van Tieghem, <i>Penicillium amagasakiense</i> Kusai, <i>Penicillium simplicissimum</i> (Ouedmans) Thom, <i>Penicillium vermiculatum</i> Dangeard
24	α -Glucosidase	<i>Aspergillus awamori</i> Nakazawa, <i>Aspergillus flavus</i> Link, <i>Aspergillus fumigatus</i> Fresenius, <i>Aspergillus niger</i> van Tieghem
25	α -D-Glucosidase	<i>Aspergillus niger</i> van Tieghem
26	β -Glucosidase	<i>Aspergillus niger</i> van Tieghem, <i>Aspergillus oryzae</i> (Ahlburg) Cohn, <i>Trichoderma reesei</i> Rifai
27	Hemicellulase	<i>Aspergillus niger</i> van Tieghem, <i>Aspergillus oryzae</i> (Ahlburg) E. Cohn, <i>Aspergillus phoenicis</i> Wisconsin, <i>Trichoderma longibrachiatum</i> Rifai, <i>Trichoderma viride</i> Persoon
28	Hesperidase	<i>Aspergillus niger</i> van Tieghem
29	Invertase	<i>Aspergillus awamori</i> Nakazawa, <i>Aspergillus niger</i> van Tieghem, <i>Aspergillus oryzae</i> (Ahlburg) Cohn
30	Lipase	<i>Aspergillus niger</i> van Tieghem, <i>Aspergillus oryzae</i> (Ahlburg) Cohn, <i>Penicillium roqueforti</i> Thom
31	Pectinase	<i>Aspergillus alliaceus</i> Thom and Church., <i>Aspergillus niger</i> van Tieghem
32	Phytase	<i>Aspergillus niger</i> van Tieghem, <i>Aspergillus ficuum</i> (Reichardt) Henn.
33	Protease	<i>Aspergillus niger</i> van Tieghem, <i>Aspergillus melleus</i> Yukawa, <i>Aspergillus oryzae</i> (Ahlburg) Cohn, <i>Aspergillus saitoi</i> , <i>Penicillium dupontii</i> Griffon and Maubl.
34	Tannase	<i>Aspergillus fumigatus</i> Fresenius, <i>Aspergillus versicolor</i> , <i>Aspergillus flavus</i> Link, <i>Aspergillus niger</i> van Tieghem, <i>Aspergillus tamarii</i> Kita, <i>Aspergillus japonicus</i> Saito, <i>Aspergillus parasiticus</i> Spaere, <i>Aspergillus oryzae</i> (Ahlburg) Cohn, <i>Penicillium charlesii</i> Thom, <i>Penicillium crustosum</i> Thom, <i>Penicillium restrictum</i> Gilman and Abbott
35	Xylanase	<i>Trichoderma reesei</i> Rifai

For example, submerged fermentation was used to produce xylanase from *T. reesei*. In that experiment, the optimal pH and temperature were 4.5 and 40 °C. This enzyme is usually used in cellulose pulp bleaching and animal feed.

Dashtban et al. (2011) performed experiments relating to the effect of different carbon sources on cellulose production by *Hypocrea jecorina* (*T. reesei*) strains using *T. reesei* QM6a, *T. reesei* QM9414, and *T. reesei* RUT-C30. Time course experiments showed that maximum cellulose activity with QM6a and QM9414 strains occurred at 120 h for the majority of the tested carbon sources, while RUT-C30 had its greatest cellulose activity at around 72 h. Maximum cellulase production was observed to be 0.035, 0.42 and 0.33 μmol glucose equivalents using microcrystalline celluloses for QM6a, QM9414, and RUTC-30, respectively; while the maximum dry weight for all three *T. reesei* strains was obtained at the highest carbon concentration level (2%). This

suggests that an increase in enzyme production is directly caused by an increase in mycelium density and may be a result of complex inducing factors in a relationship with the bioreactor configuration used.

In the course of experiments carried out in our laboratory, it was observed that during culture, wall growth of the fungi is amplified by the speed of agitation, which splashes some fragments of mycelia on the wall vessel (**Figure 1B**). This phenomenon reduces the biomass in that type of culture. In order to perform the production of biomass and the volatile, a biofilm bioreactor has been adapted, modulating plate bioreactor from the mechanical one with ten perforated stainless steel plates (**Figure 2B**) and also using an inert metallic support named packing (**Figure 3B**) on which biomass can be accumulated without growth on the vessel wall. With these results, it can be suggested that the wall growth in submerged culture will be moderated using the strategies adopted.

Table 2. Production of xylanase and 6-pentyl- α -pyrone by some filamentous fungi using different bioprocesses — Production de xylanase et de 6-pentyl- α -pyrone par les champignons filamenteux en utilisant des bioprocédés variés.

Biocompound	Production	Microorganism	Fermentation applied	Reference
Xylanase	630 IU·ml ⁻¹	<i>Trichoderma reesei</i> Rut C-30	SbmF	Xiong et al., 2004
	1,350 IU·ml ⁻¹	<i>Trichoderma reesei</i> Rut C-30	SbmF (Fed-batch)	Xiong et al., 2004
	14,790 IU·ml ⁻¹ ·h ⁻¹	<i>Aspergillus niger</i> KK2	SSF wheat raw	Park et al., 2002
	844 IU·ml ⁻¹	<i>Penicillium canescens</i> 10-10c	SbmF	Gapar et al., 1997
	7,448 IU·ml ⁻¹	<i>Penicillium canescens</i> 10-10c	SbmF	Bakri et al., 2003
	9,632 IU·g ⁻¹	<i>Penicillium canescens</i> 10-10c	SSF	Bakri et al., 2003
	9,300 IU·g ⁻¹	<i>Penicillium canescens</i> 10-10c	SSF	Assamoi et al., 2008a
	10,200 IU·g ⁻¹	<i>Penicillium canescens</i> 10-10c	SSF	Assamoi et al., 2009
	18,895 IU·g ⁻¹	<i>Penicillium canescens</i> 10-10c	SSF	Assamoi et al., 2009
	6-pentyl- α -pyrone	455 mg·l ⁻¹	<i>Trichoderma harzianum</i> Rifai	LSC
474 mg·l ⁻¹		<i>Trichoderma harzianum</i>	SbmF	Serrano-Carreón et al., 2004
3,000 mg·kg ⁻¹		<i>Trichoderma harzianum</i>	SSF	de Aroujo et al., 2002
7,100 mg·l ⁻¹		<i>Trichoderma atroviride</i> AG2755-5NM398	SbmF Ext-LSI	Shinobu et al., 2009
230 mg·l ⁻¹		<i>Trichoderma harzianum</i> IMI206040	SbmF	Rocha-Valadez et al., 2006
230 mg·l ⁻¹		<i>Trichoderma harzianum</i> IMI206040	SbmF	Rocha-Valadez et al., 2006
5,000 mg·kg ⁻¹		<i>Trichoderma harzianum</i> 4040	SSF	Ramos et al., 2008

SbmF: Submerged Fermentation; SSF: Solid Substrate Fermentation; Ext-LSI: Extractive Liquid-Surface Immobilisation.

The similar system has been used by Khalesi et al. (2014) for the production of hydrophobin HFBII by *T. reesei*. They noted that the use of a biofilm reactor has led to a significant increase of HFBII production in shorter time by comparison with a classical submerged bioreactor.

3.2. Use of *Penicillium* for the production of fungal metabolites and conidia

The main molecules produced by *Penicillium* were griseofluvin, penicillin, citric acid, koji acid, erythorbic acid, catalase, dextranase and tamase. Chávez et al. (2005) reviewed the genus *Penicillium*, particularly the species producing xylanase. They reported that the *Penicillia* were mostly saprophytic in nature, and numerous species are of particular value for humans. The best known are *Penicillium canescens* 10-10c and *Penicillium notatum* Westling, producing xylanase and the antibiotic penicillin, respectively, as well as *Penicillium roqueforti* Thom and *Penicillium camemberti* Thom, which are important in the food industry. The latter are associated with the production of particular types of cheese. A lot of *Penicillia* are soil fungi, and grow in a variety of organic substances, particularly dead plant material. They produce extracellular hydrolases such as pectic enzymes, lipases, proteases, cellulases and xylanases. The latter enzymes are known for a number of biotechnological applications. One important use in cellulose pulp biobleaching (Polizeli et al., 2005) is to eliminate lignin residues from kraft pulp. They also increase the digestibility of feed by lowering the viscosity in the intestinal tract, thus improving nutrient uptake (Twomey et al., 2003). In baking, they are added to increase the specific volume and in this way improve final flavour (Maat et al., 1992). In beer and juice processing, they are used to reduce haze formation by solubilising long chain arabinoxylans (Dervilly et al., 2002).

In studies carried out with *P. canescens* 10-10c, it was concluded

that the production of xylanase is influenced by the process used in its production, whether submerged or solid-state fermentation. The researchers who conducted the experiments in different types of fermentation noted the influence of the solid-state fermentation ($9,632 \text{ IU}\cdot\text{g}^{-1}$) (Gaspar et al., 1997; Bakri et al., 2003), or solid-state fermentation with a multi-layer system ($10,200 \text{ UI}\cdot\text{g}^{-1}$) (Assamoi et al., 2008a; Assamoi et al., 2008b; Assamoi et al., 2009). In order to improve oxygen transfer in *P. canescens* 10-10c culture, Gaspar et al. (1997) studied the influence of the agitation and aeration rates on $K_L a$ in a mono-agitated reactor; the authors concluded that a reduction in specific power is more profitable to xylanase production than an increase in $K_L a$ beyond a critical value. Too little $K_L a$ is harmful to biomass growth. This was verified when production at 200 rpm with DT4 turbine ($P = 102 \text{ W}\cdot\text{m}^{-3}$, $K_L a$ at about 20 h^{-1}) reached only $370 \text{ IU}\cdot\text{ml}^{-1}$. An optimum was found at 300 rpm with FBT8 mobile ($P = \text{about } 232 \text{ W}\cdot\text{m}^{-3}$, $K_L a$ at about 62 h^{-1}), where production reached $844 \text{ IU}\cdot\text{ml}^{-1}$. Bakri et al. (2003) performed experiments in order to evaluate the influence of carbon sources, nitrogen sources, and moisture content on xylanase production by *P. canescens* 10-10c in solid-state fermentation. The experiments started in submerged cultures using Erlenmeyer flasks (250 ml) containing 50 ml of the medium; other experiments were conducted using solid-state culture. The maximum enzyme production obtained was $7,448 \text{ IU}\cdot\text{g}^{-1}$ in submerged culture and $9,632 \text{ IU}\cdot\text{g}^{-1}$ in solid-state culture after 12 days. The yields of xylanase productivity from *P. canescens* 10-10c observed were approximately 1.5 fold higher than optimum productivities reported in the literature for some microorganisms grown in solid-state fermentation. According to the results from Assamoi et al. (2008a), Assamoi et al. (2008b), and Assamoi et al. (2009), xylanase can be produced in solid-state culture. The experiments were carried out either in Erlenmeyer flasks or reactors with *P. canescens* 10-10c. In flasks, it was observed that production was maximal after seven days, $18,895 \text{ IU}\cdot\text{g}^{-1}$. Thereafter, the authors carried out a scale-up in plastic bags and a multi-layer reactor. The results after seven days were $9,300 \text{ IU}\cdot\text{g}^{-1}$ in plastic bags and $10,200 \text{ IU}\cdot\text{g}^{-1}$ in the multi-layer reactor. These results demonstrated that many problems occur in solid-state fermentation as the scale of the process increases; in particular, transport phenomena are strongly affected by the scale of the process. In this case, oxygen transfer inside the plastic bags is expected to be very low. In order to improve the oxygen transfer effectively, it would be interesting, for example, to supplement the plastic bags with a softened forced aeration (low and intermittent humidified air flow rate). Polizeli et al. (2005) summarized data on various commercial xylanases produced in submerged and

solid-state fermentation cultures by microorganisms. It should be noted that about 80-90% of all xylanases are produced in submerged culture. Wheat bran and rice are regarded as inducers.

3.3. Secondary metabolites produced by *Aspergillus* in different bioreactor designs

The fungal metabolites produced by *Aspergillus* were penicillin, citric acid, koji acid, L_malic acid, amylase, catalase, cellulase, galactosidase, glucanase, glucosidase, hemicellulase, lipase, pectinase and protease. de Vries et al. (2001) reported that within the genus *Aspergillus*, comprising a group of filamentous fungi with a large number of species, the most important for industrial applications are some members of the group of black *Aspergilli* (*Aspergillus niger* van Tieghem and *Aspergillus tubingensis* [Schöber] Mosseray). Reclassification using molecular and biochemical techniques resulted in clear distinctions being made between eight groups of black *Aspergilli*: *A. niger*, *A. tubingensis*, *Aspergillus foetidus* (Nakazawa) Thom and Rapper, *Aspergillus carbonarius* (Bainier) Thom, *Aspergillus japonicus* Saito, *Aspergillus aculeatus* Iizuka, *Aspergillus heteromorphus* Batista and Maia, and *Aspergillus ellipticus* Raper and Fennell. Products of several of these species have attained a generally recognized as safe (GRAS) status, which allows them to be used in food and feed applications. Black *Aspergilli* have a number of characteristics that make them ideal organisms for industrial applications, such as good fermentation capabilities and high levels in protein secretion. The results obtained in a comparative study on protease yield by solid-state fermentation or submerged fermentation indicated that among the various agro-industrial residues used, wheat bran was the most suitable substrate for protease synthesis by *A. oryzae* NRRL 1808 in submerged culture as well as solid-state fermentation. The solid-state fermentation showed its superiority for enzyme production over submerged culture, since a comparative evaluation of protease yields by the two fermentation systems showed 3.5-fold greater enzyme production using the solid-state format (Sandhya et al., 2005). A mixture of rice bran, rice husk and gram hull in 5:3:2 ratio was found to be the most suitable, where maximum enzyme production occurred after 72 h and 96 h of incubation under submerged and solid-state fermentation conditions, respectively (Nehra et al., 2002). Production of xylanase by the *A. niger* KK2 mutant in solid-state fermentation reached $14,790 \text{ IU}\cdot\text{l}^{-1}\cdot\text{h}^{-1}$ (Park et al., 2002); the authors remarked during experimentation that the highest production of xylanase ($2,596 \text{ IU}\cdot\text{gds}^{-1}$) was obtained at 40°C after six days; at higher temperatures, its production decreased sharply, suggesting that the end point of fermentation should

be carefully controlled because synthesized xylanase could be degraded by non-specific proteases secreted by the fungus; the maximum value of xylanase activity achieved was 5,071 IU·g⁻¹ of rice straw, which is similar to the 5,484 IU·g⁻¹ of rice straw predicted by the model proposed. Wang et al. (2005) reported that protease could be produced from *Aspergillus oryzae* (Ahlburg) Cohn by either solid-state or submerged fermentation. However, problems exist in both processes. Besides incomplete nutrient utilization and frequent bacterial contamination, solid-state fermentation also suffers from difficulties in establishing a constant moisture content, evenly distributed aeration and effective removal of metabolic heat. In submerged fermentation, filamentous fungi exhibit various morphologies, ranging from dispersed mycelia to pellets. In some cases, dispersed growth benefits enzyme production, but in a medium of low viscosity; it is difficult to achieve absolutely dispersed mycelia, although fungal morphology varies with broth pH, inoculum size and quality, dissolved oxygen tension, mixing intensity, polymer additives and surfactant addition. For the production of xylanase, Beg et al. (2001) reported that the advantages of solid-state fermentation processes over liquid-batch fermentation include smaller volumes of liquid required for product recovery, cheap substrate, low cultivation cost for fermentation, and low risk of contamination in addition to high enzyme yield. Seye et al. (2014) carried out experiments with a fungal biofilm reactor combining the technological advantages of submerged fermentation (*i.e.*, free water facilitating the control of culture parameters) with the biological characteristics found in solid-state fermentation in order to establish effect of insecticidal activity of conidia and metabolites secreted by an entomopathogenic *A. clavatus* strain in this culture system against *Culex quinquefasciatus* larvae and adults. They concluded that the process allowed facility in recuperation and purification of conidia (confined on the solid substrate) and metabolites (contained in the liquid medium). These results confirm the usefulness of the system, which can be used in similar production for other fungi. Submerged fermentation is used more often than solid-state fermentation even this solid-state gives high production of biomolecules and conidia. Another bioreactor type that can enhance production of fungal metabolites and conidia in comparison with those produced in submerged culture is biofilm bioreactor which simulates solid-state. Gutiérrez-Correa et al. (2012) discussed the main biological mechanisms that support the development of filamentous fungal biofilms and the recent advances on their use in industrial processes with an emphasis on *Aspergillus* biofilms. They concluded that cell adhesion appears to be the trigger for biofilm formation and its main industrial features such as increased production of

metabolites and enzymes. In this sense, a great deal of information has been gained relating to the industrial uses of biofilms from *Aspergillus* and other filamentous fungi, which were originally considered as derived from cell immobilization systems. Efforts to seek an agreement are required to differentiate filamentous fungal biofilms from those cell immobilization systems based on “natural” or “passive” forces, since this would encourage deep research to understand gene regulation systems and behaviour in such a way to achieve its optimization, scale-up, and industrial production.

4. CONCLUSIONS

The aim of this paper was to highlight the process used for optimizing the production of biomolecules secreted by filamentous fungi. Traditional methods used for the production of enzymes and secondary metabolites have been focused on solid-state fermentation without controlling factors essential for microorganism growth. However, submerged fermentation is more widely used, considering the possibility to implement robust control loops allowing to ensure the reproducibility of the process. The cost of secondary metabolite is one of the main factors determining the economics of process. In some cases, solid-state fermentation offers advantages over submerged fermentation, but in general submerged cultures are used most often since the parameters can be controlled efficiently. As filamentous fungi are naturally adapted to grow on surfaces, under which conditions they show a particular physiological behaviour different from that observed in submerged fermentation, another bioreactor process named “biofilm” that can enhance the production of fungal metabolites using a synthetic support within a liquid environment should also be used in order to enhance the production of biomolecules needed. This biofilm reactor can be considered as a promising technology combining the advantages of submerged fermentation in terms of process control and the advantages of solid-state fermentation in terms of fungal biology and development.

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