

RESEARCH ARTICLE

OPTIMIZATION OF FERMENTATION CONDITIONS FOR THE ETHANOL PRODUCTION FROM SUGARCANE BAGASSE BY ZYMOMONAS MOBILIS USING RESPONSE SURFACE METHODOLOGY.

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Abstract

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The use of biomass as renewable sources of energy has increased industrial focus toward alternative fuel because of the depletion of fossil fuel reserves, the unstable panorama of the petrol prices, the increasing environmental and political pressures. The new concept of bioethanol corresponds to its production using raw cellulosic materials, such as sugarcane bagasse. The bacterium Zymomonas mobilis was shown to be extremely attractive for the ethanol second generation production from glucose of the cellulosic fraction, due to its high capacity to absorb this sugar, resulting in high ethanol productivity values. The aim of this work was to study the influence between the medium components: Yeast Extract, KH₂PO₄, (NH₄)₂SO₄, MgSO₄.7H₂O to optimize the fermentation conditions for the ethanol production from sugarcane bagasse by Z. mobilis. Initially, to make easier the accessibility of cellulases to the cellulose microfibrils, the bagasse was submitted to a pretreatment with diluted acid to fractionate and extract the hemicellulose component from the solid residue named cellulignin and then, this solid residue was pretreated using NaOH (4%) aiming at its partial delignification. Thereafter, the pretreated cellulignin underwent the action of a commercial cellulolytic preparation, allowing the conversion from cellulose to glucose. This enzymatic pretreatment occurred under temperature of 50°C for 12 hours, after which the temperature reduced to 30°C and we inoculated the system with cells of Z. mobilis. It has been used statistical experimental design to optimize the conditions of SSF, evaluating the medium components. The optimum conditions found were 12.5 g/L of Yeast Extract, 2.5 g/L of KH₂PO₄; 1.5 g/L of (NH₄)₂SO₄; 1.5 g/L of MgSO₄.7H₂O, respectively; achieving 65.3 g/L of ethanol.

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Introduction:-

For the future, completely new approaches in research and development a re-arrangement of a sustainable economy to biological raw materials, production, and economy are necessary (1). The new concept of bioethanol corresponds

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to its production from conventional raw materials such as sugarcane, beet or starch, due to their low-cost, availability and renewability. The use of biofuels can contribute to the mitigation of GHG emissions, provide a clean and therefore sustainable energy source, and increase the agricultural income for rural poor in developing countries (2). Developing countries have a comparative advantage for biofuel production because of greater availability of land, favorable climatic conditions for agriculture and lower labour costs. However, there may be other socio-economic and environmental implications affecting the potential for developing countries to benefit from the increased global demand for biofuel (3).

Ethanol production technology based on lignocellulosic biomass uses chemical and enzymatic processes for the hydrolysis of cellulose, which produces carbohydrates (sugars) and then fermented into ethanol (4). Bioethanol is by far the most widely used biofuel for transportation worldwide. Bioethanol and bioethanol/gasoline blends have a long history as alternative transportation fuels. Germany and France have used it as early as 1894 by the then incipient industry of internal combustion engines (ICEs) (5). Brazil has utilized bioethanol as a fuel since 1925. By that time, the production of bioethanol was 70 times bigger than the production and consumption of petrol (6).

The largest ethanol producing countries are USA and Brazil, responsible for the production of 54×10^6 and 21×10^6 m³ in 2011, respectively (7). Brazil utilizes sugarcane for bioethanol production while the United States and Europe mainly use starch from corn, and from wheat and barley, respectively. Sugarcane as a biofuel crop has much expanded in the last decade, yielding anhydrous bioethanol (gasoline additive) and hydrated bioethanol by fermentation and distillation of sugarcane juice and molasses (8). Brazil's estimative of sugarcane yield production averages about 617 x 10^6 tons in 2017 (9). Brazil is the largest single producer of sugarcane with about 31% of global production (10). It has nearly 9 million hectares of sugarcane under cultivation. Sugar beet crops grow in most of the EU-25 countries, and yield substantially more bioethanol per hectare than wheat.

The acid pretreatment causes the hydrolysis of hemicellulose under conditions of low-grade severity (acid concentration, temperature, time of exposure and solid: liquid relationship). After pretreatment step, two fractions are generated: a liquid containing the hemicellulose hydrolyzed and a solid called cellulignin. This solid waste needs alkaline treatment aimed at the partial removal of lignin through the structural separation of the connections between the phenolic macromolecule and cellulose, increasing thus the accessibility of enzymes to cellulose fibers. The raw materials of lignocellulosic origin contain 20% to 60% of cellulose that convert into glucose by enzymatic action. Glucose is a monosaccharide used by all major microorganisms, making it an important building block to obtain substances that ranging from fuels to polymers (11).

The research on ethanol have been targeted for the process of Saccharification and Simultaneous Fermentation (SSF), which combines in one-step enzymatic hydrolysis and fermentation of glucose coming from the cellulose hydrolysis (12). This strategy is justified because the hydrolysis products (glucose and cellobiose) inhibit the cellulase complex enzymes. The SSF process of waste materials provides the use of available low-cost substrates. These processes have been employed in various pilot plants. Moreover, the SSF process permits high productivity of ethanol, using fewer amounts of enzymes (13), when compared to those processes utilizing separated enzymatic hydrolysis and fermentation.

The most frequently used microorganism for fermenting bioethanol in industrial processes is *S. cerevisiae*, which has proved to be very robust and well suited to the fermentation of lignocellulosic hydrolysates(14). For the expansion of ethanol industry, it is necessary the searching for more competitive ethanologenic microorganisms. As they have been studied, the bacterial specie *Zymomonas mobilis* has attracted the attention of researchers and showed as a promising ethanol producer (15, 16). Due to its high fermentation potential, *Zymomonas mobilis* have been the subject of numerous studies, which results in a production of ethanol comparable to or even greater than that obtained by yeasts (17, 18, 19, 20, 21, 22). The synthesis of ethanol is conventionally performed by yeast, but the bacterial *Zymomonas sp.* has a special ability for this production, with an attractive alternative to the current global demand for fuel. When compared with *Saccharomyces cerevisiae*, high specific productivity and high tolerance to ethanol and lower production of biomass is achieved.

These microorganisms uses tiny fraction of sugar as carbon source, approximately 98% destined for the fermentation and only 2% for growth. Like much of chemoorganotrophic organisms, the bacterial *Z. mobilis* need sources of nitrogen, phosphorus, sulfur, and micronutrients for the metabolism functioning and synthesis of the cells

in a form assailable by the microorganism (23). They also require water, carbon, oxygen and growth factor (24), hence the great need for optimizing the optimal concentrations of nutrients for bacterial metabolism.

Objective:-

The aim of this work was to study the influence of the medium components (Yeast Extract, KH₂PO₄, [NH₄]₂SO₄, MgSO₄.7H₂O) on the fermentation conditions for the ethanol production from sugarcane bagasse by *Z. mobilis* using Simultaneous Saccharification and Fermentation technology.

Materials and Methods:-

Substrate: Costa Pinto Distillery (SP, Brazil) kindly provided the sugarcane bagasse (Saccharum spp.).

Pretreatment: The bagasse was primarily hydrolyzed, in order to disorganize the lignocellulosic material and remove the hemicellulose fraction. The following conditions for acid pretreatment were 1% of H_2SO_4 , solid-liquid ratio of 1:2 (g/ml), temperature of 121°C for 30 minutes (25). The hemicellulose was removed, and the remaining solid residue (cellulignin) was partially delignified by an alkaline pretreatment with 4% NaOH, using solid-liquid ratio of 1:20 (g/ml), at 121°C for 30 minutes (26). The cellulignin was washed with distilled water until the aqueous phase remained clear. Resulting solid material with increased cellulose accessibility was subjected to the enzymatic cellulose hydrolysis, which occurred using the commercial preparation Multifect (Genencor, USA) at a temperature of 50°C, 12 hours, enzyme load of 25 FPU/g and solid: liquid ratio of 3:10 (g/ml) (27). The solid fraction is the cellulose and the liquid is the fermentation medium, containing other nutrients.

Microorganism and Inoculum Preparation: The Department of Antibiotics of the Federal University of Pernambuco, Brazil, kindly provided the *Zymomonas mobilis* CP4 used in this study. The strain was grown in a liquid medium (20 g/L of glucose and 5 g/L of yeast extract) recommended by Swing & De Ley (*16*), at 30°C for 24 hours, and maintained at 4°C. Monthly transferences were performed for maintenance of cell viability. The inoculum was grown in a medium composed of glucose, 20 g/L; yeast extract, 2.5 g/L; Ammonium Sulfate, 1 g/L; Potassium Phosphate, 1 g/L and Magnesium Sulfate, 0.5 g/L. The cultures were shaken in a rotary shaker at 150 rpm, at 30°C for 20 hours. After growth, the cells were centrifuged (8000 rpm, 20 minutes), and their concentration was determined by measuring the optical density of a diluted sample at 600 nm (SPECTRUMLAB 22 PC), using a standard curve of absorbance against dry cell mass. The centrifuged cells were inoculated in the enzymatically pretreated solids.

Fermentation Assays: The fermentation medium (enzymatically pretreated solids) was supplemented with the same nutrients used for the inoculum preparation medium, without glucose, providing the technical implementation of the SSF process. Batch fermentations experiments were performed in 500mL Erlenmeyer flasks with a working volume of 100mL to define the optimum process conditions. Additionally, fermentations were carried out in a 1.5L-bioreactor (BIOFLO III, New Brunswick Scientific, USA) with control of temperature, pH and agitation. The reactor operated with a working volume of 500mL and the temperature and pH were set at 30° C and 5.0, respectively. The pH was monitored using a sterile pH electrode and controlled by adding 1M KOH. The kinetics of SSF in bioreactor was evaluated in the optimum conditions established in the shake flask experiments.

Analytical methods: Cell quantification was determined as described previously. Samples were analyzed for glucose, cellobiose and ethanol concentrations by high-efficiency liquid chromatography (HPLC) using the chromatographic system (WATERS) consisting of a HPX-87P (Bio-Rad) column, WATERS 510 pump, a refractive index detector WATERS 410 and Empower softwareTM integrator. The standard solution consists of cellobiose, glucose and ethanol concentrations of 5 g/L, 10 g/L and 15 g/L, respectively. The end of fermentation was determined through the stabilization of the production of ethanol, as verified by the reading of two consecutive and equal values of ethanol content.

Optimization of ethanol concentration: Response surface methodology (RSM) is the one suitable for identifying the effect of individual variables and for seeking the optimum conditions for a multivariable system efficiently (28). The statistical analysis of the data was performed using "Design Expert" software (7.1.6., Stat-Ease). As seen in **Table 1,** a complete factorial design 2^4 was done, where the parameters analyzed were concentrations of yeast extract (g/L), KH₂PO₄ (g/L), (NH₄)₂SO₄ (g/L) and MgSO₄.7H₂O (g/L) added to cellulignin in the beginning of hydrolysis, which along with glucose generated, provided the technical implementation of SSF.

Tuble 1. Independent variables in the experimental design.						
PARAMETERS		Real Levels				
		-α	-	0	+	$+\alpha$
Α	Yeast Extract (g/L)	0	5	12.5	20	27
В	KH_2PO_4 (g/L)	0	1	2.5	4	5.5
С	$(NH_4)_2SO_4(g/L)$	0	0.5	1.25	2	2.75
D	$MgS0_{4}.7H_{2}0(g/L)$	0	0.5	1.25	2	2.75

Table 1:- Independent variables in the experimental design.

Results and Discussion:-

This session will present the experiments concerning the use of the experimental design, evaluating the influence of medium components (yeast extract, KH_2PO_4 , $(NH_4)_2SO_4$ and $MgSO_4.7H_2O$ on the Simultaneous Saccharification and Fermentation technology. The complete 2^4 factorial design generated 30 runs (24 independent runs and 6 repetitions of central point), shown in **Table 2**.

Table 2:- 2^4 Central composite design investigating effects of adding yeast extract (g/L), KH₂PO₄ (g/L), (NH₄)₂SO₄ (g/L) and MgSO₄.7H₂O (g/L) in the ethanol production.

Yeast Extract	KH ₂ PO ₄	(NH ₄) ₂ SO4	MgS04.7H20	Ethanol	
(g/L)	(g/L)	(g/L)	(\mathbf{g}/\mathbf{L})	(g/L)	
5	1	0.5	0.5	54	
20	1	0.5	0.5	62	
5	4	0.5	0.5	45	
20	4	0.5	0.5	59	
5	1	2	0.5	38	
20	1	2	0.5	60	
5	4	2	0.5	34	
20	4	2	0.5	52	
5	1	0.5	2	32	
20	1	0.5	2	57	
5	4	0.5	2	33	
20	4	0.5	2	56	
5	1	2	2	28	
20	1	2	2	54	
5	4	2	2	48	
20	4	2	2	63	
0	2.5	1.25	1.25	30	
27.5	2.5	1.25	1.25	65	
12.5	0	1.25	1.25	33	
12.5	5.5	1.25	1.25	38	
12.5	2.5	0	1.25	34	
12.5	2.5	2.75	1.25	37	
12.5	2.5	1.25	0	34	
12.5	2.5	1.25	2.75	62	
12.5	2.5	1.25	1.25	65	
12.5	2.5	1.25	1.25	63	
12.5	2.5	1.25	1.25	62	
12.5	2.5	1.25	1.25	61	
12.5	2.5	1.25	1.25	62	
12.5	2.5	1.25	1.25	60	

The highest ethanol concentration obtained was 65 g/L, when 12.5 g/L yeast extract, 2.5 g/L KH₂PO₄, 1.25 g/L (NH₄)₂SO₄ and 0.5 g/L MgSO₄.7H₂O were used. The presence of yeast extract, rich in nitrogen, significantly increases the production of ethanol, as indicated by the comparison of runs 9 and 10. These experiments have the same concentrations of other nutrients and when yeast extract was added, ethanol production increased from 33 g/L to 57 g/L. However, the presence of high concentrations of this nutrient causes a decrease in ethanol production due

to an excess of nitrogen source (29). Soleimani *et al.* (2012) also found that ethanol production was reduced after removal of nutrients from the culture medium, such as yeast extract and peptone (30). However, some studies have shown that the use of some agro-industrial waste can substitute carbon or nitrogen sources, as observed by Patle&Lal (2008) (31).

In the other hand, Neto *et al.* (2005) found that large concentrations of yeast extract may cause a decrease in ethanol production due to an excess of nitrogen source. Thus, there is an increase in the biomass concentration, since *Zymomonas sp.* uses this compound not only as a source of nitrogen, but also as blocks for biosynthesis, implying a lower energy requirement. This fact would justify the decrease in the production of ethanol and, consequently, the lower ATP synthesis. The regulation of nitrogen is of fundamental importance in industrial microbiology, since it affects the enzymes meabolization from primary and secondary metabolism. Thus, many secondary metabolic pathways are negatively affected by growth-enhancing nitrogen sources, such as ammonium salts, as well as high concentrations of nitrogen can affect the synthesis of these enzymes (Belalch&Senez, 1965; Neto et al., 2005) (32,29).

Thus, there is an increase in biomass concentration as *Zymomonas sp.* uses this compound not only as a nitrogen source but also as building blocks for biosynthesis, resulting in a reduced need for energy, which would justify a reduction in ethanol production and, consequently, less formation of ATP. The synthesis of enzymes using other nitrogen sources is repressed until the primary substrate is completely depleted.

In experiments 17, 19, 21 and 23, without addition of yeast extract (g/L), KH_2PO_4 (g/L), $(NH_4)_2SO_4$ (g/L) and $MgSO_4.7H_2O$ (g/L), respectively, ethanol is still produced, due the presence of 44.6% carbon, 5.8% hydrogen, 44.5% oxygen, 0.6% nitrogen, 0.1% sulfur and 4.4% other elements in sugarcane bagasse (24,33). Nonetheless, there is still need for the presence of nutrients that contain chemical elements such as magnesium and phosphorus.

Cao *et al.* (*34*) have reported that KH_2PO_4 , $(NH_4)_2SO_4$, and $MgSO_4$ would influence the growth and metabolism of the yeast cells. Phosphorus is present in KH_2PO_4 , witch has an important role in metabolic pathways that are initiated with substrate phosphorylation. It is also a constituent of ATP molecules that are present in the energy mechanism of the cells. Methionine, cysteine and sulfates can supply Sulfur, present in $(NH_4)_2SO_4$ structural constituent of the cell and of great importance for the formation of proteins (*35*). $MgSO_4$ is the best sulfur source, and serves as a source of magnesium, which is responsible for the structural stability of several enzymes, but also for preventing the formation of vesicles on the outer membrane of the cell (*36*).

The presence of these nutrients in their highest levels associated with increased extract concentration, promotes increased ethanol production, as shown in the experimental 16. Therefore, there is a high production of ethanol when the concentration of KH_2PO_4 (g/L) increases in relation to increasing MgSO₄.7H₂O (g/L) and a smaller increase through its association with $(NH_4)_2SO_4$ (g/L), besides there is a decline of ethanol when the nutrients are associated at high concentrations. The increase in nutrients should occur simultaneously, as shown when experiment 1, which resulted in 54 g/L of ethanol is compared to experiment 15 where there was an increase in nutrients, except yeast extract, thus decreasing the ethanol production to 48 g/L. Therefore, a synergistic effect between the parameters analyzed is necessary.

Statistical analysis: The Statistical analysis of variance obtained by the factorial design indicated in **Table 3**, shows that the model was very significant, with (p<0.05), as well as the coefficient of total determination (\mathbb{R}^2) observed for the response to ethanol production in g/L, was 0.995, suggesting a good model fitting to experimental data. Moreover, the residue was low and not significant (p>0.05), which did not invalidate the model for predictive purposes, because the equation had a high \mathbb{R}^2 . The resulting ethanol concentration model is in equation (1).

[Ethanol]=+52.17+131.29*A+3.05*B+2.50 * C+6.59 *D-1.00*A*B+0.38*A*C+2.00 *A*D+1.50*B * C+3.63 * B * D+3.50 * C * D+8.65 * A2-5.07 * B2-5.04 * C2-2.59* D2-0.88 *A*B*C -1.25*A*B*D-1.88 * A*C*D+1.50*B*C*D-2.43 *A2*B-3.50*A2* C-8.96*A2* D-85.12*A*B2-36.42*A3 (1)

SS- Sum of Squares, DI-Degree of Heedom, MS-Medium Square.							
Source	SS	DF	Ms	F value	<i>p</i> > F		
Model	4693.84	12	391.15	78.31	< 0.0001		
Residual	74,89583	6	12,48264				
Lack of fit	60,0625	1	60,0625	20,24579	0.0064		
Pure error	14,83333	5	2,966667				
Cor. total	4505,367	29	4505,367	29	4505,367		

Table 3:-Analysis of variance in the ethanol production [Partial sum of squares]. SS= Sum of Squares: DF=Degree of Freedom: MS=Medium Square.

The variable A (yeast extract) presents the highest influence, followed by D (MgSO₄.7H₂O), which in turn is higher than B (KH₂PO₄) and finally C ([NH₄]₂SO₄), which has low values of Sum of Squares. The interaction of these four variables also resulted in high values of Fisher, indicating a great synergism between nutrients in the fermentation by *Z. mobilis*, providing increased amounts of ethanol. The interactions between nutrients complementary pairs had to be quite significant. The most influential was between BD (KH₂PO₄ and MgSO₄.7H₂O), BC (KH₂PO₄ and INH₄]₂SO₄), followed by CD ([NH₄]₂SO₄ and MgSO₄.7H₂O), as shown in **Figure 1**.



Sum of Squares

Figure 1:- Histogram of variables influence, according to the Sum of Squares.

The double interactions between nutrients analyzed for ethanol production display the interaction between KH_2PO_4 (g/L) and Yeast extract (g/L) are very low and there is no intersection between the two variables. This, in turn possesses high interactions with the variables represented by $(NH_4)_2SO_4$ (g/L) and $MgSO_4.7H_2O$ (g/L).

Figures 2 show the 3-D response surface plots demonstrate interaction between the parameters A, B, C and D, which represent yeast extract (g/L), KH_2PO_4 (g/L), $(NH_4)_2SO_4$ (g/L) and $MgSO_4.7H_2O$ (g/L), respectively. Note that in **Figure 2A** in connection with MgSO4 and $(NH_4)_2SO_4$ with KH_2PO_4 and yeast extract in their central points, ethanol production is at its optimal level when the variable D is at its center point to the higher level and the variable C is at the central point, respectively.



Figure 2 (A):- Response surface plot showing the effect of yeast extract (g/L), KH_2PO_4 (g/L), and their combined effects on ethanol concentration.

Figure 2B shows that ethanol production is at its optimal level when the variables C and D, respectively, are from their central points to their highest levels. Underscoring the results presented in the experiments, which exhibits the highest production of ethanol in the central points of the parameters, although the same values were obtained at higher levels indicating that even with increased concentrations of nutrients, with the same concentrations of enzymes, solid and cells, the concentration of ethanol is stagnant, starting inhibition by large amounts of nutrients, affecting the osmolarity of the cell or the growth of bacteria exacerbated by shifting the focus of the metabolism of ethanol production to the growth of bacteria.



Figure 2 (B):- Response surface plot showing the effect of MgSO₄.7H₂O (g/L), (NH₄)₂SO₄and their combined effects on ethanol concentration.

The volumetric productivity and production of ethanol depends, among several factors, according to the substrate used, additional nutrients, as well as the microorganism to be used. Pinilla*et al.* (2011) obtained high concentrations of ethanol (83.81 g/L), from glucose added of yeast extract, peptone and salts, after isolation of colonies grown on sugarcane molasses (*37*). In the present study, the use of sugarcane, fructose, glucose, sucrose and galactose mixtures was achieved by Wiikins (2009), reaching 43.5 g/L ethanol at the end of the fermentation process (*38*), whereas Maiti*et al.* (2011) reached 58.4 g/L from sugarcane molasses (*39*).

These results are similar to observable by Yu et al. (28), which optimized the nutrient concentrations in synthetic medium and noted a significant effect of yeast extract on biomass, associated to the carbon source. They observed an increase in Ammonium Sulfate concentration that resulted in an increase in ethanol yield, but above an optimum level the ethanol yield decreased. No significant variation was observed with the phosphate and nitrogen source, resulting in an insignificant role for the Potassium Phosphate and for the Ammonium Sulphate, as the cells can utilize nitrogen available from yeast extract (36). Excess nitrogen could lead to more biomass but less ethanol production, as shown in this work.

The experimental validation in bioreactor yielded a SSF initial glucose concentration of 85 g/L, final ethanol concentration of 65 g/L and volumetric productivity of 2.70 g.L⁻¹.h⁻¹, with a fermentation time of approximately 24 hours, at temperature of 30°C, orbital agitation at 150rpm and pH 5 (**Figure 3**).



Figure 3:- Validation of the experimental design optimum conditions (S:L=1:3; EL=25 FPU/g; X_o=4g/L) for the SSF process in bioreactor with *Zymomonasmobilis* CP4.

Conclusions:-

The experiments showed that it is possible to optimize ethanol production by examining the addition of nutrients in the fermentation medium, through the experimental designs of response surfaces. All the nutrients proved to be significant and essential to the metabolism of bacteria. The increase of yeast extract was mainly responsible for the increase of ethanol concentration, followed by KH_2PO_4 , $MgSO_4.7H_2O$ and $(NH_4)_2SO_4$, that were less significant as independent variables, but showed high levels of sum of squares, when they were involved in all the interactions studied.

The optimal conditions found were yeast extract (12.5 g/L), KH_2PO_4 (2.5 g/L), $(NH_4)_2SO_4$ (1.25 g/L) and $MgSO_4$ (1.25 g/L). It resulted in the maximum ethanol concentration of 65 g/L, with 85 g/L of SSF initial glucose concentration, reaching the highest volumetric productivity of 2.63 g/L.h, at temperature 30°C, orbital agitation at 150rpm, pH 5 in bioreactor.

It was verified, therefore, that cellulose constitutes an excellent source of carbohydrates for the execution of the process of simultaneous enzymatic hydrolysis and fermentation by *Zymomonasmobilis*, which presented promising for the production of this biofuel, due to its high capacity of glucose absorption, High specific rates of ethanol production, resulting in high productivity values. The results obtained with the present work were satisfactory, however, it is necessary to continue with the elaboration of new strategies so that the inhibitory and unsatisfactory issues placed throughout the text are circumvented, as well as the development of molecular biology techniques for the Evidence of genetic transformation, generating opportunities for future and interesting technological developments.

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