Article

Fermentation of Wild Hydrolytic Bacteria Isolated from *Opuntia ficus-indica*, and Its Effect on Flow Behavior of Cladodes Flour Culture Media

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ABSTRACT: The aim of this study was to evaluate the fermentation dynamics of two wild hydrolytic microorganisms and their effect on the flow behavior of a culture medium formulated with *Opuntia ficus-indica* cladode flour. Identified *Acinetobacter pitti* and *Bacillus subtilis* presented maximum values of specific hydrolytic activity (SHA) at 24 h of growth (0.21 ± 0.05 and 0.22 ± 0.01 IU, respectively). The apparent viscosity of cladode flour medium (CFM) measured by applying shear rates (66.7 s⁻¹–0.003 s⁻¹) in suspensions (20%) showed a significant decrease (60%) as a function of bacterial growth progressed. After fermentation, the CFM exhibited pseudoplastic (shear-thinning) behavior, which was linked to the enzymatic degradation of polysaccharides. The use of crude extracellular enzyme extracts from these wild bacteria effectively reduced medium viscosity by breaking down the plant matrix. These findings highlight the hydrolytic potential of native strains in modifying the rheological properties of cactus-based culture media, offering a low-cost alternative for biomass pretreatment and valorization in future biotechnological applications.

Keywords: Opuntia ficus-indica; Acinetobacter pittii; Kluyveromyces marxianus; Cladodes; Flow behavior



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1. Introduction

Opuntia ficus-indica is a native plant from Mexico, characterized by its oval, flattened cladodes—modified stems that perform photosynthesis and are commonly mistaken for leaves [1]. In Mexico, twenty-five states, obtain an annual production of 837,267 tons of cladodes [2], of which 70% is sold (fresh or processed) to the food and pharmaceutical industry as brines, juices, jellies, among others [3]. The remaining 30% represents a critical excess [4] which is typically discarded by producers, leading to a large quantity of daily generated waste. Due to its abundance, low cost, and renewable nature, cladode biomass represents a promising alternative substrate for microbial cultivation, enzymatic processes, and other biotechnological applications [5,6]. Mexican producers have recognized the importance of developing strategies to utilize this surplus, not only to reduce waste but also to generate added value from underused resources [7,8].

Cladodes contain complex polysaccharides such as cellulose (21.6%), hemicellulose (21%), lignin (3.6%) [9] and mucilage (33%) [10]. Cellulose and hemicellulose represent an energy-rich carbon source due to glucose molecules (C₆H₁₂O₆) bound by β -glucosidic bonds. The mucilage is a heteropolysaccharide with high water activity (>0.8) [11] that contains simple sugars (rhamnose, xylose, arabinose, glucose and galactose) bound to galacturonic acid by β -glucosidic bonds [10,12], all values expressed on a dry matter basis. The presence of this glycosidics bonds makes cladode flour (CF) susceptible to the hydrolysis process. Moreover, there is an interest in using purified and/or crude enzymes as a good hydrolyzing agent for avoiding the use of corrosive acids in this critical stage, considered as a hydrocolloid with industrial potential. The mucilage is a non-gelling agent with elastic properties similar to those of polymers such as xanthan gum and carboxymethylcellulose (CMC), with a high ability to increase the viscosity of

aqueous systems [7,13,14]. There are reports of extraction methodologies, chemical composition, functional, physicochemical and rheological properties of mucilage [7,8,10,14–18]. However, cladodes have a low total solids content (\sim 4–7% on fresh weight), which may be a limitation in fermentation processes that require high biomass input. Additionally, lignin content increases with the age of the cladode, which is why young cladodes are preferred for consumption and processing [10]. Additionally, despite the extensive characterization of mucilage, there are no reports of changes in hydrolytic activity and flow behavior due to the growth of wild hydrolytic bacteria.

The aim of this study was to evaluate the fermentation dynamics of two wild hydrolytic microorganisms and their effect on the flow behavior of a culture medium formulated with *Opuntia ficus-indica* cladode flour.

2. Materials and Methods

2.1. Vegetal Material

Opuntia ficus-indica Atlixco variety cladodes of 12 months of age were collected in an experimental field. The fresh cladodes were cut into pieces (1 cm²) using an industrial vegetable cutter (3/8 inch blade, Hobart, FP-350, Troy, OH, USA) and dried at 80 °C/24 h in a steam trays dryer (Jersa, FT-DCV-02, Guadalajara, Mexico). Subsequently, dry cladodes were ground (Pulvex, 200, Naucalpan, Mexico) to obtain a flour with a particle size of 1500 μ m, and stored in sealed plastic bags (25 °C for 1 week) for culture medium preparation and further analysis.

2.2. Cladode Medium Preparation

A suspension (250 mL) was prepared by adding cladodes flour (20%) to mineral solution (MS). The MS was prepared by the addition of distilled water (225 mL) to a mixture of mineral salts (50:50, 25 mL) consisting of mineral solution I (0.6% K₂HPO₄) and mineral solution II (1.2% NaCl, 1.2% (NH₄)₂SO₄, 0.6% KH₂PO₄, 0.12% CaCl₂ and 0.25% MgSO₄·7H₂O) [19]. All ingredients were homogenized mechanically and adjusted to pH 6.5 using a potentiometer (Oakton, pH 11 series, Singapore, Singapore). The mixture was sterilized at 121 °C/15 min.

2.3. Proximate Composition Analysis of Cladode Flour

Ash, titratable acidity, moisture and pH of cladode flour were analyzed according to AOAC 923.03, 942.15, 984.25 and 981.12, respectively [20,21]. Total carbohydrates [22], free reducing sugars [23], and minerals (Ca, K, Na, P, Mg) [24] were analyzed. Swelling capacity (SC) and water holding capacity (WHC) were determined following the method of [25], with minor modifications. Briefly, 1 g of cladode flour was mixed with 10 mL of distilled water in a graduated cylinder and allowed to stand at room temperature for 24 h. SC was expressed as the volume increase (mL water/g DM). For WHC, the hydrated sample was centrifuged at 3000 rpm for 20 min, and the retained water was measured gravimetrically (g water/g DM). For color measurements, a colorimeter (HunterLab, 4500L, Reston, VA, USA) was used with 13 g of sample considering L*, a* and b* values.

2.4. Isolation and Molecular Identification of Wild Hydrolytic Bacteria

2.4.1. Isolation

A homogenous slurry was set of decayed cladodes; an aliquot was then prepared to seed microorganisms using solid medium added with 1% CMC as a selective medium at 37 °C/24 h. The first visualized colonies were separated and re-seeded in solid media added with CMC and CFM (1%, pH 6.5).

2.4.2. Morphological Characterization and Qualitative Hydrolytic Activity Determination

The shape, size, color and texture of the obtained colonies were visualized using a stereoscope (10×, Leica, EZ4D, Singapore). The shape of the cells was determined using a microscope (100×, Olympus, CX31RBSFA, Tokyo, Japan). Congo tests were performed to ensure hydrolytic activity by observation of halos degradation according to the Teather and Wood protocol [26]. Gram tests were performed according to Bartholomew and Finkelstein (1958) [27] protocol.

2.4.3. Molecular Identification

Molecular identification of hydrolytic microorganisms was performed by analysis of the 16S ribosomal gene sequence (Thermo Fisher, Waltham, MA, USA). The genomic DNA used as a template was obtained from 5 mL of a cell culture growth in nutrient broth (Sigma, St. Lois, MO, USA) at 37 °C/24 h. For genomic DNA extraction, a CTAB

2% buffer was used [28]; subsequently, a purification with the Gen EluteTM PCR Clean-Up kit protocol (Sigma) was performed. The obtained fragments were amplified using oligonucleotides 16S (5'AGA.GTT.TGA.TCC.TGG.CTC3') and 16SR (5'CGG.GAA.CGT.ATT.CAC.CG3'). The amplified fragments were cloned into pGEM-T vector system I (Promega, Madison, WI, USA) for recombinant plasmids construction. Transformation with JM109 competent cells (Promega) was completed, and then, a purification of plasmidic DNA with GeneluteTM Plasmid Miniprep kit protocol. The pure plasmidic DNA was quantified using a Nanodrop (2000, Waltham, MA, USA), adjusted to 150 ng/ μ L and sent to sequencing. The sequencing was carried out with Macrogen Inc. (Seoul, Republic of Korea). The results of the sequences (in Fasta format) were edited excluding the vector using Geneious 5.4.8 software [29] and the resulting sequences were used to perform a BLASTn from NBCI [30] and identified with the generic and/or specific name according to the percentage of similarity to the reported in a database.

2.5. Growth of Wild Hydrolytic Bacteria in Cladode Flour Medium

Two bacteria were grown at 37 °C/6 h with agitation at 200 rpm starting from 1 mL of glycerol in 9 mL of 1% CFM as the first pass. A second growth pass was performed with the resulting 10 mL mixed with 50 mL of 20% CFM for 12 h. Finally, an aliquot was taken for the third pass, starting with a cell population of 20×10^6 cell/mL (250 mL of medium). The growth was performed for 168 h, sampling every 24 h during growth.

2.6. Protein Determination in Cladode Flour Fermentation Using Bradford Assay

Soluble protein concentration was determined throughout the fermentation process using the Bradford method [31], adapted for samples derived from cladode flour fermentation. At each sampling point (0, 6, 12, 24, 48, 72, 120, and 168 h), 5 mL of culture broth was collected and centrifuged at 10,000 rpm for 10 min at 4 °C. The supernatant was recovered and used for protein quantification.

Briefly, $100 \ \mu\text{L}$ of sample supernatant was mixed with 1 mL of Bradford reagent (Coomassie Brilliant Blue G-250, $0.01\% \ w/v$ in 4.7% ethanol and 8.5% phosphoric acid) in a test tube. After 5 min of incubation at room temperature, the absorbance was measured at 595 nm using a UV-Vis spectrophotometer (JENWAY[®] model 6715, Jenway, Cambridge, UK). A standard curve was constructed using bovine serum albumin (BSA) solutions ranging from 0 to 25 mg/mL, and the soluble protein concentration in the samples was expressed in mg/mL.

2.7. Specific Hydrolytic Activity Determination

The specific extracellular hydrolytic activity of wild bacteria was determined by correlating free reducing sugars and soluble protein, and expressed in International Units according to Equation (1):

$$IU = sugar equivalent/(protein*time)$$
(1)

where, International Units = amount of enzyme capable of releasing 1 μ mol of sugar per minute under the condition tested (IU), Sugar equivalent = free reducing sugars determined (μ mol), Protein = soluble protein determined (mg), Time = reaction time (min).

The enzyme preparation used for viscosity reduction was a crude extracellular enzyme extract, obtained directly from the culture supernatant of wild bacteria grown in CFM, without any purification steps. Hereafter, this preparation is referred to as "crude enzyme extract" throughout the text.

It is important to note that the protein values used to calculate specific activity correspond to the total extracellular protein content in the supernatant, including both enzymatic and non-enzymatic fractions. As bacterial growth increases, total protein rises due to the secretion of non-specific proteins, which may dilute the calculated IU/mg values. Therefore, specific activity is interpreted as a relative indicator of hydrolytic efficiency rather than an absolute measure of enzyme concentration.

2.8. Rheological Analysis

The viscosity of the cladode flour medium (CFM) obtained during the growth of wild bacteria was determined using a digital viscometer (SC4-16 needle, RVDV-III U; Brokfield Engineering Laboratory Inc., Middleboro, MA, USA) at 25 °C. The behavior was determined by recording the shear stress values (τ , Pa) needed to deform the samples with increasing (0.003 to 66.7 s⁻¹) shear rate values (γ , s⁻¹) and in descending order to obtain both the upward and downward curves, respectively. Data from both curves were adjusted to the Herschel-Bulkley equation (Equation (2)) characterizing the flow behavior of CFM degraded by wild bacteria.

$$\tau=\tau_0+K\gamma^n$$

where:

 $\tau_0 =$ yield stress (Pa)

n = flow behavior index (dimensionless)

K = consistency coefficient (Pa)

The parameters n and K were obtained from the log $(\tau - \tau_0)$ versus log (g) diagram. The yield stress was calculated by Casson equation $(\tau^{0.5} = \tau_0^{0.5} + \eta \gamma^{0.5})$.

2.9. Experimental Design

A factorial design 4×2 was established, evaluating the following factors (Table 1). The response variables evaluated were yield stress (τ_0), consistency coefficient (K), flow index (n), apparent viscosity (η), bacterial growth (cel/mL), total, specific hydrolytic activity (IU), and free release sugars (g/L).

Table 1. Factorial design 4×2 of the effect of fermentation time and type of microorganism on the physicochemical characteristics of a medium formulated with cladode flour.

Exp	Factors		Response Variables						
	Time (h)	Strain	τ ₀ (Pa)	K (cP)	n	η (Pa·s)	Bg	SHA	FRS
1	0	A. pitti	δ_1	β_1	α_1	Σ_1	O1	R_1	V_1
2	24	A. pitti	δ_2	β_2	α_2	Σ_2	O ₂	R_2	V_2
3	96	A. pitti	δ_3	β3	α3	Σ_3	O ₃	R_3	V_3
4	168	A. pitti	δ_4	β_4	α_4	Σ_4	O_4	R_4	V_4
5	0	B. subtilis	δ_5	β_5	α_5	Σ_5	O_5	R_5	V_5
6	24	B. subtilis	δ_6	β_6	α_6	Σ_6	O_6	R_6	V_6
7	96	B. subtilis	δ_7	β ₇	α_7	Σ_7	O ₇	\mathbf{R}_7	V_7
8	168	B. subtilis	δ_8	β_8	α_8	Σ_8	O_8	R ₈	V_8

Note: τ_0 = Yield stress; K = Consistency coefficient; n = Flow index; η = Apparent viscosity; Bg = Bacterial growth (cel/mL); SHA = Specific hydrolytic activity (IU); FRS = Free release sugars (g/L); h = Hours.

2.10. Statistical Analysis

The effect of the two parameters tested, time (0 h, 24 h, 96 h, 168 h) of fermentation, strain (*A. pitti, B. subtilis*), and their interaction, was statistically analyzed in terms of the rheological behavior. All response variables were examined with statistical software (Minitab v. 15, Inc., State College, PA, USA) with a 5% test of significance.

3. Results

3.1. Characterization of Cladode Flour

The physicochemical composition of cladode flour (CF) is presented in Table 2.

Ashes were similar (24.04%) to those reported for other authors [7,12] for cladode flours from other varieties (16–20%), probably because the cladodes of this work were part of a mineralized experimental lot where fertilization source was an inorganic fertilizer that provided nitrogen, phosphorus and potassium, based on urea, triple superphosphate and potassium chloride [32]. CF presented a higher content of Ca^{2+} , Mg^{2+} and K^+ compared with other reports [12,17,33–35] while the concentration of Na^{1+} and P^{3+} was slightly. Several researches have shown the importance of calcium and its compounds (calcium oxalate and calcium carbonate) present in the cell walls of *Opuntia ficus-indica* cladodes, because they are fundamental compounds that form the mucilage network in young cladodes (less than a year) [36].

Table 2. Proximate composition and physical characteristics of cladodes flour (dry matter) from *Opuntia ficus-indica* var. Atlixco (12 months)^a.

Parameters	Value
Ash (g/100 g)	24.04 ± 0.34
Moisture (g/100 g)	6.13 ± 0.49
pH	5.06 ± 0.01
Titratable acidity (g/100 g)	0.22 ± 0.07
Total carbohydrates (g/100 g)	18.15 ± 0.10

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(2)

Free reducing sugars (g/100 g)		4.15 ± 0.00
Swelling (mL water/g DM ^b)		10.00 ± 3.00
Water holding capacity (g water/g DM ^b)		6.80 ± 0.40
Color	L*	68.09 ± 0.35
(dimensionless)	a*	-1.82 ± 0.04
(differisioness)	b*	23.99 ± 0.22
	Ca	4.78 ± 0.01
Minerale	Mg	0.98 ± 0.01
$(\alpha/100 \alpha)$	K	3.36 ± 0.01
(g/100 g)	Na	0.006 ± 0.001
	Р	0.064 ± 0.01

Note: ^a Average of two replicates; ^b Dry Matter.

A moisture of 6.13% was detected, this value is desirable due to microbial growth and hygroscopicity is prevented (lower than 10%), improving the shelf life of the flour. Atlixco variety exhibited a high pH value (>5) compared with other varieties (*i.e., Opuntia ficus-indica amylocea* and *Opuntia ficus-indica inermis*, pH ~ 3), explained by the age of cladodes and the concentration of many organic acids, such as malic, citric and oxalic acids [25,33]. Related to hydration properties, the CF swelling power obtained (10.00 \pm 3 mL/g) was higher, while the value of water retention capacity (WRC) was similar (6.80 \pm 0.40 g/g) compared with the data obtained by Ayadi et al. (2009) [25] (7.36 \pm 0.07 mL/g and 6.85 \pm 0.03 g/g, respectively). The CF presented a better ability to bind to water molecules. Color parameters showed flour with high luminosity (L* = 68.09 \pm 0.35) and a yellow-green color (b* = 23.99 \pm 0.22, a* = -1.82 \pm 0.04). The color parameters were similar to those reported by Sáenz (1997; 2010) [12,37] (L* = 73.37, b* = 26.1 and a* = -5.20) and Ayadi et al. (2009) [25] (L* = 67.49, b* = 25.15 and a* = -8.17).

In addition, although the values of total carbohydrates and free reducing sugars represent a fraction of the flour $(18.15 \pm 0.10 \text{ g} \cdot 100 \text{ g}^{-1} \text{ and } 4.15 \pm 0.00 \text{ g} \cdot 100 \text{ g}^{-1}$, respectively), the remaining percentage up to 100% dry basis is mainly attributed to structural carbohydrates such as cellulose, hemicellulose, lignin, and pectins. This estimation is a common practice in biomass studies [7,38]. In fact, several authors have reported that *Opuntia ficus-indica* cladodes contain approximately 9–14% cellulose, 20–25% hemicellulose, and 6–14% lignin, in addition to mucilage and pectins as key components of their cell-wall architecture [39,40]. These reports support structural carbohydrate content in *Opuntia ficus-indica* cladodes flour (~60%).

3.2. Identification of Wild Hydrolytic Bacteria

3.2.1. Isolation

Producing hydrolytic enzyme strains (four) were obtained from decayed cladodes. All strains grew well at 37 °C on carboxymethylcellulose (CMC) and cladode flour medium (CFM) under aerobic conditions during the first 24 h. Microbial hydrolytic enzymes production is influenced by factors including inducer substrate, type of strain used and reaction conditions (temperature, pH, *etc.*).

3.2.2. Morphological Characterization and Qualitative Determination of Hydrolytic Activity

The microorganisms were identified as bacteria by microscopic observation (Table 3) and presented two morphologies: cocci and bacilli. An identification code was assigned to each of them: B400, B401, B402 and B403. The colonies presented circular forms (B400 and B402) and irregular forms (B401 and B403) with white, orange and yellow coloration. All bacteria had Gram (–) staining and only B400 and B401 presented hydrolytic activity, as confirmed by degradation halos in solid medium with 1% of CF. Two hydrolytic bacteria were chosen for molecular identification; specific hydrolytic activity quantification was made during growth, as well as the rheology in cladode flour medium (CFM).

C	Microscopic Form	Macroscopic Form	Colony Color	Gram Test	Congo Red _ Test ^a	Image		
Code						Microscopic (100×)	Macroscopic (10×)	
B400	Cocci	Circular and convex	Creamy and White	_	+			
B401	Coccobacillus	Irregular and flat	Creamy	_	+		0.000	
B402	Streptococci	Circular and convex	Orange	_	_			
B403	Cocci	Circular and convex	Yellow	_	_			

Table 3. Description of colonies and bacterial cells isolated from decayed cladodes.

Note: ^a Hydrolytic activity in solid cladode flour medium (1%).

3.2.3. Molecular Identification

When performing the identity search for each sequence obtained, the bacterium B400 presented 100% identity with *Acinetobacter pittii*, whereas the bacterium B401 presented 99% identity with *Bacillus subtilis*. Some studies have reported the identification of new bacteria identified as *Pseudomonas*, *Staphylococcus*, *Acinetobacter* and *Bacillus* with cellulolytic activity isolated from gut termite [41], sawdust [42] and from *Opuntia ficus-indica* [43]. The isolated bacteria from this work correspond to the genera already reported to have hydrolytic activities, specifically cellulolytic.

3.3. Growth and Specific Hydrolytic Activity Determination of Wild Bacteria

Acinetobacter pittii and Bacillus subtilis showed satisfactory growth in CFM, although they exhibited an adaptation phase of at least 24 and 48 h, respectively (Figure 1).



Figure 1. Growth curves for *A. pittii* (black square) and *B. subtilis* (gray square). Specific hydrolytic activity (IU) for *A. pittii* (black bar) and *B. subtilis* (grey bar). Free reducing sugars for *A. pittii* (black diamond) and *B. subtilis* (grey diamond). Average of two replicated. Specific activity ($IU \cdot mg^{-1}$) was calculated based on total extracellular protein. The observed decrease reflects enzyme dilution due to non-enzymatic protein accumulation, not a loss of hydrolytic capacity.

A two-phase trend in free reducing sugars was observed. In the first 24–48 h, an increase in reducing sugar concentration—particularly in *A. pittii*—coincided with the peak of extracellular hydrolytic activity, reaching values of 0.22 IU·mg⁻¹ for *B. subtilis* and 0.21 IU·mg⁻¹ for *A. pittii*. While our study did not include identification of the specific monosaccharides released or direct enzyme characterization, the hydrolytic behavior observed is consistent with the general degradation of polysaccharides in the matrix. Previous studies have reported the ability of *Bacillus subtilis* and *Acinetobacter* species to produce extracellular hydrolytic enzymes such as cellulases, pectinases, and xylanases under similar conditions [44–46]. Additionally, *Opuntia ficus-indica* cladodes have been shown to respond effectively to enzymatic mixtures containing cellulases and xylanases, facilitating pectin extraction and matrix disruption [47]. After this time, the consumption of all released sugars and the decrease of hydrolytic activity were observed, coinciding with the increase in bacterial growth. *Acinetobacter pittii* exhibited exponential growth, reaching a maximum of 4.68 ± 0.66 × 10⁹ cell/mL after 144 h, while *Bacillus subtilis* reached a final population of 3.92 ± 0.20 × 10⁹ cell/mL.

3.4. Rheological Analysis

All samples showed pseudoplastic behavior, a characteristic of compounds with high molecular weights such as mucilage $(3.3 \times 10^4 \text{ kDa})$. In general, CFM with *A. pittii* and *B. subtilis* growth (Figure 2a and Figure 2b, respectively) showed shear thinning behavior due to deformation. The viscosity curves decreased with an increasing shear rate $(0.003-66.7 \text{ s}^{-1})$. The slight decrease in viscosity was reached when $\gamma \ge 29 \text{ s}^{-1}$ (Figure 2).



Figure 2. Steady shear viscosity measurement of cladode flour medium at different times (0 h, solid circle), (6 h, solid square), (12 h, solid diamond), (24 h, solid triangle), (48 h, empty circle), (96h, empty square), (120 h, empty diamond), (144 h, empty triangle), (168 h, solid cross) for (**a**) *A. pittii* and (**b**) *B. subtilis* growth. Average of two replicates.

In addition, the diagrams showed a progressive decrease in the viscosity of the medium as a function of the growth time of wild bacteria, related to the release of hydrolytic enzymes that degrade the network of the raw polymeric material. When bacteria begin to grow using monosaccharides available in the medium, they excrete potent enzymes (exo-glucanases, endo-glucanases and β -glucosidades) that break down complex polysaccharides, causing a decrease in viscosity (Figure 3) and releasing more monosaccharides into the medium, which are consumed by microorganisms to grow exponentially. In spite the increase in the concentration of soluble protein (0.20–2.28 and 0.27–1.50 mg/mL for *A. pittii* and *B. subtilis*, respectively) as an indicative of the production of enzymes, the activity decreased because the specific hydrolytic activity is subjected to the released amount of sugars into the medium, which were consumed by the bacteria over time.

After 12 h, an increase in the viscosity was observed for medium, most notably for *B. subtilis*, which was not observed for *A. pitti* (Figure 3), presumably due to the change of pH of the medium or to the presence of divalent ions like calcium.



Figure 3. Apparent viscosity ($\gamma = 29 \text{ s}^{-1}$) for *A. pittii* (black bar) and *B. subtilis* (grey bar). Average of two replicate.

The viscosity of CFM obtained at 6 h of growth for *A. pittii* and *B. subtilis* was 1.6 ± 0.05 and 1.4 ± 0.01 Pa·s, respectively. The progressive decrease in viscosity during bacterial growth strongly suggests the secretion of hydrolytic enzymes into the medium. Although the crude enzyme extract—defined as the non-purified culture supernatant—was preliminarily tested on fresh CFM, this observation was qualitative and not quantitatively evaluated. Therefore, it is not presented as experimental evidence in this study. The viscosity reduction trends observed during fermentation remain the primary support for the enzymatic activity. The flow curves were well-fitted ($r^2 > 0.90$) by Herschel-Bulkley equation (Figure 4).





Figure 4. Flow curve of cladode flour medium (20%) for (**a**) *A. pittii* and (**b**) *B. bacillus* growth at different times (0 h, upper black circles; 24 h, black circles; 96 h, dark grey circles; and 168 h, light grey circles) with. Experimental curves: upward (increasing shear rate), close symbols, and downward (decreasing shear rate), open symbols. Modelling curves (upward: continuous line, downward: dashed line). Average of two replicates.

The ascending and descending curves of *A. pittii* and *B. subtilis* in CFM obtained during growth time (0, 24, 96 and 168 h) showed that yield stress (τ_0), viscosity (h) and consistency coefficient (K) decreased as a function of time, whereas the flow behavior index values (n) were lower when $\tau \rightarrow 0$, and increased with growth time (Table 4).

Strain	Time (h)	τ 0 (Pa)	K (cP)	n (Dimensionless)	η ^b (Pa·s)	r ²
	0	261 ± 23	49 ± 16	0.20 ± 0.13	1.42 ± 0.20	0.934
Control 6	24	256 ± 16	59 ± 12	0.22 ± 0.01	2.15 ± 0.05	0.955
Control	96	362 ± 98	50 ± 6	0.22 ± 0.03	3.32 ± 0.01	0.972
	168	248 ± 64	44 ± 1	0.24 ± 0.01	2.25 ± 0.01	0.927
	0	352 ± 151	67 ± 45	0.35 ± 0.03	2.09 ± 0.70	0.965
1	24	269 ± 32	37 ± 8	0.56 ± 0.03	1.80 ± 0.12	0.955
A. piiiii	96	175 ± 25	62 ± 9	0.53 ± 0.02	1.30 ± 0.59	0.937
	168	70 ± 17	33 ± 18	0.44 ± 0.06	0.73 ± 0.11	0.956
	0	280 ± 16	45 ± 6	0.54 ± 0.02	1.90 ± 0.05	0.94
D aubtilia	24	298 ± 46	31 ± 11	0.52 ± 0.09	1.39 ± 0.15	0.067
D. SUDIIIIS	96	149 ± 31	46 ± 14	0.47 ± 0.02	1.30 ± 0.35	0.962

Table 4. Rheological parameters (τ_0 , K and η) of upward curves ^a of cladode flour medium (20%) added with hydrolytic bacteria as a function of growth time.

Note: ^a Adjusted to Herschel-Bulkley model, average of two replicates; ^b Apparent viscosity at 29 s⁻¹; ^c Without bacteria growth.

 0.48 ± 0.01

 0.81 ± 0.04

 35 ± 1

 76 ± 10

168

Flow index values obtained for *A. pittii* and *B. subtilis* (0.56 ± 0.03 and 0.54 ± 0.02 , respectively) were higher compared to the control (culture medium without microorganisms) (0.22 ± 0.01) indicating a higher fluidity and a loss of the pseudoplasticity of CFM and therefore, a degradation of the raw material by the hydrolytic bacteria. Statistical analysis indicated that there were significant differences in flow index due to strain, time and the interaction of both factors (Table 5).

Table 5. Probability of individual and interaction effects on studied factors (strain and time) on the rheological parameters of cladode flour medium.

Eastans	<i>p</i> -Values of Independent Variables							
Factors	τ ₀ (Pa)	K (cP)	η (Dimensionless)	H ^a (cP)				
		Individual factor	rs					
A: Strain	<0.001 ^b	0.090	<0.001 ^b	<0.001 ^b				
B: Time	me <0.001 ^b <0.001 ^b		<0.001 ^b	<0.001 ^b				
Interaction								
AxB	<0.001 ^b	0.316	<0.001 ^b	<0.001 ^b				

Note: ^a Apparent viscosity at 29 s⁻¹; ^b Significant effect.

Both model curves (upward and downward) showed that CFM with *A. pittii* and *B. subtilis* exhibited a close to zero thixotropic area despite its high-water absorption capacity due to the hydrocolloid nature (30 g/100 g of mucilage).

4. Discussion

The fermentation of *Opuntia ficus-indica* cladode flour medium (CFM) by native hydrolytic bacteria led to significant modifications in the physicochemical properties of this system, particularly in the reduction of viscosity and the development of a pseudoplastic flow profile. Consistent with our findings, recent studies have reported that enzymatic hydrolysis during fermentation significantly decreases the viscosity of the medium. For instance, Song et al. (2022) [48] observed very low viscosities (~14.5 cP) in *Opuntia* humifusa extracts fermented with cellulolytic enzymes in contrast to considerably higher viscosities of the unfermented controls. This effect reflects the well-established pseudoplastic nature of *Opuntia* mucilage: a viscous biopolymer that exhibits shear-thinning behavior, with a flow index n < 1 that becomes more pronounced at higher concentrations. The enzymatic activity of *Bacillus subtilis* and *Acinetobacter pittii*, which reached maximum specific hydrolytic activities (SHA) of 0.22 ± 0.01 IU and 0.21 ± 0.05 IU, respectively, at 24 h, revealed a pivotal role in the hydrolysis of polysaccharides present in the medium. In particular, *Bacillus subtilis* is a robust cellulolytic bacterium; Malik and Javed (2021) [49] reported that *B. subtilis* strains can produce high levels of cellulases (e.g., CMCase ≈ 2.4 IU/mL) capable of degrading lignocellulose. Similarly, *Acinetobacter pittii*, isolated from decomposing cladodes, exhibited strong cellulolytic activity on cactus tissues. In a previous study, López-Domínguez et al. (2019) [50] reported up to 0.67 IU/mL of total cellulase activity (FPase) and 0.23 IU/mL of endoglucanase activity within 24 h when using *A. pittii* in a 20% cladode suspension.

0.958

The reduction in viscosity (~60%) observed after fermentation of the cactus flour suspension may be attributed to hydrolytic activity targeting the fiber polysaccharides. While this aligns with reports on microbial degradation of mucilage, our study specifically confirmed enzymatic activity using carboxymethyl cellulose (CMC) as a model substrate.

Although both bacterial strains exhibited similar SHA values, the progression of enzymatic activity over time revealed differences in metabolic dynamics, with *B. subtilis* achieving higher early-stage activity, which may be advantageous for short-cycle bioprocesses. The sustained activity observed up to 48 h suggests a stable production of hydrolytic enzymes, compatible with longer fermentation periods. The enzyme activity data correlate with the 60% reduction in viscosity of the 20% cladode flour suspension, supporting the hypothesis that enzymatic breakdown of fiber and mucilage leads to structural disintegration of the medium. In line with previous studies where *Bacillus subtilis* strains exhibited both amylolytic and cellulolytic activity during the microbial saccharification of plant-based residues [51], the present work highlights the applicability of native strains for enzymatic degradation and biomass valorization of complex substrates.

The composition of *Opuntia* cladodes, rich in mucilage, insoluble fiber, and calcium-bound polysaccharides, influences the substrate's susceptibility to hydrolysis. The acidity and pH of cladodes are partially determined by the presence of organic acids generated via crassulacean acid metabolism (CAM), an adaptation of cacti to arid environments that also affects the fermentability of the substrate [33]. Recent studies indicate that water retention in fiber-rich flours is governed mainly by branched, highly polar polysaccharides (e.g., pectins, hemicelluloses) rather than by cellulose or lignin [38,52]. These non-cellulosic fibers possess abundant hydroxyl and carboxyl groups, conferring much higher water-binding capacity than the crystalline cellulose or relatively hydrophobic lignin [52]. In *Opuntia* cladode flour, the native mucilage—a high-molecular-weight heteropolysaccharide rich in galacturonic acid and neutral sugars—chelates Ca²⁺ to form extensive cross-linked, gel-like networks that trap water and impart high viscosity [47,53]. When microbial cell wall hydrolases (e.g., cellulases, pectinases, hemicellulases) are applied, these polysaccharide matrices are cleaved and the previously bound water is released, leading to a dramatic loss of viscosity and altered rheological behavior [5]. In the present study, although only endoglucanase activity was directly evaluated using CMC as a substrate, a comparable decrease in viscosity was observed, suggesting that similar hydrolytic mechanisms may be involved. However, the presence of pectinolytic or hemicellulolytic activity remains a plausible hypothesis that warrants further investigation.

From a rheological standpoint, CFM displayed marked pseudoplastic behavior after bacterial fermentation, characterized by a decrease in the flow behavior index (n < 1). This non-Newtonian response is consistent with previous studies using enzymatic hydrolysis on plant-based suspensions. Kim et al. (2013) [54], for instance, reported a 70% reduction in viscosity using Viscozyme on 20% cladode suspensions, confirming the effectiveness of multi-enzyme cocktails in degrading mucilage and cellulose. In this study, a comparable decrease in viscosity was achieved using crude supernatants from wild bacteria, suggesting their potential for fiber degradation. Although the enzymatic spectrum of the bacterial extracts was not fully characterized, the reduction was consistent with microbial action on polysaccharides.

The enzymatic performance observed, mainly reflected by viscosity loss and endoglucanase activity, suggests that pectin degradation might also occur, although no direct pectinolytic assays were performed. This hypothesis is supported by prior studies on microbial action over *Opuntia* mucilage [5,6]. This process involves the cleavage of galacturonic acid-rich domains within the mucilage matrix, facilitating breakdown and access to underlying polysaccharides. Furthermore, the comparatively lower enzymatic activity detected in *Acinetobacter pittii* relative to *A. anitratus* [55] may be attributed to the complex architecture of the culture medium based on cladode flour (CFM), where mucilage layers can sterically hinder access to glycosidic bonds within cellulose microfibrils [56].

Finally, while the results are promising, variability in cladode composition depending on harvest season, maturity, and postharvest treatment could influence the reproducibility of enzymatic degradation. Future studies should address the standardization of the substrate and evaluate the scalability of bacterial fermentation to optimize hydrolytic enzyme production and polysaccharide breakdown. Nevertheless, this work demonstrates that *O. ficus-indica* cladodes, in combination with native hydrolytic bacteria, represent a viable and sustainable option for biomass valorization through enzymatic pretreatment in the context of green biotechnologies.

5. Conclusions

Bacteria isolated from decayed cladodes were identified as *Acinetobacter pittii* and *Bacillus subtilis*. Both had a maximum hydrolytic activity on a culture medium based on cladodes flour during the first 24 h, with no significant difference. All samples presented non-Newtonian behavior, and the upward and downward curves were well fitted to

Herschel-Bulkley equation. In the culture medium, the consistency (K) decreased and presented a loss of pseudoplasticity (n) with the addition of microorganisms and growth time, concluding that the raw material was degraded positively by the effect of extracellular enzymes. Statistically, it was shown that the interaction of both factors (type of strain and time) had a significant influence. Although specific enzymatic pathways were not characterized, the observed hydrolytic trends and sugar release suggest the action of extracellular enzymes. However, further studies are necessary to confirm the type and specificity of the enzymatic activities involved. Statistically, the interaction between strain type and incubation time had a significant influence on both flow behavior and sugar release. Considering the low total solids content of cladodes, future optimization strategies should explore pretreatment (e.g., drying or concentration) or co-substrate approaches to enhance fermentation efficiency and improve substrate availability in biotechnological applications.

Author Contributions

Conceptualization, M.O.R.-S., I.M.R.-B. and C.M.L.-D.; Methodology, I.M.R.-B. and C.M.L.-D.; Software, C.M.L.-D. and K.A.A.-B.; Validation, M.O.R.-S., I.M.R.-B. and C.M.L.-D.; Formal Analysis, M.O.R.-S. and I.M.R.-B; Investigation, I.M.R.-B. and K.A.A.-B.; Resources, M.O.R.-S.; Data Curation, C.M.L.-D.; Writing—Original Draft Preparation, M.O.R.-S. and C.M.L.-D.; Writing—Review & Editing, C.M.L.-D., K.A.A.-B. and I.M.R.-B.; Visualization, M.O.R.-S.; Supervision, M.O.R.-S., I.M.R.-B.; Project Administration, M.O.R.-S., I.M.R.-B.; Funding Acquisition, I.M.R.-B.

Ethics Statement

Not applicable.

Data Availability Statement

The datasets generated and analyzed during the current study are available from the corresponding author upon reasonable request.

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Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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