



Article Biological Hydrogen Production by Dark Fermentation in a Stirred Tank Reactor and Its Correlation with the pH Time Evolution

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Abstract: Dark fermentation is a hydrogen generating process carried out by anaerobic spore-forming bacteria that metabolize carbon sources producing gas and short-chain acids. The process can be controlled, and the hydrogen harvested if bacteria are grown in a reactor with favorable conditions. In this work, bacteria selected from natural sources were grown with a defined culture media, while pH was monitored, with the aim of relating the amount of generated hydrogen to the increase in hydron ion concentration. Therefore, a model based on the acid-base species mass balance is proposed and solved to estimate the lag phase time and measure the hydrogen production efficiency and kinetics. Hydrogen production in a stirred batch reactor was performed for 150–200 h, at given operating conditions using a previously defined growth media, to validate the model. Using the proposed model, the cumulated moles of produced hydrogen correlate well with those predicted from the pH curve. Hence, the modified Gompertz model parameters, largely used for describing the hydrogen generation kinetics by dark fermentation, were estimated from the pH curve and from the experimentally measured generated hydrogen. Satisfactory agreement was found, thus, validating the method.

Keywords: biohydrogen; dark fermentation; pH influence; Gompertz model

1. Introduction

The energy matrix is of deep concern to humanity. Modern life has become energy intensive [1], and energy problems are very disruptive in our society [2]. We depend on energy for transportation, food production, goods manufacturing, communications, heating, cooling, and entertainment [3]. The most common energy sources we use today are fossil fuels, which are non-renewable. Oil will slowly become scarce and increasingly expensive before running out. Another downside of using fossil fuels is that they are the most important source of humanity-driven climate change [4,5]. Additionally, because oil reserves are localized in a few countries, their availability is centralized and is often the source of international conflicts [6]. Hence, alternative energy sources to fossil fuels are becoming increasingly important. Sustainable fuels play a particularly important role in this scenario since the transition into a sustainable economy should also mean energy security. The research community has, thus, strongly devoted itself to developing alternative energy from renewable sources. The transition to more sustainable and carbon-free energy sources



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Copyright: © 2022 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). has brought hydrogen into the spotlight, as a clean energy vector. It is one of the candidates for diversifying the energy matrix due to its high energy yield and clean emissions. Then, hydrogen production processes are extensively investigated, particularly using renewable resources [7,8]. A clean and sustainable way of producing hydrogen is from biomass.

Biomass refers to all masses of living organisms, such as bacteria, fungi, plants, animals, or their remains [9]. Therefore, biomass can be transformed and absorbed by the Earth's ecosystems as part of the carbon cycle. Energy derived from biomass can then be considered sustainable [10]. Hydrogen can be produced from biomass using natural bacterial communities. The bioprocess by which anaerobic bacteria ferment carbohydrates releasing H₂ is called dark fermentation. This process can be carried out in rather simple fermentation tanks and potentially works with direct carbon sources, such as fermentable sugars, but also degrading second-generation biomass from crops or animal residues from livestock, among others [11]. Hydrogen-producing bacteria can be collected from several natural sources. In a previous contribution, the authors reported evidence on the feasibility of using bacterial communities from different sources [12,13]. The one extracted directly from a wastewater treatment plant on board a navy ship was selected, considering the convenience of in situ production for ship maintenance. Before its usage, the source was submitted to a heat pre-treatment to eliminate hydrogen trophic bacteria and retain the producers, which were mainly *Clostridium* sp. spore-forming bacteria [14,15].

The conditions inside the bioreactor require an optimized media to promote bacterial growth, since hydrogen is a byproduct of their metabolism, along with carbon dioxide. The gas mixture can be easily collected and used as a green energy vector. Although it can be stored, in situ consumption for feeding an onboard proton exchange membrane (PEM) fuel cell is an attractive alternative [16]. Moreover, it can also be devised for being used in hydrogenation processes, if any is carried out on board.

Dark fermentation is a stage in the natural degradation of organic material, which usually ends up in methane and carbon dioxide, which are greenhouse gases [17]. However, by eliminating the non-hydrogen-producing bacteria through a pre-treatment, it is possible to produce hydrogen with high efficiency, mixed only with carbon dioxide [15,18]. The gas mixture produced, which contains only hydrogen and carbon dioxide in a ratio of around 2:1, is very convenient for a PEM fuel cell operation, with almost no need for a downstream purification step. Moreover, the gas has no carbon monoxide, as in gas reforming, which poisons PEM fuel cells electrocatalysts [19].

The amount of gas produced is less than in a methanogenic fermentation because the carbon atoms that would form methane and carbon dioxide mostly become shortchain carboxylic acids in the effluent. Mainly, acetic and butyric acids are formed [20,21]. Therefore, the dark fermentation of selected hydrogen-producing bacteria strongly modifies the acidity of the cultivation media [22–24]. As the pH lowers and the acid concentration increases in the culture media, hydrogen production starts slowing down, showing pH has an inhibitory effect [11,25]. The pH and the developed hydrogen is, thus, strongly linked, and controlling pH has been proposed to improve hydrogen generation [24,26–28]. Although there are works focusing on the influence of pH, a quantitative relationship between pH variation and hydrogen generation that allows obtaining kinetic information on H₂ production has not been proposed. The pH curve can be readily measured with an appropriate sensor and monitored during the whole fermentation. Therefore, this work aims at exploiting the information provided by the pH curve to infer the hydrogen production efficiency and kinetics.

2. Results and Discussion

2.1. Hydrogen Measure and Production

Typical results of the moles of produced hydrogen superimposed on the evolution of the pH in the reactor as a function of time are shown in Figure 1. The generated hydrogen is expressed as the number of moles divided by the time during which the gas was collected. As observed, the largest hydrogen production is associated with a sharp decrease in pH, in agreement with the generally accepted mechanism that suggests simultaneous production of short-chain acids. Acetic and butyric acids have been found during dark fermentation of different starting materials when *Clostridium* species are dominant, as in this case [14].



Figure 1. The pH (left) and hydrogen production (right) evolution over time.

2.2. Relation between the Evolved Hydrogen and the pH Curve

In this section, a model oriented to explain the relationship between the observed production of H_2 as a function of the pH evolution over time is developed and resolved. Considering the simplified acetogenic reaction path [20,29], Equation (1) indicates that the evolution of hydrogen is linked to the formation of acid species. According to Equation (1), as hydrogen is generated, acetic acid and carbon dioxide are produced. Although other acids can be formed, leading to a lower ratio of hydrogen to acid per gram of substrate, we assume this route in the analysis to be representative. If other routes are apparent, they will be reflected in a lower estimated efficiency.

Substrate +
$$H_2O \rightarrow 2AcOH + 2CO_2 + 4H_2$$
 (1)

2.3. Correlation between pH Evolution and Hydrogen Production

The monitored pH curve shows an initial period of constant value, corresponding to the lag phase of the microorganism culture. Then, the culture activity starts with an abrupt change in pH towards acidic conditions until several buffer stages are reached. The buffer stages are likely due to the action of phosphate, dicarboxylic acids, and by the mixture of saturated carboxylic acids, dominated by acetate (pKa = 4.75) from the media.

Considering a generic situation where m weak bases (W_b^{n-}) are used as alkalizing agents, and that the generation of acetic acid and carbon dioxide are linked to hydrogen generation, the mass balance given by Equation (2) is proposed to describe the change in the observed molar concentration of protons, $[H^+]_{obs'}$ in the media. Since the formed acids are generally weak acids, they will remain partially dissociated. Variation of the molar proton concentration would depend on the generated acetic acid and on the dissociation constant and will be linked to the variation in acetate concentration. The generated carbon dioxide is supposed to be totally released to the gaseous phase together with hydrogen. Additionally, for pH < 7, carbonic acid is hardly dissociated.

$$\Delta [\mathrm{H}^{+}]_{\mathrm{obs}} = \Delta [\mathrm{AcO}^{-}] - \sum_{i=1}^{m} \Delta [\mathrm{HW}_{\mathrm{b}_{i}}^{1-\mathrm{n}}] - 2 \frac{\Delta \mathrm{n}_{\mathrm{H2-gen}}}{\mathrm{V}_{\mathrm{R}}} + 2 \frac{\Delta \mathrm{n}_{\mathrm{H2-cons}}}{\mathrm{V}_{\mathrm{R}}}$$
(2)

The generation and consumption of molecular hydrogen are assumed to take place, mediated by biochemical or abiotic means, through the reaction given by Equation (3).

$$2\mathrm{H}^{+} + 2\mathrm{e}^{-} \Leftrightarrow \mathrm{H}_{2} \tag{3}$$

It is also assumed that the entire change in acetic acid concentration and acid-base carbon dioxide species are generated by dark fermentation and stoichiometrically linked to hydrogen generation, as expressed in Equation (4).

$$\Delta[\text{AcOH}] = \Delta[\text{CO}_2] = \frac{\Delta n_{\text{H2-gen}}}{2V_{\text{R}}} = \frac{P_{\text{H2}}\Delta t}{2V_{\text{R}}}$$
(4)

where P_{H2} is the production of hydrogen due to dark fermentation. Since in the culture medium there is hydrogen degradation activity, it is proposed that:

$$\Delta n_{\rm H2-gen} - \Delta n_{\rm H2-cons} = \eta P_{\rm H2} \Delta t \tag{5}$$

where η is the net efficiency of molecular hydrogen production. Given that adenosyl phosphates are produced and consumed by biological activity, it is reasonable to consider that the concentration of inorganic phosphates present in the medium is constant regardless of whether they are located within or outside the cells. In addition, since the pH of the time series, shown in Figure 1, is between 8 and 5, the dominant phosphate acid-base balance is its second protonation (Equation (6)):

$$\begin{aligned} \left[Phos \right]_{o} &= \left[H.PO_{4}^{2-} \right] + \left[H_{2}PO_{4}^{-} \right] \\ \frac{\left[H.P.O_{4}^{2-} \right]}{\left[H_{2}PO_{4}^{-} \right]} &= 10^{-pKa2+pH} = \frac{K_{a2}}{\left[H^{+} \right]} \\ &= > \left[H_{2}PO_{4}^{-} \right] = \frac{\left[Phos \right]_{o}}{\left(1 + \frac{K_{a2}}{\left[H^{+} \right]} \right)} \end{aligned}$$
(6)

Analogously, for the remaining acid-base species present in the culture, $\{Wb_i\}$, a simplified ionic balance (Equation (7)) can be performed considering a dominant acid-base couple in the case of polyprotic species:

$$\begin{split} \left[Wb_{i} \right]_{o} &= \left[Wb_{i}^{-n_{i}} \right] + \left[HWb_{i}^{1-n_{i}} \right] \\ \frac{\left[Wb_{i}^{-n_{i}} \right]}{\left[HWb_{i}^{1-n_{i}} \right]} &= 10^{-pKa_{i}+pH} = \frac{K_{ai}}{\left[H^{+} \right]} \\ &= > \left[HWb_{i}^{1-n_{i}} \right] = \frac{\left[Wb_{i} \right]_{o}}{\left(1 + \frac{K_{ai}}{\left[H^{+} \right]} \right)} \end{split}$$
(7)

The variation in acetate concentration is linked to the amount of acetic acid produced, through the acetic acid constant. Acetic acid will partially dissociate, generating acetate. Although there is acetate in the medium, which would regulate the dissociation degree, Equation (1), in this case, is representative of any of the short chain carboxylic acids formed. Hence, we assume a variation of acetate concentration as the one expected in pure water and that the dissociation constant of acetic acid, K_{AcOH}, represents a lumped constant for other weak acids formed, since they are generally in the same order (~10⁻⁵ M) [24]. The variation in acetate concentration is proposed to be approximately related to the produced acetic acid through Equation (8).

$$\Delta[\text{AcO}^{-}] \approx \frac{1}{2} \left(\text{K}_{\text{AcOH}} \right)^{1/2} \Delta[\text{AcOH}]$$
(8)

By conveniently replacing each term of the balance given in Equation (2) as a function of the proton concentration and hydrogen production, Equation (9) is obtained.

$$\Delta \left[\mathrm{H}^{+} \right]_{\mathrm{obs}} = \mathrm{K}_{\mathrm{AcOH}}^{1/2} \frac{\mathrm{P}_{\mathrm{H2}} \Delta t}{4 \mathrm{V}_{\mathrm{R}}} - \Delta \left\{ \frac{\left[\mathrm{Phos} \right]_{\mathrm{o}}}{\left(1 + \frac{\mathrm{K}_{\mathrm{a2}}}{\left[\mathrm{H}^{+} \right]_{\mathrm{obs}}} \right)} + \sum_{i=1}^{\mathrm{m}} \frac{\left[\mathrm{Wb}_{i} \right]_{\mathrm{o}}}{\left(1 + \frac{\mathrm{K}_{\mathrm{ai}}}{\left[\mathrm{H}^{+} \right]_{\mathrm{obs}}} \right)} \right\} - \frac{2 \eta \mathrm{P}_{\mathrm{H2}}}{\mathrm{V}_{\mathrm{R}}} \Delta t \tag{9}$$

where m is the number of acid-base equilibria other than acetate, carbonate, and phosphate. Rearranging Equation (9) and at the limit when $\Delta t \rightarrow 0$:

$$\frac{\partial}{\partial t} \left\{ \frac{\left[\text{Phos} \right]_{o}}{\left(1 + \frac{K_{a2}}{\left[\text{H}^{+}\right]_{obs}} \right)} + \sum_{i=1}^{m} \frac{\left[\text{Wb}_{i} \right]_{o}}{\left(1 + \frac{K_{ai}}{\left[\text{H}^{+}\right]_{obs}} \right)} \right\} = \left\{ \frac{\left[\text{Phos} \right]_{o} K_{a2}}{\left(K_{a2} + \left[\text{H}^{+}\right]_{obs} \right)^{2}} + \sum_{i=1}^{m} \frac{\left[\text{Wb}_{i} \right]_{o} K_{ai}}{\left(K_{ai} + \left[\text{H}^{+}\right]_{obs} \right)^{2}} \right\} \frac{\partial \left[\text{H}^{+} \right]_{obs}}{\partial t} \tag{10}$$

)

$$\frac{\partial [H^+]_{obs}}{\partial t} = \left(\frac{1}{4}K_{AcOH}^{1/2} - 2\eta\right)\frac{P_{H2}}{V_R} - \left\{\frac{[Phos]_o K_{a2}}{\left(K_{a2} + [H^+]_{obs}\right)^2} + \sum_{i=1}^m \frac{[Wb_i]_o K_{ai}}{\left(K_{ai} + [H^+]_{obs}\right)^2}\right\}\frac{\partial [H^+]_{obs}}{\partial t}$$
(11)

$$\left\{1 + \frac{[\text{Phos}]_{o}K_{a2}}{\left(K_{a2} + [\text{H}^{+}]_{obs}\right)^{2}} + \sum_{i=1}^{m} \frac{[\text{Wb}_{i}]_{o}K_{ai}}{\left(K_{ai} + [\text{H}^{+}]_{obs}\right)^{2}}\right\} \cdot \frac{\partial [\text{H}^{+}]_{obs}}{\partial t} = \left(\frac{1}{4}K_{AcOH}^{1/2} - 2\eta\right)\frac{P_{H2}}{V_{R}}$$
(12)

Numerically integrating Equation (12) over time in the observed pH range, where the major acid-base equilibrium is the one of phosphate, the following expression is approximately valid:

$$\sum_{\lambda}^{t_{f}} \left\{ 1 + \frac{[\text{Phos}]_{o} K_{a2}}{\left(K_{a2} + [\text{H}^{+}]_{obs}\right)^{2}} \right\} \cdot \frac{\Delta [\text{H}^{+}]_{obs}}{\Delta t} \Delta t = \int_{\lambda}^{t_{f}} \left(\frac{1}{4} K_{\text{AcOH}}^{1/2} - 2\eta \right) \frac{P_{\text{H2}}}{V_{\text{R}}} dt = \frac{\left(\frac{1}{4} K_{\text{AcOH}}^{1/2} - 2\eta \right)}{V_{\text{R}}} H$$
(13)

where λ is the lag phase time and *H* is the molar cumulative hydrogen production; the left-hand side of Equation (13) indicates the numerical integration. Aimed at facilitating notation, the function f_H is defined and indicates the numerical integral when $\Delta t \rightarrow 0$.

$$f_{H} = \sum_{\lambda}^{t_{f}} \left\{ 1 + \frac{\left[\text{Phos} \right]_{o} K_{a2}}{\left(K_{a2} + \left[H^{+} \right]_{obs} \right)^{2}} \right\} \cdot \frac{\Delta \left[H^{+} \right]_{obs}}{\Delta t} \Delta t \xrightarrow{\Delta t \to 0} \left\{ 1 + \frac{\left[\text{Phos} \right]_{o}}{\left(1 + K_{a2} / \left[H^{+} \right]_{obs} \right)} \right\} \left[H^{+} \right]_{obs}$$
(14)

 f_H can be obtained entirely from the bulk pH time series. Since the pH sampling period used is exceedingly small, compared with the characteristic time of the dark fermentation, the numerical cumulative sum can approximate the integral, thus, simplifying the calculation of the f_H function. The value obtained for f_H is close to the proton concentration, $|H^+|_{obs'}$ since the correction factor is close to 1.

Statistical analysis of the experimental results and the developed models were implemented in MATLAB R2021b (The MathWorks Inc., Natick, MA, USA).

2.4. Lag Time Calculation

The lag time can be estimated directly from the pH curve. It is observed that during an initial period, the pH presents a constant value, corresponding to the lag phase of the microorganism culture. Then, the intersection between a forward extrapolation of the initial pH with the back extrapolation of a function describing the pH decay would be a robust estimator of the lag phase time, λ . The culture activity starts with an abrupt change in pH towards acidic conditions until several buffer stages are reached, by a mixture of species and saturated carboxylic acids, dominated by the acetate (pKa = 4.75) from the media. To find the lag phase time, we make an exponential adjustment (Figure 2) of the pH curve between 7.2 and 5.4. The time obtained from the exponential curve fit intersection with the pH at the lag phase is an unbiased measure of lag phase time. This direct adjustment can give an accurate estimation directly from experimental information without the need of assuming any kinetic model.



Figure 2. Typical decay of pH and exponential fit of the pH time series during the onset of fermentative activity. Estimation of the lag phase time for the experiment shown in Figure 1.

2.5. A Measure of Hydrogen Production Efficiency

Taking the results shown in Figure 1 and representing the measured cumulated moles of produced hydrogen in the same figure, superimposed with the time evolution of the function f_H , described in Equation (14), a correlation between the two variables is evidenced (Figure 3). Even if the scales do not match, the trends are remarkably similar, indicating a strong relation despite the scale factor. Therefore, we propose to level scales through an estimated efficiency obtained by fitting Equation (13).



Figure 3. *f_H* function (left) and cumulated hydrogen production (right).

Figure 4 illustrates the goodness of fit of a linear model relating the f_H function and the cumulated moles of produced hydrogen. Hence, we postulate that the coefficient η of Equation (13) obtained by comparing the experimental hydrogen production to the synchronic values of the f_H function would provide a reasonable estimate of the process relative efficiency. Knowing the effective volume of the reactor, 5000 mL, and converting the hydrogen production from volumetric to molar, η can be obtained (Figure 4) from the linear fit parameters, forcing the ordinate at the origin to cross zero, as expressed in Equation (13).



Figure 4. Linear fit of f_H function vs. cumulative hydrogen production for the experiment shown in Figure 1.

Table 1 lists the calculated efficiencies determined from the proposed model and statistical figures that describe the goodness of fit. Additionally, values of the lag time, estimated directly from the pH curve, as described in Section 2.4, are indicated for each experiment. The lag time is generally between 13 and 16 h, except in Batch V, which also led to the lowest efficiency and the worst fit. Hence, a mean value for the explored conditions would be 17 ± 7 h. The estimated efficiency is quite low, around 0.04%, indicating many metabolic routes consuming protons, apart from the one leading to hydrogen gas formation. Nevertheless, it should be kept in mind that the scale factor is given by the acetic acid dissociation constant as a lumped factor. Although the value of this factor modifies the calculated efficiency, it provides a quantitative mean to compare results obtained with different media. The estimated values for the efficiency coefficient are also quite regular, always between 0.029% and 0.043%, leading to a mean value of 0.037% \pm 0.006% for the conditions explored in this work.

Batch	λ (h)	Slope (L ⁻¹)	r ²	η (%)
Ι	13	$4.8 imes10^{-5}$	0.9452	0.041
II	13	$7.7 imes10^{-5}$	0.9553	0.033
III	13	$3.8 imes10^{-5}$	0.8934	0.043
IV	16	$5.0 imes10^{-5}$	0.9788	0.040
V	29	$9.4 imes 10^{-5}$	0.8728	0.029
Mean	17	$6.1 imes 10^{-5}$	0.9291	0.037
Std. dev.	7	$2.3 imes10^{-5}$	0.0443	0.006

Table 1. Summary of the lag time estimated directly from the pH curve, and of the efficiency determined from the linear fit of f_H vs. H plots.

2.6. Kinetic Parameters—Modified Gompertz Model

The biological production of H_2 has been largely modeled according to the modified Gompertz model [30–32], since it encompasses the bacterial cell growth coupled with a parsimonious parametrization of the hydrogen production with more direct physical meaning [33,34]. The modified Gompertz model has the expression given in Equation (15), where *H* is the cumulated hydrogen produced, which can be given in moles, liters, or moles per reactor volume [35]. Parameter A has the same units and indicates the maximum

amount of hydrogen that can be produced, interpreted as the affinity towards hydrogen production. R_m is the maximum hydrogen production rate, and λ is the lag phase time.

$$H = \mathbf{A} \cdot \mathbf{e}^{-\mathbf{e}^{\left[\frac{\mathbf{R}_{m,\mathbf{e}}}{\mathbf{A}}(\lambda-\mathbf{t})+1\right]}}$$
(15)

The fitting can be further simplified from three to two parameters if the lag phase time is estimated directly from the pH curve with the procedure described in Section 2.4.

Given the approximately linear relation between the f_H function and H highlighted in Section 2.5, it is worth exploring the possibility of estimating the modified Gompertz model parameters from the pH curve. Equation (13) relates the cumulated hydrogen production to the function (f_H), which is entirely obtained from the pH curve. The parameters can be obtained from a nonlinear fitting or, since the modified Gompertz model is a bi-exponential function, an exponential fitting over a semi-log plot of the f_H function vs. time, starting after the lag phase, can be obtained, as indicated in Equation (16).

$$\ln(f_H) = \ln\left(\frac{\left(\frac{3}{2} - 2\eta\right)H}{V_r}\right) = -e^{\left[\frac{R_m \cdot e}{A}(\lambda - t) + 1\right]} + \ln\left(\frac{\left(\frac{3}{2} - 2\eta\right)A}{V_r}\right)$$
(16)

$$\ln(f_H) = -e^{[\alpha(-t)+1]} + \beta \tag{17}$$

$$\alpha = \frac{R_{m}.e}{A} \tag{18}$$

$$\beta = \ln\left(\frac{A}{V_{R}}\left[\frac{3}{2} - 2\eta\right]\right) \tag{19}$$

For comparison, the experimental cumulated hydrogen production was also fitted to the modified Gompertz model, taking the logarithm of Equation (15):

$$\ln(H) = -e^{\left[\frac{K_{m} \cdot e}{A}(-t) + 1\right]} + \log(A)$$
(20)

Tables 2 and 3 list the parameters fitted from all the experiments under the conditions explored, which were defined based on a previous piece of work [36]. Table 2 reports the values when the parameters A, Rm, and λ are fitted, and Table 3 provides A and R_m obtained while setting the lag phase time as the value obtained from the pH curve with the procedure described before.

Table 2. Summary of modified Gompertz model parameters.

Batch	fit	A (mol)	R _m (mol/h)	λ (h)	r ²
Ι	$_{f_H}^{ m H}$	$\begin{array}{c} 0.85 \pm 0.39 \\ 0.84 \pm 0.01 \end{array}$	$\begin{array}{c} 0.011 \pm 0.003 \\ 0.018 \pm 0.001 \end{array}$	$\begin{array}{c} 15.2\pm6.1\\ 26.7\pm0.5\end{array}$	0.9748 0.9782
II	${ m H} f_{H}$	$\begin{array}{c} 0.44 \pm 0.07 \\ 0.52 \pm 0.01 \end{array}$	$\begin{array}{c} 0.011 \pm 0.003 \\ 0.011 \pm 0.001 \end{array}$	$\begin{array}{c} 12.9\pm5.3\\ 26.7\pm0.5\end{array}$	0.9938 0.9781
III	${ m H} f_{H}$	$\begin{array}{c} 2.39 \pm 1.02 \\ 0.98 \pm 0.01 \end{array}$	$\begin{array}{c} 0.017 \pm 0.002 \\ 0.016 \pm 0.001 \end{array}$	$\begin{array}{c} 30.0\pm0.1\\ 19.2\pm0.4\end{array}$	0.9771 0.9886
IV	${ m H} f_{H}$	$\begin{array}{c} 0.99 \pm 0.03 \\ 1.05 \pm 0.001 \end{array}$	$\begin{array}{c} 0.010 \pm 0.001 \\ 0.011 \pm 0.001 \end{array}$	$\begin{array}{c} 12.2 \pm 2.5 \\ 22.2 \pm 0.3 \end{array}$	0.9986 0.9953
V	${ m H} f_{H}$	$\begin{array}{c} 0.87 \pm 0.19 \\ 1.23 \pm 0.01 \end{array}$	$\begin{array}{c} 0.014 \pm 0.008 \\ 0.008 \pm 0.001 \end{array}$	$\begin{array}{c} 20.8 \pm 16.7 \\ 23.2 \pm 1.0 \end{array}$	0.9713 0.9685

Figures 5 and 6 illustrate the goodness of fit of the results of two experiments, the best and the worst fit. The parameters determined both from the experimentally measured moles of hydrogen (Figures 5a,c and 6a,c) and from the moles calculated through the pH curve with Equation (13) ((Figures 5b,d and 6b,d) are coincident. Hence, it is inferred that the pH curve can be successfully used to get kinetic information on hydrogen production

without measuring the gas evolved if a reasonable estimation of the efficiency coefficient is available.

Batch	Fit	A (mol)	R _m (mol/h)	λ (h)	r ²
Ι	Н	0.85 ± 0.40	0.010 ± 0.002	13	0.9712
	f_H	0.98 ± 0.04	0.012 ± 0.001	13	0.9474
	Н	0.44 ± 0.02	0.011 ± 0.001	13	0.9997
11	f_H	0.61 ± 0.02	0.007 ± 0.001	13	0.9473
III	Н	1.97 ± 1.04	0.015 ± 0.008	13	0.9376
	f_H	1.05 ± 0.02	0.014 ± 0.001	13	0.9800
IV	Н	0.98 ± 0.04	0.011 ± 0.001	16	0.9973
	f_H	1.07 ± 0.001	0.009 ± 0.001	16	0.9922
V	Н	0.85 ± 0.20	0.018 ± 0.005	29	0.9518
	f_H	1.20 ± 0.01	0.009 ± 0.001	29	0.9469

Table 3. Summary of modified Gompertz model parameters, setting the lag time.



Figure 5. Comparison of the measured moles of H_2 in Batch IV (**a**,**c**) and the moles of hydrogen estimated from the corresponding pH curve using the function f_H and the relation given by Equation (13) (**b**,**d**) vs. the values predicted by the modified Gompertz model with three fitted parameters (**a**,**b**) or two fitted parameters and the lag phase time estimated from the pH curve, as described before (**c**,**d**).

Figure 5 correspond to Batch IV, the experimental results shown in Figures 1 and 3, and for which the estimated lag phase time method has been shown in Figure 2. This batch experiment, carried out for 180 h led to more than 20 L, around 1 mole, of hydrogen until the generation stopped, due to sucrose consumption. The Gompertz model successfully describes the hydrogen production with three fitted parameters (Figure 5a) and restricting

the lag phase time to the one obtained with the procedure given in Section 2.4 (Figure 5c). The figure also illustrates the moles of hydrogen inferred from the pH curve through Equation (13), using the fitted efficiency coefficient given in Table 1 (blue points with 5 min sampling period in Figure 5b,d). Since the method successfully estimates the produced hydrogen, the parameters of the Gompertz model are close to those obtained from the measured gas.



Figure 6. Comparison of the measured moles of H_2 in batch V (**a**,**c**) and the moles of hydrogen estimated from the corresponding pH curve using the function f_H and the relation given by Equation (13) (**b**,**d**) vs. the values predicted by the modified Gompertz model with three fitted parameters (**a**,**b**) or two fitted parameters and the lag phase time estimated from the pH curve, as described before (**c**,**d**).

Figure 6 correspond to Batch V, which was also carried out for 180 h, and again led to approximately 20 L, around 0.8 moles, of hydrogen until the generation stopped. The conditions and the media were the same as for Batch IV, but the experimental hydrogen gas was measured less frequently. Nevertheless, the Gompertz model satisfactorily describes the hydrogen production both with three fitted parameters (Figure 6a) and restricting the lag phase time to the one obtained with the procedure given in Section 2.4 (Figure 6c). The pH curve in this experiment crossed through a period of buffered pH, which is reflected in the estimated hydrogen production (blue points with 5 min sampling period in Figure 6b,d). This variation led to a worse fit and the parameters are still in the same order. It is worth-while mentioning that buffering periods were also reflected in less hydrogen generation, supporting the link between pH and hydrogen gas production.

Figure 7 illustrates the goodness of fit of the model calculated with the results shown in Figure 1, using both the parameters fitted directly from the volume of hydrogen produced, and estimated with the hydron ion mass balance. Although the fitting is better if the experimentally measured moles of hydrogen are used, the estimation arising from the

moles of hydrogen calculated through the mass balance from the monitored pH is also good. There is a high correlation between the experimental H values with those predicted with the modified Gompertz model fit over the f_H function.



Figure 7. Comparison of experimentally measured moles of H_2 vs. the values predicted by the modified Gompertz model using parameters fitted from the experiments and the pH curve using the function f_H and the relation given by Equation (13).

Figure 8 illustrates the plot of estimated vs. measured hydrogen obtained using the fitted parameters given in Table 3. The result indicates that the modified Gompertz model provides a good description of the hydrogen production rate. Comparing the parity plots calculated with parameters fitted from experiments or from the pH curve, the mean absolute error increases from 7% to 12%. It is also worth mentioning that the modified Gompertz parameters can also be obtained from the initial period of the f_H function. Hence, it is likely that the pH values in the initial active stages of the dark fermentation could be used to predict the hydrogen productivity of the culture using the mass balance.



Figure 8. Cont.



Figure 8. Comparison of experimental moles of H₂ vs. those predicted by the modified Gompertz model using parameters fitted from the experiments and the pH curve.

Table 4 compares the modified Gompertz model parameters reported in the literature for hydrogen production by dark fermentation from different sources. Results obtained in this work are in the order of those achieved using other sources. Mu et al. [19] working with a mixed culture of anaerobic bacteria obtained from a citrate-producing wastewater achieved similar values of A and higher values of Rm. The sludge was heated at 102 °C for 90 min to eliminate methanogenic hydrogen trophic bacteria, concentrating the spore forming bacteria responsible for hydrogen production. The work was also carried out in a 5 L batch stirred tank but regulating the pH in 5.5. More recently, Turhal et al. [24] found values in the order of the ones obtained in this work, if they are referred to the reactor working volume, using a heat-treated anaerobic sludge fed with a mixture of melon and watermelon with different solid concentration. Additionally, Del Angel Acosta et al. [30] obtained parameters in the same order if they are referred to the reactor working volume was produced with a heat-treated sludge. The authors used corn and brewery wastewater starting with different initial pH values (from 5.8 to 12.3) according to the mixture or setting the initial pH to 6.

Table 4. Summary of Gompertz model parameters estimated from different works aimed at obtaining hydrogen from effluents and solid wastes.

Conditions							Gompertz Parameters *		
Inoculum	Substrate	V _{R.} (L)	Т (°С)	pН	Agitation	A (mmol H ₂ /L Reactor)	R _m (mmol H ₂ /L.h)	λ (h)	
Preheated anaerobic sludge from citrate producing wastewater	Sucrose	5	30–45	Regulated at 5.5	120 rpm	168 ± 7	15 ± 1	9.8 ± 0.3	[30]
Heat-shocked digester sludge	Microcrystalline Cellulose	- 0.12	37	7.0	0	1.63 ± 0.07	0.29 ± 0.02	4.6 ± 0.3	- [30,37,38] - **
<i>Clostridium</i> -rich sludge from a pig manure digester	Rice organic waste			(initial)	1.5 rpm	30 ± 2	0.17 ± 0.01	54 ± 2	
	Potato organic waste	-			-	25 ± 3	0.55 ± 0.06	110 ± 10	
Compost-based and low-pH inocula	Glucose and sucrose solutions	0.15–0.175	22–30	7–8.5 (initial)	Unstirred vs. 160 rpm	49–120	2.7–13.7	18–60	[39]

Conditions						Gompertz Parameters *			Ref.		
Inoculum	Substrate	V _{R.} (L)	Т (°С)	рН	Agitation	A (mmol H ₂ /L Reactor)	R _m (mmol H ₂ /L.h)	λ (h)			
Sea sediment clostridia	Mineral salt—glucose	0.04	30	3.25–4 (final)	- None -	17	0.030	15.9	[33]		
	Ferrihydrite amended					20	0.033	15.2			
connitanty	Ferrihydrite addition					21	0.033	17.9			
	Melon and watermelon fruit mixture with 0.74 to 37 g/L total Solid concentration, with		36	5.5–6			37 gTS/L				
Heat-treated anaerobic sludge		0.08				218 (wi) 176 (woi)	15.7 (wi) 3.6 (woi)	3.65 (wi) 27.3 (woi)	- - [24] -		
							0.74 gTS/L				
	(wi) and without inoculation (woi)					16 (wi) 6.7 (woi)	0.5 (wi) 0.19 (woi)	4.1 (wi) 76.4 (woi)			
	Mineral salt—glucose	0.1	37	N/D		32.4	0.49 ± 0.05	4	[40]		
	Fe ₃ O ₄ addition					40.7	0.93 ± 0.20	3.8			
DSM 10,702	Lactate addition					38.0	0.72 ± 0.04	4.8			
	Lactate + Fe ₃ O ₄ addition					51.4	0.76 ± 0.13	3.6			
Anaerobic sludge blanket	Brewery waste and corn waste mixtures	0.08	35	Variable or 6 (initial)	150 rpm	106–167	0.6–2.1	1.5–122	[41]		
Ship wastewater Clostridia, selected by thermotolerant evolution	Minanal activity and the			7–8.5 (initial) 4–4.5 (final)	50 rpm -	204 ± 68 ***	2.6 ± 0.6 ***				
	media	5	30			196 ± 44 ****	2.2 ± 0.6 ****	13–29	This work		

 Table 4. Cont.

* expressed divided by the reactor working volume ** data from references [26,27] analyzed in reference [30]; *** from H, **** from the pH curve.

3. Materials and Methods

3.1. Bacterial Sludge

The bacterial source for this work has been a sample of residual water from a water treatment chamber on board an Argentine Navy ship, the frigate "Libertad". The sludge is collected directly from the chamber into a 5 L jerrycan and stored at room temperature. To seed a 5 L rector, 500 mL of this sludge is collected in an Erlenmeyer flask and heated at 75 °C for 50 min to rid the sludge of non-spore-forming species. After this pre-treatment, it is ready to pour into the reactor tank. The performance of this bacterial biomass from the water treatment plant of a ship has been compared to other sources in previous work [12], showing outstanding hydrogen production. Genetic analysis of 16S rRNA on the untreated sludge has shown that it is composed of over 25 different bacterial families, but this is notoriously reduced by the temperature pretreatment. After such pretreatment, it is mostly composed of *Clostridium* sp., which are well known hydrogen producers [14,21].

3.2. Culture Medium and Fermentation Process

The 500 mL of pre-treated sludge is added to a stirred reactor where a 4.5 L culture medium has been prepared. This culture medium is a variant of the one proposed by Logan et al. [42]. This medium contains distilled water, ammonium bicarbonate (NH₄HCO₃) in the amount of 2 g/L, potassium phosphate (KH₂PO₄) at 1.2 g/L, and several other nutrients and micronutrients in the amount of 0.2 g/L, which are magnesium sulfate (MgSO₄), manganese sulfate (MnSO₄), sodium molybdate (Na₂MoO₄), calcium chloride (CaCl₂), and ferric chloride (FeCl₃). The carbon source is sucrose in a 25 g/L concentration.

Clostridium species ferment the carbohydrate while producing carbon dioxide (CO_2) and H_2 gas released into the reactor's headspace. They also produce a mix of acids, mostly acetic and butyric acids, that are released into the culture medium. As a result, acidity increases while H_2 is being produced. At a certain point, low pH affects bacterial metabolism, and, to avoid this effect, a weak base is initially added to the culture medium.

In this case, sodium acetate (NaC₂H₃O₂) in 0.05 M concentration has been chosen (6.8 g/L of the trihydrate salt) as neutralizing agent.

Once the culture medium and the inoculum were introduced into the reactor vessel, the cover was screwed in place, the stirring was set at 50 rpm, and an anaerobic atmosphere was generated by bubbling nitrogen (N₂) for 5 min. The temperature was adjusted with a heating jacket at 37 °C. A pH sensor was attached to the lid registering with a sample period of five minutes. The reactor was a Sartorius Biostat A plus in batch mode.

After the N_2 had bubbled out, an acrylic container made of three equal cylinders, connected in parallel, was attached to the vent to capture all gases produced in the reactor. The cylinders were full of water. As the gas was produced, the water was removed from the container, passing on to a water reserve. The volume of removed water was the gas volume produced by the bacterial biomass. The gas volume was calculated by measuring the cylinder volume. Each cm of the three cylinders container was equal to 0.72 L (each cylinder height was 20 cm, radius 8.7 cm). The starting time of the reactor was recorded as the point when the gas container was attached.

3.3. Hydrogen Measure and Production

As the fermentation starts, the bacterial spores detect optimal conditions and start budding, ending the lag phase. Once grown, the biomass starts metabolizing sucrose, and cell replication occur. In this phase, pH shows a steep decline while gases accumulate. The gas stored in the container corresponds to all gases produced by the biomass, which means a mix of H₂ and CO₂. This gas mixture has been evaluated by chromatography and has shown to be composed of only these two gas species, with a 60–70% proportion of H₂. The authors developed a new quantitative method to estimate the proportions of these gases in previous work [16]. It has been shown that using a syringe with a known volume of gas mix and consuming the H₂ in a highly selective proton-exchange membrane (PEM) fuel cell leaves only the volume of CO₂, which can be directly read with the syringe. A simple volume balance results in the amount of H₂ present in the sample. This method is simple, cost-effective, and straightforward.

4. Conclusions

Valuable information on hydrogen production efficiency and kinetics can be obtained by analyzing the pH time evolution monitored during dark fermentation by spore-forming bacteria in a batch stirred tank reactor. On the one hand, the pH curve can provide a direct estimation of the lag phase time without assuming any kinetic model. Since the exponential period of culture growth produces short chain acids, it leads to a sharp pH decay. Hence, from the intersection of the initial constant pH value with an exponential fit describing the pH decay, a robust lag phase time estimator can be obtained.

Furthermore, from a model based on the mass balances of the hydron ion, describing the link between the hydrogen produced by dark fermentation and the evolution of pH during the process, a measure of the hydrogen efficiency is proposed. The model is developed, solved, and successfully compared with experiments.

The agreement between the experimentally measured moles of hydrogen and the values estimated from the hydron ion balance indicates that it is a reactor performance descriptor worth being explored. Moreover, it can be used to estimate parameters of models, describing the hydrogen generation kinetics, such as the widely applied modified Gompertz model. The pH values in the initial active stages of the dark fermentation could be used to predict the hydrogen productivity of the culture. There is a high correlation between the experimental *H* values with those predicted with the modified Gompertz model fit over the f_H function. A fair similarity arose in the affinity factor, A, and the estimated R_m parameter predicted independently from both results. Therefore, it can be concluded that hydron-related kinetics is strongly linked to molecular hydrogen production, and the monitored pH when no pH regulation is imposed allows estimating relevant information of the process.

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Nomenclature

Symbol	Name	Units
aη	Ratio of H and f_H according to Equation (13)	L^{-1}
α	Rate parameter of $\ln(f_H)$ vs. t non-linear fit	h^{-1}
А	Affinity towards hydrogen production	Mol
β	Independent parameter of $\ln(f_H)$ vs. t non-linear fit	-
e	Base of natural logarithm	-
f_H	Cumulated hydron conversion function	mol/L
Η	Cumulated hydrogen production	Mol
$[H^+]$	Hydron concentration	mol/L
К _а	Acid dissociation equilibrium constant	
n	Number of moles	Mol
n _i	Electric charge of the i-th base anion	-
r	Correlation coefficient	-
R _m	Maximum hydrogen production rate	mol/h
pН	Negative base 10 logarithm of hydron concentration	-
P _{H2}	Hydrogen production	mol/h
[Phos] _o	Analytic phosphate concentration	mol/L
t	Time	Н
VR	Reactor working volume	L
Wb	Weak base	-
η	Efficiency towards hydrogen production	-
λ	Lag phase time	Н

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