

Characterization by spectrometric methods of chitin produced from white shrimp shells of *Parapenaeus longirostris* by *Lactobacillus helveticus* cultivated on glucose or date waste

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ABSTRACT

Abstract: Chitin recovery by lactic acid fermentation was considered. It was previously shown that the use of glucose led to 61% of demineralization and 42% of deproteinization; while and unlike glucose, the inoculation of *L. helveticus* in date waste (200 g L⁻¹ reducing sugars) led to a simultaneous demineralization and deproteinization of shrimp shells, 83 and 61 %, respectively. The purity of end-products were characterized by infrared (FTIR), X-ray diffraction and X-ray fluorescence.

Analysis of the chemical composition of fermentation end-product by X-ray fluorescence showed that residual minerals are in trace except CaO and P₂O₅. The infrared spectra and the X-ray diffractograms of both samples compared to those of pure chitin showed a strong similarity. FTIR of fermented shells showed the appearance of characteristic peaks of α -chitin and the disappearance of calcite peaks. In addition and especially for the shells fermented using *L. helveticus* cultivated on date waste, a similarity with those of pure chitin should be noted. These results confirmed the efficiency of biological chitin recovery, and that a synthetic carbon source (glucose) can be replaced by a natural one, date waste, allowing a valorization of this latter. The relevance of subsequent works to improve yields of demineralization and deproteinization was therefore confirmed.

I. Introduction

Chitin is a major component in the supporting tissues of organisms such as crustaceans, fungi and insects. Next to cellulose, it is the most abundant biopolymer found in nature [1]. Chitin is structurally similar to cellulose [2,3] In the

Exoskeletons of crustaceans, chitin is intimately associated with Other Materials Such as proteins, lipids, calcium carbonate and pigments [3]. Natural, non toxic, biopolymer chitin is now widely produced commercially from crab and shrimp waste shells [4].

Because of their linear (1,4)- β -N-acetylglucosaminoglycan structure with two hydroxyl groups and an acetamide group (Figure 1), native chitin in crustacean shells are highly crystalline with strong hydrogen bonding, and are arranged as α -chitin microfibrils in an antiparallel fashion [5].

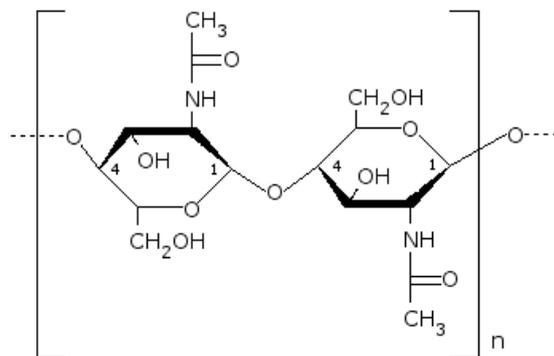


Figure 1. Chemical structure of chitin

X-ray diffraction studies reveal that chitin occurs in three polymorphic forms, α -, β -, and γ -chitins, which differ in the arrangement of molecular chains within the crystal cell [6, 7, 8]. α -chitin showing parallel chain packing is the most abundant chitin in nature (shrimps, crabs), β -chitin presents antiparallel chain and occurs in squid pens [9, 10, 11], while γ chitin presents a mixture of α and β chitins [7]. Because chitin possesses many beneficially biological properties such as biocompatibility, biodegradability, hemostatic activity, and wound healing property, much attention has been paid to its biomedical applications [12]. Its main derivative, chitosan, has been used as a dietary supplement due to its effective lipid binding and hypocholesterolemic properties and as a film-forming agent [13]. Chitin, chitosan and their derivatives can be used as antimicrobial, emulsifying, thickening and stabilizing agents in food industry [13].

Recently, biological processes for chitin production have been reported using organic acids producing bacteria [11] and enzymes [14] for the bio-demineralization and the bio-deproteinization of crustacean shells. Lactic acid fermentation of shrimp (*Penaeus monodon*) waste for chitin recovery was studied with added carbohydrate such as lactose or cassava extract as a natural energy source [15] and date waste for extraction of chitin from head and shell of shrimp *Parapenaeus longirostris* using *Lactobacillus helveticus* [16]. Raw heads of Africa river prawn (*Macrobrachium vollohovenii*) were fermented with *Lactobacillus plantarum* using cane molasses [17]. Lactic acid fermentation for demineralization has been also reported for shrimp waste [18],

crayfish exoskeleton [19], scampi waste [20] and prawn waste.

In this study, chitin was extracted from shrimp (*Parapenaeus longirostris*) shells by lactic acid fermentation using *Lactobacillus helveticus* cultivated on glucose or date waste. The biological extraction of chitin requires two main steps bio-deproteinization and bio-demineralization. The calcium and the protein in the shell waste were mainly dissolved by organic acids and proteases produced by microorganisms. The Efficiency of fermentation depends on factors such as the shell content [21].

The aim of this work was to study the efficiency of bio-demineralization and bio-deproteinization. The products were characterized by spectrometric methods (IR, XRD, X-ray fluorescence). Infrared spectroscopy (FTIR) has been proposed as the quickest and most effective way to compare the properties of chitin from different sources (animal and fungal). While the XRD and X-ray fluorescence are used to determine the crystal structure [22] and the quantity of components, respectively.

II. Materials and methods

II.1. Biological Material

This work was carried out using the biological material cited in table 1.

Crude shells, demineralized and deproteinized shells fermented on glucose and date's juice were harvested [16,23] (table 1) and analyzed in order to study the difference between end-products of each step. Analysis allowed comparing the appearance and disappearance of peaks characteristics of some elements such as protein and calcite.

II.2. Infrared spectra (IR)

To spectroscopically characterize the biopolymer contained in the samples, FTIR measurements were performed and the IR spectra were recorded on a Bruker IFS113vFTIR-spectrometer (Bruker AXS S.A.S., Champs-sur-Marne, France), in the 4000–500 cm^{-1} spectral region at a 2 cm^{-1} resolution. The samples were ground, mixed with KBr powder and then compressed into pellets.

II.3. X-ray diffraction (XRD)

X-ray diffraction measurements were applied to determine the crystallinity of the raw shell, chitin and the end-products. Their patterns were recorded using XRD analysis by means of a monochromatic Cu Ka Philips X'Pert Pro diffractometer (PANalytical, Limeil-Brévannes, France). The 2θ angle was scanned between 5° and 40° and the counting time was 15 s at each angle step. The voltage was 40 Kv and the intensity 50 mA. The intensities of the peaks at (110) lattice (I_{110} , at $2\theta \sim 20^\circ$ corresponding to the maximum intensity)

and at 2θ~16° (amorphous diffraction) were used to calculate the crystallinity indice I_{CR} following equation [24]:

$$I_{CR} = \frac{I_{110} - I_{am}}{I_{110}} \times 100 \quad (1)$$

Table 1. Characterization of the biological material used in this work

	Crude shell		Fermented shell			
	[16] 1	[23] 2	[16]	[16]	[23]	[23]
			DP ^a 80 g L ⁻¹ glucose 30°C	DM ^b 300 g L ⁻¹ glucose 35°C	DP and DM ^c 300 g L ⁻¹ glucose 30°C	DP and DM ^c 200 g L ⁻¹ reducing sugars 30°C
pH	8-9	8-9	76 % DP	20.6% DP	42.14% DP	83% DP
Humidity (%)	3.25	3.25	0% DM	53% DM	61% DM	61% DM
Ashs (%)	35.5	27				
Proteins (%)	27.61	12				
Chitin (%)	23.61	23				
Pigments and lipids (%)	13.28	35				

^aDeproteinized shell

^bDemineralized shell

^cSimultaneous demineralization and deproteinization

III. Results and Discussion

Characteristics of shells

The physico-chemical analysis of powder shells was based on the determination of protein content, minerals, chitin and water content. The obtained results are summarized in Table 1. It can be observed that for the same species, shell composition displayed variations. The first sample of crude shells (Table 1) shows a strong mineralization (35.5%) and significant protein content (27%); while the second crude sample had a low mineral level (27%) and a low protein content (12%). Infrared spectroscopy is a technique, which has the advantage of inexpensive method for identification of chemical groups such as (NH, CO) present on the surface of raw material. The spectra of the two shells are shown in figure 2.

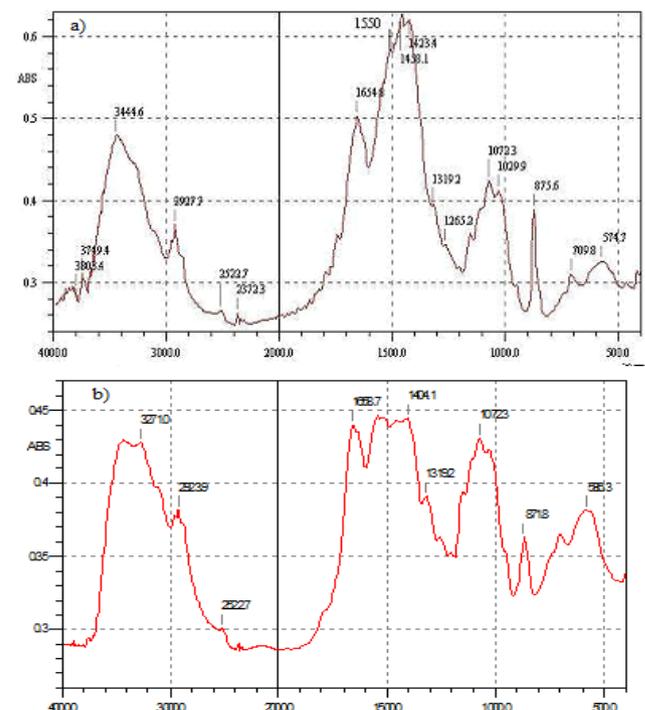


Figure 2. Infrared spectra of raw shell (sample 1 – Table 1) (a) and (sample 2 – Table 1) (b)

The infrared spectrum (a) shows that in addition of characteristics peaks of calcite (871 cm⁻¹, 1404cm⁻¹, 2522cm⁻¹), there are characteristic peaks of α-

chitin (1658cm^{-1} , 1319cm^{-1} , 1072cm^{-1} , 586cm^{-1}) (Fig. 2).

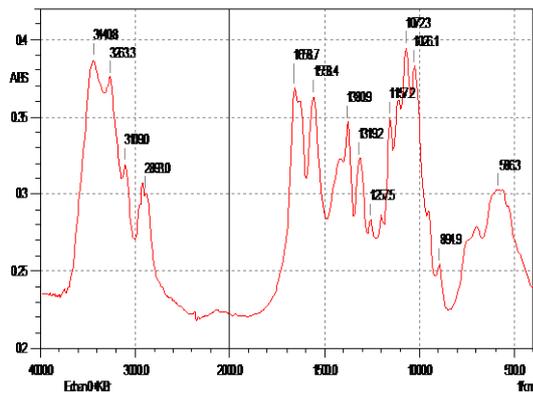


Figure 3. Infrared spectrum of pure chitin

While the spectrum of the first sample of raw shell shows the appearance of a peak characteristic of chitin (COC) stretching at 1072cm^{-1} and a strong band at 1654cm^{-1} , which corresponds to the (C = O) stretching, characteristic of amide I band and peak at 1550cm^{-1} corresponding to (NH) stretching characteristic of amide II. This difference may be attributed to the high rate of minerals and proteins present in the the first sample of raw shell.

Comparing with the spectrum of pure chitin (Fig.3), figure 1, shows the absence of calcite peaks and the presence of other intense absorption peaks at 1157cm^{-1} , 1380cm^{-1} , 1558cm^{-1} , 2893cm^{-1} , 3109cm^{-1} and 3263.3cm^{-1} which are characteristic of α -chitin [25, 26, 27, 28].

Analysis of end-products after lactic acid fermentation of the first sample of shrimp shells using *Lactobacillus helveticus* cultivated on 300g L^{-1} glucose at 35°C (53% demineralization) and on 80g L^{-1} glucose at 30°C (76% deproteinization) [16] gave infrared spectra shown in figure 4.

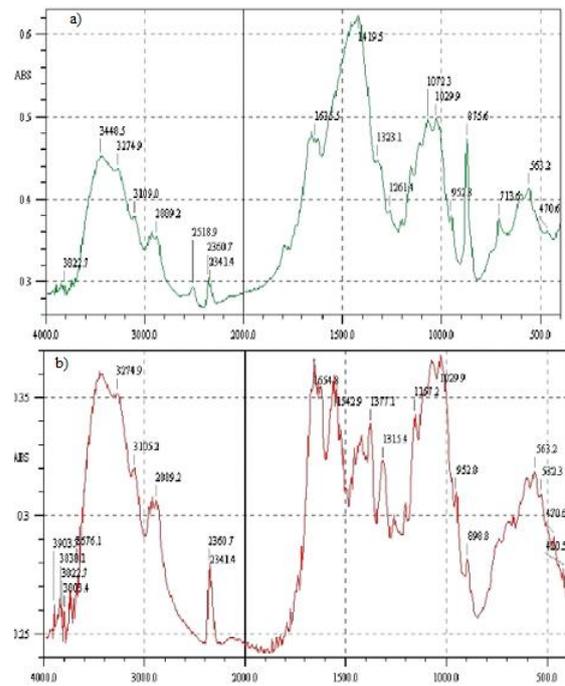


Figure 4. Infrared spectra of deproteinized (76% with 80g L^{-1} glucose at 30°C) (a) and demineralized (53% with 300g L^{-1} glucose at 35°C) (b) shrimp shells

The persistence of characteristic bands of calcite in the spectrum of deproteinized shells (Fig.4a) can be noted. New peaks at 2893cm^{-1} , 3109cm^{-1} which correspond to (CH) stretching also appeared, which are characteristic of α -chitin [29]. Contrarily, the spectrum displayed in Figure 4b shows a disappearance of bands characteristic of calcite, with appearance of intense characteristic bands of α -chitin at 2893cm^{-1} and 3109cm^{-1} (CH) which have been masked by calcite bands [29].

In order to improve the obtained results, another round of extraction of chitin by *Lactobacillus helveticus*, cultivated on glucose or date waste was carried out [23]. The physicochemical analysis of the raw shell showed lower protein content (Table 1). The use of glucose, with initial concentration of 300g L^{-1} at 35°C , as fermentation medium for chitin recovery led to 61 and 42% of demineralization and deproteinization, respectively; while, the use of date waste corresponding to 200g L^{-1} of reducing sugars at 30°C led to simultaneous efficient DM and DP, 61 and 83% [23]. The obtained products were characterized by infrared spectroscopy. The spectra obtained are shown in Figure 5.

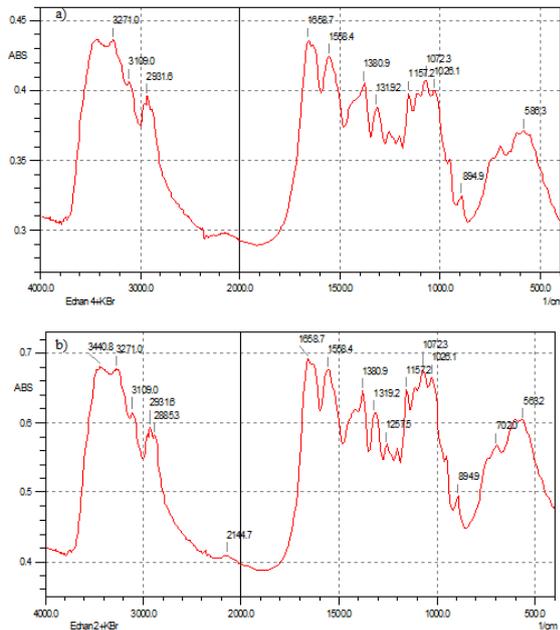


Figure 5. Infrared spectra of fermented shell (a) using *L. helveticus* growing on date waste (200g L⁻¹ reducing sugars at 30°C) (a) and on glucose (300g L⁻¹ at 35°C) (b).

Infrared spectra of the fermented shells with glucose (Fig.5a) and juice date (Fig.5b) show the disappearance of characteristic calcite peaks, and intense bands were found at 1658 and 1558 cm⁻¹. Both are typical for amides and corresponded to stretching vibrations of carbonyl groups, including in chitins (C=O usually called amide I) and to NH (amide II) respectively [29]. In addition, the spectra were lacking a signal at 1540 cm⁻¹, where proteins normally would give rise to absorption [3] and as well as the appearance of new peaks corresponding to α -chitin. The determination of the different absorption peaks and the different bands characteristics of chemical groups, observed in each sample, show a significant similarity between pure chitin and fermented shells (Table 2).

Table 2. The characteristic peaks of the samples [25, 26, 27, 28]

pure α -chitin	Fermented shell (with glucose 300g L ⁻¹ at 35 °C)	fermented shell (with juice date 200g L ⁻¹ reducing sugars at 30 °C)	crudeshell	The chemical groups
-	563	-	-	
586	-	586	586	
-	702	-	-	COC stretching
-	-	-	871	calcite
894	894	894	-	COC bridge (glucosidic linkage)
1026	1026	1026	-	COC stretching
1072	1072	1072	1072	COC stretching
1157	1157	1157	-	Glucose linkage / COC bridge [asymmetric (antisymmetric) stretching]
1257	1257	-	-	CH stretching
1319	1319	1319	1319	Amide III band
1380	1380	1380	-	CH stretching
-	-	-	1404	Very strong CO stretch CaCO ₃ [25]
1558	1558	1558	-	Amide II band
1658	1658	1658	1658	C = O-NH-CH ₃ stretching (amide I, α -chitin)
-	-	-	2522	Calcite
-	2144	-	-	CH = C + C elongationelongation (Bertrand, 2006)
-	-	-	-	CH stretching
-	2885	-	-	CH stretching
-	-	-	-	CH stretching

2893	-	-	-	Very strong CH stretching
-	-	-	2923	Amide Band II
-	2931	2931	-	Intermolecular C (2 ₁)NH ... O = C
3109	3109	3109	-	(7 ₃) hydrogen bonds
3263	-	-	-	NH stretching
-	3271	3217	3271	Intermolecular OH3 CH ₂ OH and
3440	3440	-	-	hydrogen bonds5)

From Table 2, it can be concluded that the characteristic peaks of α -chitin appeared close to those observed after fermentation using *L. helveticus* growing on date waste.

The diffraction pattern of crude shrimp shell (Fig.6), shows the presence of two intense peaks $2\theta = 19.39^\circ$ and $2\theta = 29.53^\circ$ which was typical of the calcium carbonate present in the carapace (5,30), as well as a low peak at $2\theta = 9.16^\circ$.

In the case of shrimp material, six signals were identified: at $2\theta = 9.2^\circ$, $2\theta = 12.55^\circ$, $2\theta = 19.15^\circ$, $2\theta = 20.8^\circ$, $2\theta = 22.95^\circ$ and $2\theta = 26.15^\circ$ [30].

Zhou et al. [26] showed that α -chitin has 2θ diffraction peaks at 9.3° , 19.4° and 26.3° . The work carried out by Stawski et al. [28] showed that for α -chitin, the maximum were located at $2\theta = 20.5^\circ$ and $2\theta = 40.5^\circ$.

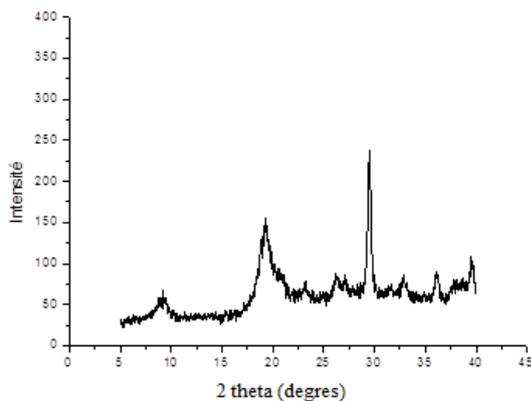


Figure 6. Spectrum of X-ray diffraction of raw shell

The characteristic peaks of pure chitin can be observed in both samples, namely obtained after shrimp shell treatment using *L. helveticus* growing on glucose (Fig.7a) or date waste (Fig.7b).

The interesting results obtained using the shell harvested in 2008 compared to those obtained using the shell harvested in 2005, pushed us to analyze by X-ray diffraction and X-ray fluorescence, only the end products of fermentation of the shell harvested in 2008.

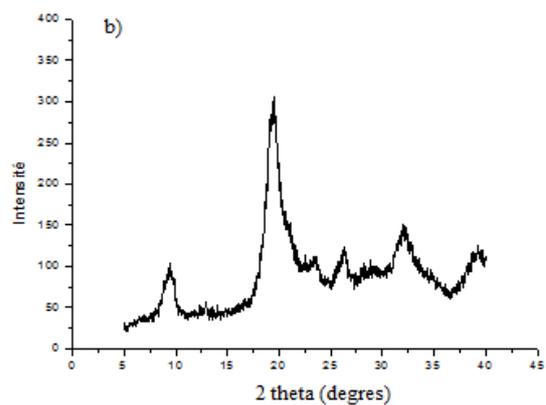
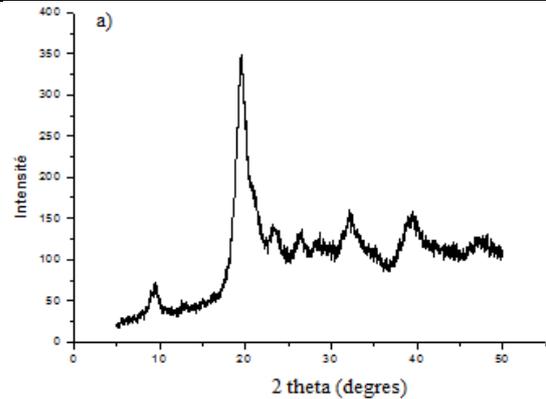


Figure 7. Spectrum of X-ray diffraction of the carapace treated by *L. helveticus* cultivated on glucose (a) or the date waste (b)

The X-ray diffractograms of both samples chitin (glucose and date waste) (Fig.7) were largely similar to α -chitin diffractograms described in the literature.

It was found that I_{CR} of the chitin from crab shells and rice-field crab shells (Table 3) were slightly higher than that of the chitin recovered in the present study. This difference may be attributed to the effect of interactions between chitin and impurities explained by the relatively low rates of demineralization and deproteinization.

The chemical composition of the raw shell and the two products previously recovered [23] was determined by X-ray fluorescence. The obtained results are collected in Table 4, and showed that the majority of minerals present in the raw shell were

Table 3. I_{CR} values of different samples of chitin

Sample	I_{CR} (%)	2θ (°)
Chitin (glucose)	77.15	9.30°, 19.45°, 23.35°, 26.42°, 32.33°, 39.41°
Chitin (date waste)	72.18	9.55°, 19.49°, 23.45°, 26.25°, 32.14°, 39.39°
α -chitin (from crab shells) [32]	85.0	
chitin from Cicada sloughs (insect) [24]	89.7	
Rice-field crab shells [24]		
Brown shrimp chitin (<i>Penaeus aztecus</i>) [33]	91.7	
Pink shrimp chitin (<i>Penaeus durarum</i>)	64	
	66.6	

removed, since they were at trace levels in the fermented shells except for calcium oxide CaO and phosphate P_2O_5 . This elimination is due to the production of lactic acid by lactobacilli. This production caused acidification and lowering of the pH to around 4.7 [23] which caused the

solubilization of the minerals initially present in in the raw shell and consequently demineralization. While the proteolytic enzymes secreted in the fermentative medium by the lactic acid bacterium ensure its deproteinization.

Table 4. Chemical composition of raw and fermented shells

Compounds	Raw shell	Fermented shell	
		on 200 g L ⁻¹ date juice at 30°C	on 300 g L ⁻¹ glucose at 35°C
Na ₂ O	1.899	0.004	0.018
MgO	1.739	0.014	0.022
Al ₂ O ₃	0.233	0.037	0.045
SiO ₂	2.202	0.04	0.171
P ₂ O ₅	5.323	5.96	5.42
CaO	21.636	9.014	8.9

IV. Conclusion

In order to check for the nature of the residual minerals, the elemental composition of the collected samples was analyzed. The X-ray fluorescence results showed that most elements were in trace. This analysis shows the elimination of 50% calcium oxide initially present in the carapace, which proved the efficiency of the biological treatment.

FTIR of fermented shells showed the appearance of characteristic peaks of α -chitin, and the disappearance of calcite peaks. In addition and especially for the shells fermented using *L. helveticus* cultivated on date waste, a similarity with those of pure chitin should be noted.

Finally and according to X-ray diffractograms of chitins obtained after biological treatment of shells, the appearance of three major characteristic peaks

of α -chitin should be noted, with however some impurities due to residual minerals.

The above results allowed confirming that a natural one, date waste, which also constituted a valorization of this waste, can replace a synthetic carbon source (glucose).

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