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REVIEW ARTICLE

Biobutanol Production from Plant Biomass

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Abstract:

Microbiological conversion of biosphere renewable resources to produce useful products, in particular biofuels, is currently one of the pressing problems of biotechnology. To establish a microbiological production of biobutanol at an industrial scale, strains with high-yield solvent production on plant biomass as a cheap substrate are needed.

This paper summarizes the main outcomes of the authors' original research focused on a) obtaining new butanol-producing strains of *Clostridium* genus, b) testing different sources of non-food raw material as a substrate for fermentation. A comparison of different methods of biomass pretreatment and their efficiency for the accumulation of butanol in the liquid medium is also reported.

In particular, the efficiency of butanol production by *C. acetobutylicum* strains isolated or mutagenized by the authors on a) ground green rapeseed, switchgrass, sweet sorghum, soybean, wheat biomass; b) components of switchgrass after thermobaric hydrolysis and c) paper mill sludge from the pulp as substrates is reported. This paper also highlights the progress made concerning substrate pre-treatment and optimization of cultivation conditions to increase butanol production. Finally, future directions to optimize the different biotechnological steps leading to butanol production are discussed.

Keywords: Biobutanol, Biotechnology, *Clostridium*, Plant biomass, Strain producer, Butanol production.

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1. INTRODUCTION

Biobutanol is considered one of the most promising renewable biofuels [1]. Due to its high calorific value and low hydrophilicity, butanol is more like gasoline than ethanol. However, a large scale industrial production of biobutanol is still restrained by low efficiency of its production technology [2]. Butanol is a principal end product of acetone-butanol-ethanol (ABE) fermentation by certain species of anaerobic, spore-forming gram-positive microorganisms. Among them, *Clostridium acetobutylicum* can mostly produce butanol and acetone.

Biobutanol, as well as bioethanol, can be obtained by both fermentations of traditional sugar- or starch-containing raw materials (1st generation biobutanol) and fermentation of lignocellulosic raw materials from biomass (2nd generation biobutanol). Biobutanol produced in the process of ABE fermentation of biomass by *Clostridia* spp. has the same characteristics as the butanol obtained by chemical synthesis

[3, 4]. The main factor influencing the profitability of the traditional ABE fermentation is the relatively high cost of starch- (corn, wheat, millet, *etc.*) and sugar- (molasses, sorghum, *etc.*) containing substrates [5 - 9]. It is this factor and the ability of saccharolytic *Clostridia* to use various carbohydrates as substrates that have triggered the research for the production of biobutanol from alternative and renewable raw materials [10, 11].

Earlier works [12 - 15] on fermentation of various carbohydrates for the production of acetone and butanol demonstrated that some sugars such as glucose, fructose, mannose, sucrose, lactose, starch and dextrose are utilized comprehensively, while galactose, xylose, arabinose, raffinose, melezitose, inulin and mannitol are utilized only partially, and trehalose, rhamnase and melibiose are virtually unutilized. It is known that lignocellulose is a principal structural component of numerous agricultural and wood waste products. A significant amount of residual biomass can potentially be recycled for a variety of products, including biofuel [16]. Bioconversion of lignocellulosic waste for the production of biofuel could make a significant contribution to overcoming

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numerous environmental and economic problems [17 - 26].

Over the past two decades, significant progress has been made in the butanol biotechnology, especially in the cultivation process through (1) butanol-titer improvement, (2) by-products isolation or elimination, (3) increased tolerance of bacteria to butanol concentration and (4) utilization of not only pure sugars, but also lignocellulosic biomass hydrolysates as substrates for bacteria. The use of renewable raw materials - lignocellulosic biomass - and the possibility of partial or complete replacement of food raw materials with non-food raw materials is important progress on the way to establishing production of biobutanol at an industrial scale [27 - 31]. Here we review recent advancements on the use of non-food plant raw materials as a substrate for fermentation and compare effectiveness of different methodologies of green biomass pretreatment for the accumulation of butanol in a liquid medium.

2. SUBSTRATE AND ITS COMPONENTS

Lignocellulose, as the main component of almost all plant biomass, is the most common renewable resource and practically unlimited cultivation substrate for the production of biofuels and other valuable chemicals [11, 14, 16, 17]. Lignocellulose is composed of lignin, hemicellulose and cellulose. The percentage ratios of the lignocellulose components may vary depending on the type of raw material. Each of the lignocellulose components (hemicellulose, cellulose and lignin) can, if properly treated, be used in the production of biofuels.

Hemicelluloses are heteropolysaccharides commonly found in woody parts of plants, cereal straw with cellulose [32, 33]. Depending on the content of hemicelluloses of certain monosaccharides, they are called mannans, galactans or pentosans (xylans, arabans). The most important and common in nature are pentoses - arabinose and xylose, and especially hexoses - glucose, fructose, galactose, mannose. Hemicellulose can be split by xylase enzymes to form pentose sugars. Some *Clostridia* sp. synthesize xylanase and therefore are able to use xylan, a major component of hemicellulose, as an essential carbon source. Hemicelluloses are decomposed by three sets of enzymes - the first set includes xylanase, mannanase, galactase, arabinase, the second and the third sets contain cellulase and β -glucanase, respectively [11, 14, 16, 17, 33].

Cellulose is a linear homopolysaccharide of β -D-glucopyranose in the form of long filaments joined by β -1,4-glycosidic bonds [33, 34]. Its structural unit is disaccharide cellobiose made up of β -glucose residues. The association of cellulose molecules in a stable crystalline structure led to the formation of microfibrils with 15 - 45 chains. At the microscopic level, there are microfibrils that form a complete fibril. Cellulose in the paracrystalline structure contains both amorphous and crystalline regions. The degree of polymerization is one of the main factors affecting cellulose bioconversion.

Cellulose is digested by cellulase, which acts to split molecules of cellulose into cellobiose and finally to glucose.

Cellulose chains are very stable due to the interchain hydrogen and van der Waals interaction between pyranose rings that form crystalline regions [35]. The cellulose crystalline regions are interspersed with amorphous ones, which are degradable much easier. The cellulases responsible for the hydrolysis of cellulose are composed of a complex of proteins that have different specific activities for the hydrolysis of glycosidic bonds. Depending on their specific activities, cellulases are divided into three main classes: endoglucanases, cellobiohydrolases (exoglucanases) and β -glucosidases. Endoglucanases accidentally attack the cellulose chains in the unprotected positions of the amorphous region and create new ends, while the exoglucanases digest the polymer chain, both from the reducing and non-reducing ends, producing the main product - cellobiose. Endoglucanases exhibit high specific activity against soluble cellulose derivatives, such as carboxymethylcellulose, and low (or none at all) activity for microcrystalline cellulose as compared to exoglucanases. Synergistic action of both enzymes is required for the efficient cellulose degradation [35].

Lignin, an other component of lignocellulolytic biomass, has only recently attracted the interest of scientists. Lignin is a natural polymer that is located in cell walls and intercellular space and holds cellulose fibers [11, 14, 16, 17, 33, 36, 37]. Lignin can be isolated from plant tissues in various ways. It dissolves in both acids and alkalis. Protolignin is almost insoluble in organic solvents. Lignin, if isolated in different ways, differs in its composition and properties. In the chemical sense, lignin is a general and conditional concept - there can be no two fully identical lignins in nature. Numerous lignin formulas can be found in the literature [36, 37], but International Lignin Institute recommended to use the structural lignin formula [11]. As a rule, a lignin molecule is indefinitely large and contains multiple functional groups. A common structural constituent of all types of lignin is phenyl propane derivatives. According to the modern concepts, lignin is a complex three-dimensional polymer of aromatic nature, formed as a result of polycondensation of several monolignols - cinnamyl alcohols (paracumarone, coniferyl and sinapic).

Modern methods of analysis allow to determine with sufficient accuracy the constituencies (individual macrocomponents) that are part of plant biomass [27]. In order to identify the carbon sources that are part of the biomass, the components of switchgrass (*Panicum virgatum*), rapeseed (*Brassica napus*), sweet sorghum (*Sorghum saccharatum*), wheat (*Triticum aestivum*), soybean (*Glycine max*) and paper mill sludge were determined Fig. (1) and published earlier [2, 27, 31, 38 - 40]. The green biomass of soybean, rape and wheat was provided by Institute for Agricultural Engineering and Electrification (National Academy of Agrarian Sciences of Ukraine), switchgrass and sweet sorghum - by M.M. Hryshko National Botanic Garden (National Academy of Sciences of Ukraine), components of switchgrass (cellulose, arabinogalactan and lignin) after thermobaric hydrolysis - by V.P. Kukhar Institute of Bioorganic Chemistry and Petrochemistry (National Academy of Sciences of Ukraine) and paper mill sludge - by Kyiv Cardboard and Paper Mill.

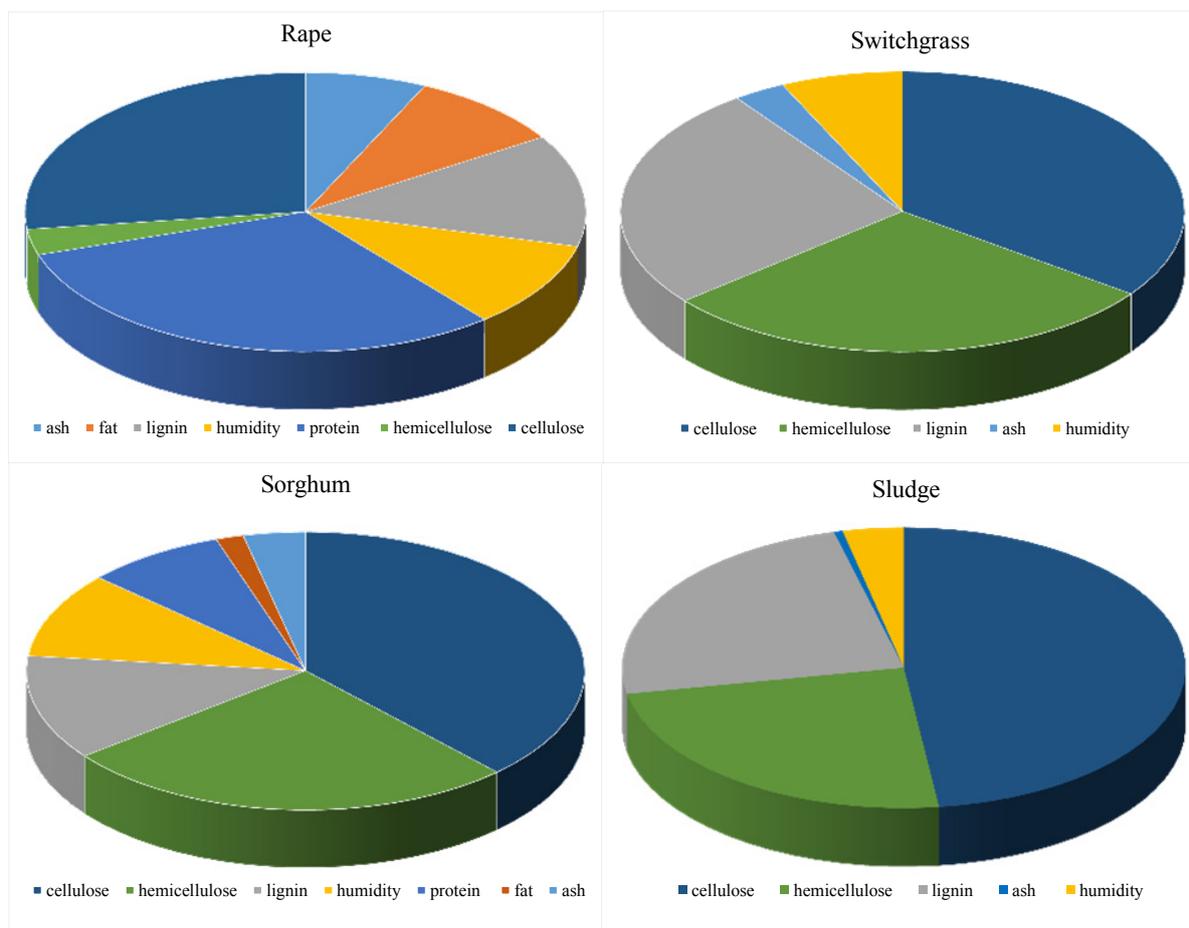


Fig. (1). The components of switchgrass, rape, sorghum biomass and sludge.

The thermobaric hydrolysis of lignocellulosic raw materials was performed with specially constructed equipment at a temperature of 200 C for 10-15 min [31]. Dry ground switchgrass biomass (8% humidity) was placed in a reactor, some part of the water was added and hydrolysis was performed. The result was a mushy mixture, which was either immediately used as a substrate for the fermentation or fractionated according to the scheme shown in Fig. (2). Cellulose (35%), hemicellulose (29%) and lignin (26%) were found to be the main components of the biomass of switchgrass; for rape - protein (31%), cellulose (27%), hemicellulose (3%) and lignin (13%); for sludge - cellulose (48%), lignin (24%) and hemicellulose (24%) [31, 40]. The identified biomass composition allows to deduce that *Clostridium* bacteria are able to grow and convert the biomass of these species [2]. Therefore, in all these experiments, the butanol-producing strains *C. acetobutylicum* IMB B-7407 (IFBG C6H - parental strain) and *Clostridium* sp. IMB B-7570 (IFBG C6H 5M - mutant strain) from the collection of SI Institute of Food Biotechnology and Genomics (hereinafter referred to as IFBG Collection) were used.

3. PRODUCING STRAINS

Different strains of *C. acetobutylicum* absorb the vast majority of sugars present in the hemicelluloses of wood and its hydrolysates, in particular, xylose, which is the main component of hemicellulose. The bioconversion of xylose by *C. acetobutylicum* was investigated on dried grass [41 - 43]. The bioconversion of β -glucans and xylans was shown by *C. acetobutylicum*-6, *C. acetobutylicum*-7 and *C. acetobutylicum* VKPM B-4786 strains which had high hemicellulase activity. Such conversion increased with an increasing proportion of dry grass in the cultivation medium [44, 45].

Isolation of new strains of *C. acetobutylicum* is essential for efficient bioconversion of cellulosic waste. As reported previously [46 - 49], 20 strains of *Clostridium* spp. were screened for their cellulolytic activity and only two *C. acetobutylicum* strains, namely NRRL B527 and ATCC 824, demonstrated cellulolytic activity and the ability to produce extracellular enzymes. Previous studies [15] explored the possibility of producing solvents (butanol, methanol and ethanol) from cellulosic biomass using the *C. acetobutylicum* P 262 strain and suggested technology for butanol production from cellulose.

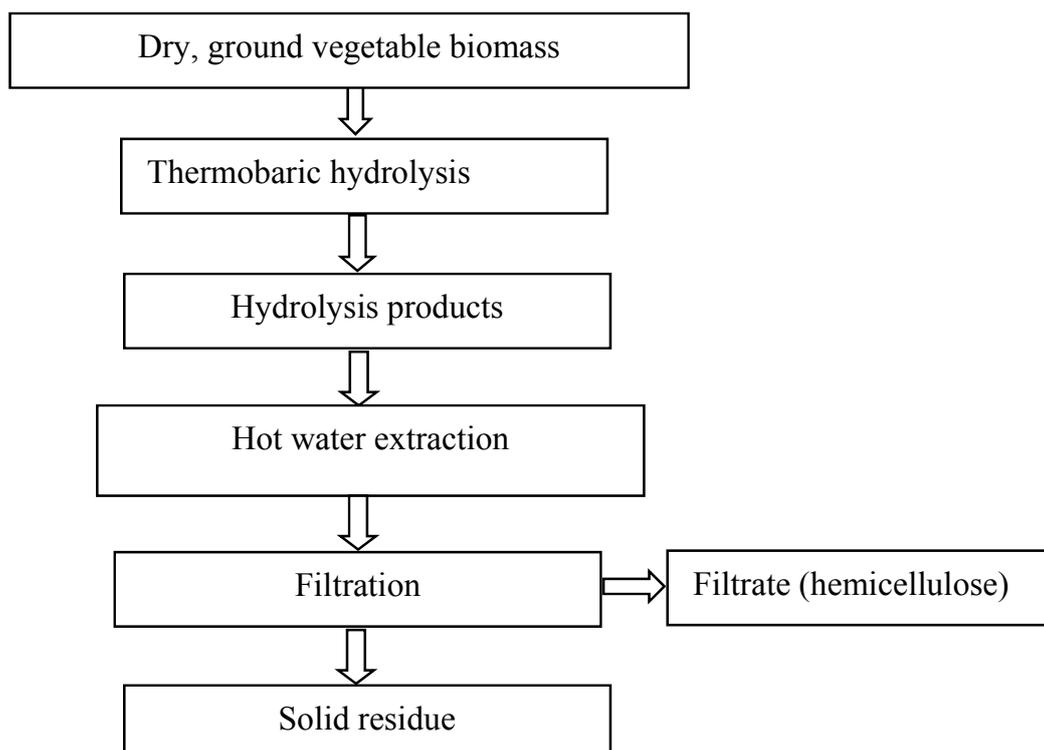


Fig. (2). Scheme of fractionation of lignocelluloses biomass after thermobaric autohydrolysis.

Some authors have also shown that the substrate composition influences the composition and enzymatic activity of *C. cellulovorans* cellulosome [50 - 52]. Bacterial genes encoding cellulase can be either randomly distributed, as in *C. thermocellum*, or clustered on the chromosome, as in *C. cellulovorans*, *C. cellulolyticum* and *C. josui* [53 - 57]. Other authors have identified the genes that encode putative cellulolytic components in a chromosome of *C. acetobutylicum* ATCC 824 strain [58].

To summarise our findings, it can be concluded that currently, the main obstacle to the commercialization of the cellulose enzymatic hydrolysis is the high cost of enzymes. This is further complicated by the fact that the low activity of cellulases requires a large number of lignocellulolytic enzymes [59]. In our view, the strategies to improve the economic feasibility of the hydrolysis of raw materials can be divided into four groups:

- (1) Screening organisms with new enzymes [60 - 64];
- (2) Improvement of existing industrial organisms and techniques for using their enzymes [65, 66];
- (3) Improvement of operations related to substrate selection, cultivation conditions, enzyme processing and process reorganization [67 - 70] and
- (4) Creation of genetically modified plant cultures [71].

Previous studies have shown that high hemicellulase activity observed in *C. acetobutylicum-6*, *C. acetobutylicum-7* and *C. acetobutylicum* VKPM B-4786 strains made it possible to use plant biomass for the butanol production instead of corn or wheat flour as a traditional substrate [44]. Moreover, repeated “passages” of *C. thermocellum* 5 CT strain with high

cellulolytic activity enabled the authors [60] to obtain thermostable enzymes.

Several studies were conducted earlier [72 - 80] to determine the ability of new strains to digest different components of lignocellulosic biomass. *Clostridium* cultures IFBG C6H and IFBG C7P were selected from the IFBG collection. They were introduced into the synthetic medium with sugars that are part of the hemicelluloses. It was shown that the selected strains can metabolise mannose, glucose, cellulose, arabinose and xylose. The maximum accumulation of butanol (1.9 g/l) was obtained by using xylose as a carbon source in the cultivation of IFBG C6H strain and cellulose with IFBG C7P strain (1.3 g/l). Similar results were obtained by the authors in the process of fermentation on model mixtures consisting of yeast extract and peptone with the sugars: cellulose, arabinose, glucose, xylose, cellobiose. Earlier research was conducted [76 - 83] to compare the butanol production efficiency of the above mentioned new strains with that of the old ones. The results demonstrated that the new strains could digest a wide range of raw materials. Due to the ability of these strains to dissimilate cellulose, their cultivation continued with cellulosic raw materials added to the medium.

To confirm the cellulose fermentation ability of IFBG C6H and IFBG C7P strains, the latter were plated on a filter paper. After 72 h of cultivation, transparent areas appeared on the filter paper, and after 120 h, the destruction of the paper around the colonies was observed. These tests demonstrated the bioconversion of non-food raw materials (filter paper) by 90% and 60% for IFBG C6H and IFBG C7P strains, respectively. On cellulose rich substrate, both strains produced acetic and butyric acids at a concentration of approximately 4 g/l and produced almost no alcohol. These results allowed to suppose

the presence of cellulases in *C. acetobutylicum* IMB B-7407 (IFBG C6H - parent strain), *Clostridium sp.* IMB B-7570 (IFBG C6H 5M - mutant strain) and the possibility of using cellulose as a carbon source.

To study the ability of the strains of *Clostridium sp.* to assimilate lignocellulosic raw materials, a study was conducted by cultivating them on different lignocellulosic-rich biomasses/wastes (rape, soybean, wheat, switch grass, sawdust of various tree species, stock, fallen leaves). The dried green biomass was used without further pre-treatment. All strains assimilated the biomass and converted it to butanol Fig. (3). However, the accumulation of butanol was low (i.e., less than 1 g/l).

4. SUBSTRATE PREPARATION

Pre-treatment of lignocellulosic raw material is essential to facilitate its bioconversion. The pre-treatment is usually performed in several technological steps. After drying, the basic pre-treatment step for all types of lignocellulosic raw materials is grinding [84]. The procedures and extent of grinding determine the temperature regime of subsequent water-heat treatment and the extent of the fermented carbohydrates release. The higher the degree of dispersion of the raw materials, the lower the boiling point and sugar content, while the ABE solvents yield is increased. This is due to the partial cellulose degradation and the formation of additional fermentable sugars [2].

Effect of the variation of grinding degree of rape and switchgrass biomass on butanol accumulation by IMB B-7570 strain was investigated earlier [38 - 40]. Without plant biomass grinding, butanol accumulated at a concentration of up to 0.1 g/l. With prior grinding of rape biomass, butanol accumulation in the culture fluid increased, the highest level of butanol accumulation was observed at the grinding degree of 200 mesh (0.076 mm). We believe that the improvement in substrate bioavailability may be due to the fact that grinding decreases

the number of crystalline cellulose zones and increases the number of amorphous zones, which are easily fermentable by the bacterial enzymes. In *Clostridium* bacteria, the enzymes that breakdown cellulose are part of the extracellular multiprotein complex - cellulosome.

A recent study was conducted to test the accumulation of butanol by IMB B-7570, IFBG C4B, and IFBG C6H strains using ground rape biomass as a substrate [27]. Three major products of the ABE process - acetone, butanol and ethanol - were identified in the fluid of these cultures. The highest accumulation of butanol (2.3 g/l) was shown for *Clostridium sp.* IMB B-7570 strain; acetone was present in a small amount (0.5 g/l), and ethanol in the amount of 0.1 g/l. The cultivation of *Clostridium sp.* IMB B-7570 (IFBG C6H 5M), *C. acetobutylicum* IIB B-7407 (IFBG C6H) and *C. tyrobutylicum* IFBG C4B strains on sorghum biomass, juice and bagasse was also performed. It was shown that the least accumulation of butanol (0.5 g/l) was achieved by *C. tyrobutylicum* IFBG C4B on ground sorghum biomass, whereas the highest accumulation (8.2 g/l) by *C. acetobutylicum* IMB B-7407 (IFBG C6H) on sorghum juice.

One of the possible subsequent lignocellulosic biomass pre-treatment steps is the temperature treatment to release sugars. This treatment can be carried out either at a relatively low temperature (135-145°C) for one or more hours or at a higher temperature (165-175°C) for 2 to 5 min, or in a stage wise manner - from higher temperature to lower [84]. The raw material preparatory treatments can be conducted in a variety of ways, for example with organosol [85, 86], alkali [87], sodium hydroxide, acid [88, 89], ionic liquid [90 - 92], high-frequency heating [93], sharp steam [94, 95], lime [96] or ammonia [97]. The most promising of them is Organosol preparation and treatment with the help of acute steam. Such processing increases the sugar yield, reduces the amount of insoluble lignin and increases the content of easily fermentable cellulose [85, 86].

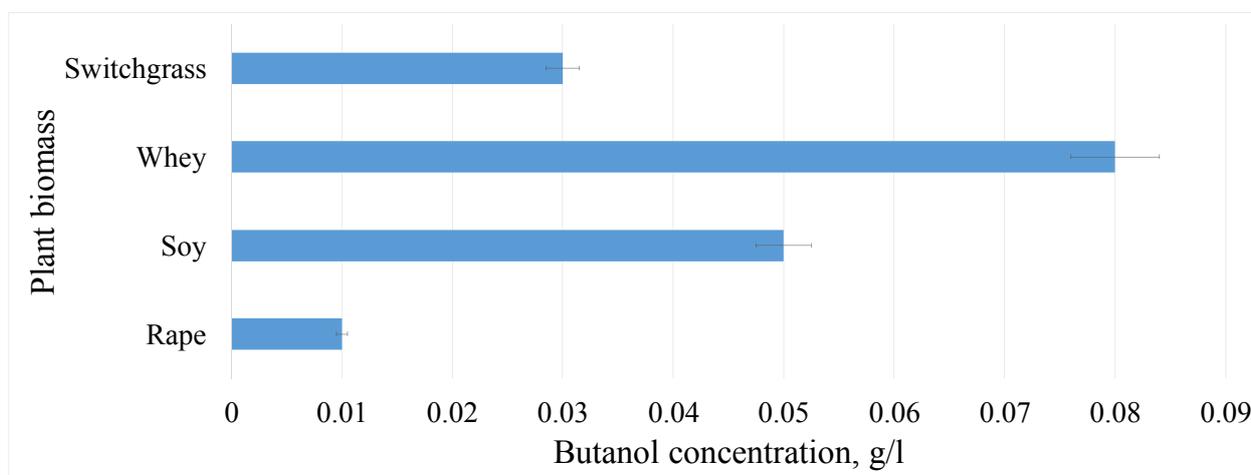


Fig. (3). Butanol accumulation on native plant biomass by *Clostridium sp.*

Another possible step of raw material preparation is through the hydrolysis of sugars to pentoses or hexoses. Such hydrolysis can be performed with the use of alkalis, acids or enzymes. Enzymatic hydrolysis involves the use of both enzymes and enzyme-producing organisms. The catalytic activity of enzymes is characterized by “turnover number”, that is, the number of moles of substance converted per time unit per enzyme mole. The activity of enzymes depends on temperature, concentration of water ions, presence of activators and inhibitors. The detailed mechanism of lignocellulose enzymatic hydrolysis has not yet been elucidated.

Significant progress has been made in the study of lignocellulolytic genes of organisms known to show hydrolytic properties (including bacteria and fungi) [98, 99]. Currently, more than 14,000 mushroom species are known to exhibit cellulolytic activity, and only some of them are used in industrial technologies [100]. *Trichoderma reesei* is one of the most common organisms for cellulose and hemicellulose degradation, although it does not have enzymes for lignin degradation [98 - 100]. The major role in lignin processing is played by basidiomycetes, living on both live and deadwood. Much of protolignin is degraded by brown rot fungi, which cause the hydrolysis of polysaccharides [11]. The effective destructors of lignin are white rot fungi, among which *Phanerochaete chrysosporium* is studied the best [101, 102]. These are species used in industrial technologies for the production of lignin degrading enzymes, as well as for direct use during the lignocellulose bioconversion [103, 104].

Polymeric lignin degradation occurs under the action of extracellular enzymes - oxidoreductases, in particular ligninperoxidase, Mn-peroxidase and laccase. In addition, the ligninolytic fungi complex includes pyranosoxidases, glucose oxidases, glyoxal oxidase and cellobiodehydrogenase. *Daedalea flavida*, *Phlebia fascicularia*, *P. floridensis* and *P. radiata* fungi have not been studied deeply enough and they are used only for the lignin separation from raw materials and for the selective lignin degradation [105]. *Cellulomonas*, *Pseudomonas*, *Thermomonospora* and *Microbispora* actinomycetes, as well as bacteria with a surface-bound cellulose complex, such as *Clostridium thermocellum* and *Ruminococcus*, have a slight lignin digestion potential [106]. Organisms with high lignin degradation activity could be used as a source of the gene pool for engineering those organisms with slight lignin digestion potential. Currently, the cost of enzymes is one of the major obstacles to exploit the hydrolysis of cellulose by an enzymatic approach. Due to the low cellulase activity of enzymes, it is necessary to increase the bioavailability of cellulose (destroy its crystalline regions) by mechanical treatment [107] or hydrolysis.

Fractionation of the lignocellulosic switchgrass biomass onto its components following the thermobaric treatment produces water-soluble substances (Table 1) and stock (Table 2). The feasibility of using switchgrass lignocellulose components as substrates for butanol synthesis was tested by cultivation of the IFBG C6H and IFBG C6H 5M strains on various switchgrass components and sludge medium [31].

It was found that the products initially obtained with the thermobaric hydrolysis of switchgrass need to be purified from

furfural, formed during the thermobaric treatment and that acts as an inhibitor of the microorganisms. It was also shown that switchgrass sludge contains a higher proportion of cellulose in comparison to the switchgrass biomass [31]. From the results, it is clear that the obtained components (cellulose, arabinogalactan, lignin) can be used to produce butanol. It should be noted that, due to the sludge thermobaric hydrolysis, furfural was formed in a small amount, which did not affect the viability of the microorganisms and did not require additional separation. In this case, the use of thermobaric hydrolysis increased the accumulation of target product twofold [31].

Table 1. Millet switchgrass biomass composition in resulting of autohydrolysis.

Components	original Biomass, %	Biomass after autohydrolysis, %
Cellulose	46,7±0,5	53,9±0,5
Hemicellulose	23,0±1,0	10,6±1,0
Soluble substances	7,7±0,5	11,5±0,5
Lignin	13,8±0,2	14,7±0,2
Resins and fats	2,0±10	2,0±10
Ash	5,4±0,5	5,4±0,5
Others	1,4±0,2	1,9±0,2
Total	100,0	100,0

Table 2. Sludge composition in resulting of autohydrolysis.

Components	Original composition, %	Composition after autohydrolysis, %
Cellulose	34,0±0,5	37,0±0,5
Hemicellulose	3,7±1,0	1,5±1,0
Soluble substances	3,2±0,5	6,9±0,5
Lignin	8,5±0,2	3,5±0,2
Resins and fats	2,3±10	2,3±10
Ash	47,8±0,5	48,0±0,5
Others	0,5±0,2	0,8±0,2
Total	100,0	100,0

5. OPTIMIZATION OF CULTIVATION PARAMETERS

An important aspect of the ultimate cost of butanol production is also the cost of the substrate and its concentration in the fermentation medium. The efficiency of butanol accumulation using different concentrations of switchgrass and rape biomasses in the fermentation medium was investigated earlier [27, 38]. The data showed that the accumulation of butanol in the culture fluid increased with the increase of the rape biomass concentration from 5 to 10 g/l. Further increase in the amount of dry ground rape biomass from 15.0 to 30.0 g/l decreased the accumulation of butanol. The results of study showed that the bioavailability of the substrate decreases with the increase in carbon concentration. The highest accumulation of butanol (2.9 g/l) in the culture fluid was obtained with a dry ground rape biomass concentration of 10.0 g/l. and the raw materials pre-treated by autoclaving for 2 hours at a pressure of 2 atm. Further increase in the processing time or pressure did not significantly affect the production of butanol.

However, carbon is not the only essential parameter

affecting the growth of microorganisms. It is known that optimization of cultivation parameters such as temperature, pH and incubation time can increase the accumulation of butanol in cultural fluid several times [108, 109]. The influence of cultivation parameters on butanol accumulation was investigated, using switchgrass as a substrate. The effect of temperature of cultivation on butanol accumulation was investigated on IFBG C6H and IFBG C7P strains. The basic cultivation parameters of these strains were optimized, as described earlier [38]. It was shown that the temperature increase from $28\pm 1^\circ\text{C}$ to $36\pm 1^\circ\text{C}$ led to an increase in the butanol accumulation for both strains, temperature increase to $38\pm 1^\circ\text{C}$ reduced the butanol yield in the culture fluid. All further studies were performed at $35\pm 1^\circ\text{C}$.

It is well known that the pH of the medium significantly affects butanol accumulation. In our experiments, the effect of pH on butanol production was investigated using the IFBG C6H and IFBG C7P strains. It was found that the maximum butanol accumulation (3.5 g/l) occurred at pH 7.0 with IFBG C6H strain and 2.5 g/l at pH 6.0 with IFBG C7P strain. Moreover, pH adjustment from 5 to 7 did not affect butanol accumulation by the IFBG C7P strain. At the beginning of the fermentation, the bacterial count was determined by the quality and quantity of the bacterial inoculum. The number of cellulosomes increased following the increase of bacterial count in the fermentation medium. For this reason, to standardize the amount of bacteria introduced into the inoculum, the liquid nutrient media with water-soluble carbon sources were used.

The effect of the seed material concentration on butanol accumulation was investigated using lignocellulosic biomass as a substrate and the IMB B-7570 strain. It was established that with the increase of the seed material concentration from 5 to 10% of the fermentation mixture volume, the butanol synthesis increased [2]. The increase of inoculum concentration up to 15-20% caused a negative effect on butanol accumulation. Further increase in seed material concentration added to the fermentation mixture generally suppressed the butanol synthesis in the medium. The inhibition of butanol synthesis by the high inoculum concentration may be due to an increase of the primary metabolites of the ABE fermentation (butyric, lactic and acetic acids) in the accumulation medium. The optimum concentration of the inoculum introduced into the fermentation medium was 10%. Under such conditions, the largest amount of butanol accumulated was 2.5 g/l.

Since ABE is a biphasic process, the time of cultivation is very important for optimisation of butanol accumulation conditions. The impact of changing cultivation times on butanol accumulation in the fluid was investigated using IFBG C6H and IFBG C7P strains. The maximum butanol accumulation was obtained after 72 h of cultivation for both IFBG C6H and IFBG C7P strains on a synthetic medium. Further elongation of the cultivation time did not produce positive effects on butanol synthesis. Optimization of the technological parameters by testing different types of plant material allowed to increase the butanol accumulation by 20% (2 g/l) with strain IFBG C6H and by 50% (4.3 g/l) with the mutant strain IFBG C6H 5M [40].

6. USE OF SYNTHESIS PRECURSORS

ABE fermentation is known to be a type of butyric acid fermentation; therefore, the addition of precursors of this process, such as lactic, acetic and butyric acids, to increase the accumulation of butanol, has been investigated.

The corresponding acids were introduced into the enzymatic medium, first individually, and then in the mixture, and for each case, the cultivation was performed [40]. The obtained data indicate that the use of synthesis precursors increased butanol accumulation. Slight stimulation of butanol accumulation, as compared to control (3.5 g/l), was observed in the case of acetic acid (3.7 g/l) and serum (3.9 g/l). A significant 1.7-fold increase in butanol concentration was due to the use of a mixture of all three precursors, with the conversion of sugars almost doubling from 32% to 58%. The addition of precursors to the fermentation medium increased the economic coefficients of fermentation, but did not affect biomass accumulation and specific growth rate of culture. It should be noted that the ratio of butanol concentration to biomass increased from 1.2 to 2 with the use of precursors. At the same time, the biomass to consumed substrate ratio decreased from 0.38 to 0.29, respectively. Such a change in kinetic parameters as a result of precursors addition indicates an increase in butanol accumulation due to substrate conversion. To determine the optimal concentration of precursors, the cultivation with different combinations of concentrations of precursors was performed. The butanol synthesis precursors introduced into the fermentation medium increased the final concentration of end-product differently. The optimal option was a mixture of lactic, acetic and butyric acids in a ratio of 1: 1: 3, respectively. The addition of a mixture of these acids increased the concentration of the target product by 17% [40].

7. WASTE DISPOSAL

An important issue related to butanol production technology is the production of waste disposal. The utilization (evaporation) of the acetone-butyl bard is quite costly. To overcome this challenge, we considered an option to return the acetone-butyl distillery waste to the cultivation process [40]. The use of "reverse" distillery waste to reduce production waste was successful. The concentration of the reverse distillery wastes up to 60% of the volume of the fermentation mixture did not significantly affect the synthesis of butanol by *C. acetobutylicum* IFBG C6H 5M strain. Further increase in the concentration of distillery waste in medium decreased accumulation of butanol. This finding makes it possible to return up to 60% of the distillery waste to the enzymatic medium. The remaining part of the distillery waste was separated into decantate and sludge. The decantate contained 80-85% of the distillery waste, 2% of solids, 0.7 g/l of reducing substances and had a pH 4. The decantate was used instead of water in further fermentation ("reverse" water), and the sludge can be used as animal feed or as a fertilizer [40].

CONCLUSION

Overall, our studies demonstrate the possibility to produce butanol efficiently from plant biomass. The accumulation of

solvents (alcohols) directly depends on the characteristics of the butanol-producing bacterial strains and the stages of biomass pre-treatment (preparation) - drying, grinding, sterilization, type of hydrolysis. In order to develop commercial biobutanol technology using lignocellulosic raw materials, it is, however, necessary to improve significantly the efficiency of the bioconversion of lignocellulosic raw materials by selecting high yield bacterial strains and optimizing the efficiency of pre-treatment methods of the biomass.

CONSENT FOR PUBLICATION

Not applicable.

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None.

CONFLICT OF INTEREST

The author declares no conflict of interest, financial or otherwise.

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REFERENCES

- Cheng C, Bao T, Yang ST. Engineering *Clostridium* for improved solvent production: recent progress and perspective. *Appl Microbiol Biotechnol* 2019; 103(14): 5549-66. [http://dx.doi.org/10.1007/s00253-019-09916-7] [PMID: 31139901]
- Tigunova OO, Beiko NY, Andriash HS, Pryiomov SH, Shulga SM. Domestic butanol-producing strains of the *Clostridium* genus. *Biotechnol Acta* 2017; 10(1): 34-42. [http://dx.doi.org/10.15407/biotech10.01.034]
- Jiménez-Bonilla P, Wang Y. *In situ* biobutanol recovery from clostridial fermentations: a critical review. *Crit Rev Biotechnol* 2017; 14(1): 4-6. [http://dx.doi.org/10.1080/07388551.2017.1376308] [PMID: 28920460]
- Tigunova O, Shulga S, Blume Y. Biobutanol as an alternative type of fuel. *Cytol Genet* 2013; 47(6): 51-71. [http://dx.doi.org/10.3103/S0095452713060042] [PMID: 24437198]
- Lin Z, Liu H, Wu J, Patakova P, Branska B, Zhang J. Effective continuous acetone-butanol-ethanol production with full utilization of cassava by immobilized symbiotic TSH06. *Biotechnol Biofuels* 2019; 12: 219. [http://dx.doi.org/10.1186/s13068-019-1561-1] [PMID: 31534478]
- Kolesinska B, Fraczyk J, Binczarski M, et al. Butanol synthesis routes for biofuel production: trends and perspectives. *Materials* 2019; 12(350): 1-2. [http://dx.doi.org/10.3390/ma12030350]
- Tian L, Conway PM, Cervenska ND, et al. Metabolic engineering of *Clostridium thermocellum* for *n*-butanol production from cellulose. *Biotechnol Biofuels* 2019; 12(186): 186. [http://dx.doi.org/10.1186/s13068-019-1524-6] [PMID: 31367231]
- Benali M, Ajao O, Mehdi NE, Restrepo AM, Fradj N, Boumghar Y. Acetone-butanol-ethanol production from Eastern Canadian yellow birch and screening of isopropanol-butanol-ethanol-producing strains. *Ind Biotechnol (New Rochelle NY)* 2019; 15(3): 188-201. [http://dx.doi.org/10.1089/ind.2019.0002]
- Niglio S, Marzocchella A, Rehmann L. Clostridial conversion of corn syrup to Acetone-Butanol-Ethanol (ABE) via batch and fed-batch fermentation. *Heliyon* 2019; 5(3):e01401 [http://dx.doi.org/10.1016/j.heliyon.2019.e01401] [PMID: 30963127]
- Al-Shorgani NKN, Al-Tabib AI, Kadier A, Zamil MF, Lee KM, Kalil MS. Continuous butanol fermentation of dilute acid-pretreated deoiled rice bran by *Clostridium acetobutylicum* YM1. *Sci Rep* 2019; 9(1): 4622. [http://dx.doi.org/10.1038/s41598-019-40840-y] [PMID: 30874578]
- Shulga SM, Tigunova OO, Blume YB. Lignocellulose as an alternative source for obtaining of biobutanol. *Biotechnol Acta* 2013; 6(2): 9-21. [http://dx.doi.org/10.15407/biotech6.02.009]
- Jing Q, Wang H, Huang H, Qureshi Q. Acetone-butanol-ethanol (ABE) fermentation of soluble and hydrolyzed sugars in apple pomace by *Clostridium beijerinckii* P26. *Fuel* 2019; 244: 536-44. [http://dx.doi.org/10.1016/j.fuel.2019.01.177]
- Birgen C, Dürre P, Preisig HA, Wentzel A. Butanol production from lignocellulosic biomass: revisiting fermentation performance indicators with exploratory data analysis. *Biotechnol Biofuels* 2019; 12: 167. [http://dx.doi.org/10.1186/s13068-019-1508-6] [PMID: 31297155]
- Raganati F, Procentese A, Olivieri G, Gotz P, Salatino P, Marzocchella A. Kinetic study of butanol production from various sugars by *Clostridium acetobutylicum* using a dynamic model. *Biochem Eng J* 2015; 99(15): 156-66. [http://dx.doi.org/10.1016/j.bej.2015.03.001]
- Jones DT, Woods DR. Acetone-butanol fermentation revisited. *Microbiol Rev* 1986; 50(4): 484-524. [http://dx.doi.org/10.1128/MMBR.50.4.484-524.1986] [PMID: 3540574]
- Howard RL, Abotsi E, Jansen van Rensburg EL, Howard S. Lignocellulose biotechnology issues of bioconversion and enzyme production. *Afr J Biotechnol* 2003; 2(12): 602-19. [http://dx.doi.org/10.5897/AJB2003.000-1115]
- Malherbe S, Cloete TE. Lignocellulose biodegradation: fundamentals and applications. *Rev Environ Sci Biotechnol* 2003; 1: 105-14. [http://dx.doi.org/10.1023/A:1020858910646]
- Yadav RC, Devi NL. Biomass burning, regional air quality, and climate change. *Earth Systems and Environmental Sciences, Encyclopedia of Environmental Health*. 2nd ed. Waltham, MA: Elsevier 2018.
- Tran TTA, Le TKP, Mai TP, Nguyen DQ. Bioethanol production from lignocellulosic biomass. *IntechOpen* 2019.
- Bhat MK. Cellulases and related enzymes in biotechnology. *Biotechnol Adv* 2000; 18(5): 355-83. [http://dx.doi.org/10.1016/S0734-9750(00)00041-0] [PMID: 14538100]
- Sun Y, Cheng J. Hydrolysis of lignocellulosic materials for ethanol production: a review. *Bioresour Technol* 2002; 83(1): 1-11. [http://dx.doi.org/10.1016/S0960-8524(01)00212-7] [PMID: 12058826]
- Li X, Chang S, Liu R. Industrial Applications of cellulases and hemicellulases: microbial production and application. In: *Fungal Cellulolytic Enzymes*. 2018; pp. 267-82. [http://dx.doi.org/10.1007/978-981-13-0749-2_15]
- Beauchemin KA, Morgavi DP, McAllister TA. The use of enzymes in ruminant diets. *Animal Nutr Nottingham Univ Press* 2001; pp. 296-322.
- Beauchemin KA, Colombatto D, Morgavi DP, Yang WZ. Use of exogenous fibrolytic enzymes to improve animal feed utilisation by ruminants. *J Anim Sci* 2003; 81(2): 37-47. [http://dx.doi.org/10.2527/2003.8114_suppl_2E37x]
- Subramaniyan S, Prema P. Biotechnology of microbial xylanases: enzymology, molecular biology, and application. *Crit Rev Biotechnol* 2002; 22(1): 33-64. [http://dx.doi.org/10.1080/07388550290789450] [PMID: 11958335]
- Beg QK, Kapoor M, Mahajan L, Hoondal GS. Microbial xylanases and their industrial applications: a review. *Appl Microbiol Biotechnol* 2001; 56(3-4): 326-38. [http://dx.doi.org/10.1007/s002530100704] [PMID: 11548999]
- Tigunova OO, Andriash HS, Beiko NY, Zaharova OG, Pryiomov SH, Shulga SM. Rape biomass (*Brassica napus*) as a raw materials for biobutanol production. *Biotechnol Acta* 2019; 12(1): 75-81. [http://dx.doi.org/10.15407/biotech12.01.075]
- Rajagopalan G, He J, Yang K-I. One pot fermentation of agricultural residues to produce butanol and hydrogen by *Clostridium* strain BOH3. *Renew Energy* 2016; 85: 1127-34.

- [29] Borah AJ, Roy K, Goyal A, Moholkar VS. Mechanistic investigations in biobutanol synthesis via ultrasound-assisted ABE fermentation using mixed feedstock of invasive weeds. *Bioresour Technol* 2019; 272: 389-97. [http://dx.doi.org/10.1016/j.renene.2015.07.051]
- [30] Lin PP, Mi L, Morioka AH, *et al.* Consolidated bioprocessing of cellulose to isobutanol using *Clostridium thermocellum*. *Metab Eng* 2015; 31: 44-52. [http://dx.doi.org/10.1016/j.biortech.2018.10.063] [PMID: 30388576]
- [31] Tigonova OO, Beiko NE, Kamenskyh DS, *et al.* Lignocellulosic biomass after explosive autohydrolysis as substrate for butanol obtaining. *Biotechnol Acta* 2016; 9(4): 28-34. [http://dx.doi.org/10.15407/biotech9.04.028]
- [32] Dhyani V, Bhaskar T. *Biofuels: Alternative Feedstocks and Conversion Processes for the Production of Liquid and Gaseous Biofuels (Second Edition) Biomass, Biofuels, Biochemicals*. 2019; pp. 217-44.
- [33] Lampugnani ER, Khan GA, Somssich M, Persson S. Building a plant cell wall at a glance. *J Cell Sci* 2018; 131(2): jcs207373 [http://dx.doi.org/10.1242/jcs.207373] [PMID: 29378834]
- [34] Desvaux M. *Clostridium cellulolyticum*: model organism of mesophilic cellulolytic clostridia. *FEMS Microbiol Rev* 2005; 29(4): 741-64. [http://dx.doi.org/10.1016/j.femsre.2004.11.003] [PMID: 16102601]
- [35] Lopez Contreras AM. Utilization of lignocellulosic substrates by solvent-producing clostridia. PhD thesis Wageningen University, Wageningen The Netherlands. 2003; 144.
- [36] Huang J, Fu S, Gan L. *Lignin chemistry and application*. Elsevier 2019; p. 255.
- [37] Bogolitsin KG, Lunina VV. Physical chemistry of lignin. *Moskva, Akademkniga* 2010; p. 489.
- [38] Tigonova OO, Shulga SM. New producer strains of biobutanol. II. Renewable lignocellulose fermentation. *Biotechnol Acta* 2014; 7(4): 54-60. [http://dx.doi.org/10.15407/biotech7.04.054]
- [39] Tigonova OO, Shulga SM. Using by mutant strains *C. acetobutylicum* ligno-cellulosic material as a substrate. *Microb Biotechnol* 2015; 3: 35-44. [http://dx.doi.org/10.18524/2307-4663.2015.3(31).53656]
- [40] Tigonova OO, Shulga SM. New producer strains of biobutanol. III. Methods increased accumulation of butanol from switchgrass *Panicum virgatum* L. biomass. *Biotechnol Acta* 2015; 8(4): 65-8. [http://dx.doi.org/10.15407/biotech8.04.092]
- [41] Xue C, Cheng C. Butanol production by *Clostridium*. *Adv Bioenergy* 2019; 4: 35-77. [http://dx.doi.org/10.1016/bs.aibe.2018.12.001]
- [42] Salleh MSM, Ibrahim MF, Roslan AM, Abd-Aziz S. Improved biobutanol production in 2-l simultaneous saccharification and fermentation with delayed yeast extract feeding and *in-situ* recovery. *Sci Rep* 2019; 9(1): 7443. [http://dx.doi.org/10.1038/s41598-019-43718-1] [PMID: 31092836]
- [43] Passoth V, Sandgren M. Biofuel production from straw hydrolysates: current achievements and perspectives. *Appl Microbiol Biotechnol* 2019; 103(13): 5105-16. [http://dx.doi.org/10.1007/s00253-019-09863-3] [PMID: 31081521]
- [44] Zverlov VV, Berezina O, Velikodvorskaya GA, Schwarz WH. Bacterial acetone and butanol production by industrial fermentation in the Soviet Union: use of hydrolyzed agricultural waste for biorefinery. *Appl Microbiol Biotechnol* 2006; 71(5): 587-97. [http://dx.doi.org/10.1007/s00253-006-0445-z] [PMID: 16685494]
- [45] Berezina OV, Sineoky SP, Velikodvorskaya GA, Schwarz W, Zverlov VV. Extracellular glycosyl hydrolase activity of the *Clostridium* strains producing acetone, butanol, and ethanol. *Appl Biochem Microbiol* 2008; 44(1): 42-7. [http://dx.doi.org/10.1134/S0003683808010079]
- [46] Zhao T, Tashiro Y, Sonomoto K. Smart fermentation engineering for butanol production: designed biomass and consolidated bioprocessing systems. *Appl Microbiol Biotechnol* 2019; 103(23-24): 9359-71. [http://dx.doi.org/10.1007/s00253-019-10198-2] [PMID: 31720773]
- [47] Hallac BB, Ragauskas AJ. Analyzing cellulose degree of polymerization and its relevancy to cellulosic ethanol Review. *Biofuels Bioprod Biorefin* 2011; 5: 215-25. [http://dx.doi.org/10.1002/bbb.269]
- [48] Herrera LM, Braña V, Franco Fraguas L, Castro-Sowinski S. Characterization of the cellulase-secretome produced by the Antarctic bacterium *Flavobacterium* sp. AUG42. *Microbiol Res* 2019; 223-225: 13-21. [http://dx.doi.org/10.1016/j.micres.2019.03.009] [PMID: 31178046]
- [49] Qureshi N, Liu S, Ezeji NC. Cellulosic butanol production from agricultural biomass and residues: recent advances in technology. *Advanced Biofuels and Bioproducts*. Springer 2012; pp. 247-65.
- [50] Doi RH, Tamaru Y. The *Clostridium cellulovorans* cellulosome: an enzyme complex with plant cell wall degrading activity. *Chem Rec* 2001; 1(1): 24-32. [http://dx.doi.org/10.1002/1528-0691(2001)1:1<24::AID-TCR5>3.0.CO;2-W] [PMID: 11893054]
- [51] Schwarz WH. The cellulosome and cellulose degradation by anaerobic bacteria. *Appl Microbiol Biotechnol* 2001; 56(5-6): 634-49. [http://dx.doi.org/10.1007/s002530100710] [PMID: 11601609]
- [52] Gal L, Pages S, Gaudin C, *et al.* Characterization of the cellulolytic complex (cellulosome) produced by *Clostridium cellulolyticum*. *Appl Environ Microbiol* 1997; 63(3): 903-9. [http://dx.doi.org/10.1128/AEM.63.3.903-909.1997] [PMID: 9055408]
- [53] Bélaich JP, Tardif C, Bélaich A, Gaudin C. The cellulolytic system of *Clostridium cellulolyticum*. *J Biotechnol* 1997; 57(1-3): 3-14. [http://dx.doi.org/10.1016/S0168-1656(97)00085-0] [PMID: 9335163]
- [54] Kakiuchi M, Isui A, Suzuki K, *et al.* Cloning and DNA sequencing of the genes encoding *Clostridium josui* scaffolding protein CipA and cellulase CelD and identification of their gene products as major components of the cellulosome. *J Bacteriol* 1998; 180(16): 4303-8. [http://dx.doi.org/10.1128/JB.180.16.4303-4308.1998] [PMID: 9696784]
- [55] Tamaru Y, Karita S, Ibrahim A, Chan H, Doi RH. A large gene cluster for the *Clostridium cellulovorans* cellulosome. *J Bacteriol* 2000; 182(20): 5906-10. [http://dx.doi.org/10.1128/JB.182.20.5906-5910.2000] [PMID: 11004194]
- [56] Qureshi N, Ezji TC. Butanol, 'a superior biofuel' production from agricultural residues (renewable biomass): recent progress in technology. *Biofuels Bioprod Biorefin* 2008; 2(4): 319-30. [http://dx.doi.org/10.1002/bbb.85]
- [57] López-Contreras AM, Martens AA, Szijarto N, *et al.* Production by *Clostridium acetobutylicum* ATCC 824 of CelG, a cellulosomal glycoside hydrolase belonging to family 9. *Appl Environ Microbiol* 2003; 69(2): 869-77. [http://dx.doi.org/10.1128/AEM.69.2.869-877.2003] [PMID: 12571006]
- [58] Nölling J, Breton G, Omelchenko MV, *et al.* Genome sequence and comparative analysis of the solvent-producing bacterium *Clostridium acetobutylicum*. *J Bacteriol* 2001; 183(16): 4823-38. [http://dx.doi.org/10.1128/JB.183.16.4823-4838.2001] [PMID: 11466286]
- [59] Walker LP, Wilson DB. Enzymatic hydrolysis of cellulose: An overview. *Bioresour Technol* 1991; 36: 3-14. [http://dx.doi.org/10.1016/0960-8524(91)90095-2]
- [60] Yastremskaya LS, Vasilchenko OA. Selection of anaerobic cellulolytic thermophilic strain *Clostridium thermocellum* 5CT. *Biotechnologia* 2011; 4(2): 80-5. [in Ukr.].
- [61] Hongoh Y, Sharma VK, Prakash T, *et al.* Genome of an endosymbiont coupling N₂ fixation to cellulolysis within protist cells in termite gut. *Science* 2008; 322(5904): 1108-9. [http://dx.doi.org/10.1126/science.1165578] [PMID: 19008447]
- [62] Jindou S, Xu Q, Kenig R, *et al.* Novel architecture of family-9 glycoside hydrolases identified in cellulosomal enzymes of *Acetivibrio cellulolyticus* and *Clostridium thermocellum*. *FEMS Microbiol Lett* 2006; 254(2): 308-16. [http://dx.doi.org/10.1111/j.1574-6968.2005.00040.x] [PMID: 16445761]
- [63] Willquist K, Zeidan AA, van Niel EWJ. Physiological characteristics of the extreme thermophile *Caldicellulosiruptor saccharolyticus*: an efficient hydrogen cell factory. *Microb Cell Fact* 2010; 9(89): 89. [http://dx.doi.org/10.1186/1475-2859-9-89] [PMID: 21092203]
- [64] Mullin DA, Velankar HR. Isolated bacteria, methods for use, and methods for isolation. WIPO Patent WO2012021678A2 2012.
- [65] Brás JL, Cartmell A, Carvalho AL, *et al.* Structural insights into a unique cellulase fold and mechanism of cellulose hydrolysis. *Proc Natl Acad Sci USA* 2011; 108(13): 5237-42. [http://dx.doi.org/10.1073/pnas.1015006108] [PMID: 21393568]
- [66] Jennert KCB, Tardif C, Young DI, Young M. Gene transfer to *Clostridium cellulolyticum* ATCC 35319. *Microbiology* 2000; 146(Pt 12): 3071-80. [http://dx.doi.org/10.1099/00221287-146-12-3071] [PMID: 11101665]
- [67] Lu Y, Zhang Y-HP, Lynd LR. Enzyme-microbe synergy during

- cellulose hydrolysis by *Clostridium thermocellum*. Proc Natl Acad Sci USA 2006; 103(44): 16165-9.
[http://dx.doi.org/10.1073/pnas.0605381103] [PMID: 17060624]
- [68] Guedon E, Payot S, Desvaux M, Petitdemange H. Carbon and electron flow in *Clostridium cellulolyticum* grown in chemostat culture on synthetic medium. J Bacteriol 1999; 181(10): 3262-9.
[http://dx.doi.org/10.1128/JB.181.10.3262-3269.1999] [PMID: 10322031]
- [69] Desvaux M, Guedon E, Petitdemange H. Cellulose catabolism by *Clostridium cellulolyticum* growing in batch culture on defined medium. Appl Environ Microbiol 2000; 66(6): 2461-70.
[http://dx.doi.org/10.1128/AEM.66.6.2461-2470.2000] [PMID: 10831425]
- [70] Hibino S, Minami Z. Bacterial preparation for agricultural use. US Patent N 5,733,355 1998.
- [71] Ohmiya K, Sakka K, Kimura T, Morimoto K. Application of microbial genes to recalcitrant biomass utilization and environmental conservation. J Biosci Bioeng 2003; 95(6): 549-61.
[http://dx.doi.org/10.1016/S1389-1723(03)80161-5] [PMID: 16233456]
- [72] Biebl H. Fermentation of glycerol by *Clostridium pasteurianum*--batch and continuous culture studies. J Ind Microbiol Biotechnol 2001; 27(1): 18-26.
[http://dx.doi.org/10.1038/sj.jim.7000155] [PMID: 11598806]
- [73] Xiao H, Gu Y, Ning Y, et al. Confirmation and elimination of xylose metabolism bottlenecks in glucose phosphoenolpyruvate-dependent phosphotransferase system-deficient *Clostridium acetobutylicum* for simultaneous utilization of glucose, xylose, and arabinose. Appl Environ Microbiol 2011; 77(22): 7886-95.
[http://dx.doi.org/10.1128/AEM.00644-11] [PMID: 21926197]
- [74] Sivagnanam K, Raghavan VGS, Shah M. L. Hetlich R, Verberkmoes NC, Lefsrud MG. Characterization of *Clostridium acetobutylicum* protein interaction network from butanol fermentation. J Anal Bioanal Tech 2012; S3
[http://dx.doi.org/10.4172/2155-9872.S3-002]
- [75] López-Contreras AM, Gabor K, Martens AA, et al. Substrate-induced production and secretion of cellulases by *Clostridium acetobutylicum*. Appl Environ Microbiol 2004; 70(9): 5238-43.
[http://dx.doi.org/10.1128/AEM.70.9.5238-5243.2004] [PMID: 15345405]
- [76] Andrade JC, Vasconcelos I. Continuous cultures of *Clostridium acetobutylicum*: culture stability and low-grade glycerol utilisation. Biotechnol Lett 2003; 25(2): 121-5.
[http://dx.doi.org/10.1023/A:1021911217270] [PMID: 12882286]
- [77] López-Contreras AM, Smidt H, van der Oost J, Claassen PA, Mooibroek H, de Vos WM. *Clostridium beijerinckii* cells expressing *Neocallimastix patriciarum* glycoside hydrolases show enhanced lichenan utilization and solvent production. Appl Environ Microbiol 2001; 67(11): 5127-33.
[http://dx.doi.org/10.1128/AEM.67.11.5127-5133.2001] [PMID: 11679336]
- [78] Xiao H, Li Z, Jiang Y, et al. Metabolic engineering of D-xylose pathway in *Clostridium beijerinckii* to optimize solvent production from xylose mother liquid. Metab Eng 2012; 14(5): 569-78.
[http://dx.doi.org/10.1016/j.ymben.2012.05.003] [PMID: 22677452]
- [79] Watsom JE. Pentose sugar utilisation in *Clostridium beijerinckii* NCIMB 8052 for biobutanol production: genetic and physiological studies PhD Thesis. Edinburgh Napier University 2012. <https://www.napier.ac.uk/~media/worktribe/output-190857/jwatsonphdthesis2012.pdf>
- [80] Linggang S, Phang LY, Wasoh H, Abd-Aziz S. Acetone-butanol-ethanol production by *Clostridium acetobutylicum* ATCC 824 using sago pith residues hydrolysate. BioEnergy Res 2013; 6(1): 321-8.
[http://dx.doi.org/10.1007/s12155-012-9260-9]
- [81] Priyomov S, Tigunova O. Phylogenetic analysis of butanol producing strains compared 16S rRNA gene sequence. Naukovi dopovidi NUBiP Ukrainy 2015. Available from: http://nd.nubip.edu.ua/2015_8/index.html
- [82] Tigunova OO, Shulga SM. Using by mutant strain *C. acetobutylicum* ligno-cellulosic material as substrate. Mikrobiologija i biotehnologija 2015; : 35-44.
[http://dx.doi.org/10.18524/2307-4663.2015.3(31).53656(in Ukr.)]
- [83] Tigunova OO, Shulga SM. Butanol yield increase of *C. acetobutylicum* mutant strain on non-food raw material Proc Int Sci Conference «Biological Resources and Novel Biotechnologies of Biofuel Production» Kyiv. 142-48.
- [84] Olijnichuk ST, Levandovskij LV, Shevchenko VI. Technological regulations of ethanol production from starch-containing raw materials. Kyiv, TOV «Matritsia» 2000; p. 142.
- [85] Cateto C, Hu G, Ragauskas A. Enzymatic hydrolysis of organosolv Kanlow switchgrass and its impact on cellulose crystallinity and degree of polymerization. Energy Environ Sci 2011; 4: 1516-21.
[http://dx.doi.org/10.1039/c0ee00827c]
- [86] Sannigrahi P, Miller SJ, Ragauskas AJ. Effects of organosolv pretreatment and enzymatic hydrolysis on cellulose structure and crystallinity in Loblolly pine. Carbohydr Res 2010; 345(7): 965-70.
[http://dx.doi.org/10.1016/j.carres.2010.02.010] [PMID: 20307873]
- [87] Wang Z, Keshwani DR, Redding AP, Cheng JJ. Alkaline pretreatment of coastal bermudagrass for bioethanol. ASABE Annual Int Meeting; Rhode Island Convention Center Providence, Rhode Island.
- [88] Sun Y, Cheng JJ. Dilute acid pretreatment of rye straw and bermudagrass for ethanol production. Bioresour Technol 2005; 96(14): 1599-606.
[http://dx.doi.org/10.1016/j.biortech.2004.12.022] [PMID: 15978993]
- [89] Laser M, Schulman D, Allen SG, Lichwa J, Antal MJ Jr, Lynd LR. A comparison of liquid hot water and steam pretreatments of sugar cane bagasse for bioconversion to ethanol. Bioresour Technol 2002; 81(1): 33-44.
[http://dx.doi.org/10.1016/S0960-8524(01)00103-1] [PMID: 11708754]
- [90] Li C, Knierim B, Manisseri C, et al. Comparison of dilute acid and ionic liquid pretreatment of switchgrass: Biomass recalcitrance, delignification and enzymatic saccharification. Bioresour Technol 2010; 101(13): 4900-6.
[http://dx.doi.org/10.1016/j.biortech.2009.10.066] [PMID: 19945861]
- [91] Arora R, Manisseri C, Li C, et al. Monitoring and analyzing process streams towards understanding ionic liquid pretreatment of switchgrass (*Panicum virgatum* L.). BioEnergy Res 2010; 3: 134-45.
[http://dx.doi.org/10.1007/s12155-010-9087-1]
- [92] Zhao H, Baker GA, Cowins JV. Fast enzymatic saccharification of switchgrass after pretreatment with ionic liquids. Biotechnol Prog 2010; 26(1): 127-33.
[http://dx.doi.org/10.1002/btpr.331] [PMID: 19918908]
- [93] Hu Z, Wang Y, Wen Z. Alkali (NaOH) pretreatment of switchgrass by radio frequency-based dielectric heating. Appl Biochem Biotechnol 2008; 148(1-3): 71-81.
[http://dx.doi.org/10.1007/s12010-007-8083-1] [PMID: 18418741]
- [94] Sassner P, Mårtensson CG, Galbe M, Zacchi G. Steam pretreatment of H(2)SO(4)-impregnated Salix for the production of bioethanol. Bioresour Technol 2008; 99(1): 137-45.
[http://dx.doi.org/10.1016/j.biortech.2006.11.039] [PMID: 17223555]
- [95] Sassner P, Galbe M, Zacchi G. Steam pretreatment of Salix with and without SO₂ impregnation for production of bioethanol. Appl Biochem Biotechnol 2005; 121-124: 1101-17.
[http://dx.doi.org/10.1385/ABAB:124:1-3:1101] [PMID: 15930584]
- [96] Li M, Foster C, Kelkar S, et al. Structural characterization of alkaline hydrogen peroxide pretreated grasses exhibiting diverse lignin phenotypes. Biotechnol Biofuels 2012; 5(1): 38.
[http://dx.doi.org/10.1186/1754-6834-5-38] [PMID: 22672858]
- [97] Bals B, Rogers C, Jin M, Balan V, Dale B. Evaluation of ammonia fibre expansion (AFEX) pretreatment for enzymatic hydrolysis of switchgrass harvested in different seasons and locations. Biotechnol Biofuels 2010; 3(1): 1.
[http://dx.doi.org/10.1186/1754-6834-3-1] [PMID: 20047650]
- [98] Esterbauer H, Steiner W. Production of *Trichoderma* cellulase in laboratory and pilot scale. Bioresour Technol 1991; 36: 51-65.
[http://dx.doi.org/10.1016/0960-8524(91)90099-6]
- [99] Jorgensen H, Erriksson T. Purification and characterization of five cellulases and xylanases from *Penicillium brasilianum* IBT 20888. Enzyme Microb Technol 2003; 32: 851-61.
[http://dx.doi.org/10.1016/S0141-0229(03)00056-5]
- [100] Nieves RA, Ehrman CI. Technical communication: survey and commercial cellulose preparations suitable for biomass conversion to ethanol. World J Microbiol Biotechnol 1998; 14: 301-4.
[http://dx.doi.org/10.1023/A:1008871205580]
- [101] Akin DE, Rigsby LL, Sethuraman A, Morrison WH III, Gamble GR, Eriksson KE. Alterations in structure, chemistry, and biodegradability of grass lignocellulose treated with the white rot fungi *Ceriporiopsis subvermispora* and *Cyathus stercoreus*. Appl Environ Microbiol 1995; 61(4): 1591-8.
[http://dx.doi.org/10.1128/AEM.61.4.1591-1598.1995] [PMID: 7747973]
- [102] Gold MH, Alic M. Molecular biology of the lignin-degrading basidiomycete *Phanerochaete chrysosporium*. Microbiol Rev 1993;

- 57(3): 605-22.
[<http://dx.doi.org/10.1128/MMBR.57.3.605-622.1993>] [PMID: 8246842]
- [103] Ruggeri B, Sassi G. Experimental sensitivity analysis of a trickle bed bioreactor for lignin peroxidases production by *Phanerochaete chrysosporium*. *Process Biochem* 2003; 38(12): 1669-76.
[[http://dx.doi.org/10.1016/S0032-9592\(02\)00199-1](http://dx.doi.org/10.1016/S0032-9592(02)00199-1)]
- [104] Bosco F, Ruggeri B, Sassi G. Performances of a trickle bed reactor (TBR) for exoenzyme production by *Phanerochaete chrysosporium*: influence of a superficial liquid velocity. *Chem Eng Sci* 1999; 54: 3163-9.
[[http://dx.doi.org/10.1016/S0009-2509\(98\)00365-0](http://dx.doi.org/10.1016/S0009-2509(98)00365-0)]
- [105] Arora DS, Chander M, Gill PK. Involvement of lignin peroxidase, manganese peroxidase and laccase in the degradation and selective ligninolysis of wheat straw. *Int Biotechnol Biodegrad* 2002; 50: 115-20.
[[http://dx.doi.org/10.1016/S0964-8305\(02\)00064-1](http://dx.doi.org/10.1016/S0964-8305(02)00064-1)]
- [106] Miller RC Jr, Gilkes NR. Similarities between bacterial and fungal cellulases systems. *Proc of the 6th Int Conference on Biotechnology in the Pulp and Paper Industry: Advances in Applied and Fundamental Research*. 531-42.
- [107] Tigunova OO, Shulga SM. New producer strains of biobutanol. I. Isolation and identification. *Biotechnol Acta* 2013; 6(1): 97-104.
[<http://dx.doi.org/10.15407/biotech6.01.097>]
- [108] Mauerhofer LM, Pappenreiter P, Paulik C, Seifert AH, Bernacchi S, Rittmann SKR. Methods for quantification of growth and productivity in anaerobic microbiology and biotechnology. *Folia Microbiol (Praha)* 2019; 64(3): 321-60.
[<http://dx.doi.org/10.1007/s12223-018-0658-4>] [PMID: 30446943]
- [109] Al-Shorgani NKN, Shukor H, Abdeslahian P, Kalil MS, Yusoff WMW, Hamid AA. Enhanced butanol production by optimization of medium parameters using *Clostridium acetobutylicum* YM1. *Saudi J Biol Sci* 2018; 25(7): 1308-21.
[<http://dx.doi.org/10.1016/j.sjbs.2016.02.017>] [PMID: 30505175]

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