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# Ultrasonics Sonochemistry



journal homepage: www.elsevier.com/locate/ultson

# Effects of ultrasound-assisted chitosan grafted caffeic acid coating on the quality and microbial composition of pompano during ice storage

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ARTICLE INFO

Keywords: Ultrasound Chitosan grafted caffeic acid High-throughput sequencing technology Microbial composition

#### ABSTRACT

The effects of ultrasound-assisted chitosan grafted caffeic acid coating on the quality and microbial composition of fresh pompano (*Trachinotus ovatus*) fillets during ice storage for 24 days were evaluated. Samples were treated by distilled water (CK), ultrasound (US), chitosan grafted caffeic acid coating (G), and chitosan grafted caffeic acid coating with ultrasound-assisted (USG). Results showed that samples treated with USG could inhibit the formation of corrupt substances such as TVB-N, TBA, biogenic amines (BAs), hypoxanthine (Hx), and hypoxanthine riboside (HxR) when compared to the CK group. The results of high-throughput sequencing technology observed that the major bacteria genus of fresh samples was *Acinetobacter*. The diversity of bacterial communities at the initial stage was more diverse than that at the end of stage. With the extension of storage time, the USG treatment could maintain the microbial diversity. The dominant microbiota was *Shewanella* and *Brochothrix* in the CK group streated groups was effectively decreased. The microbial communities of samples in all treatments were changed during storage. At the end of storage, there was a significant difference in bacterial composition between the CK and treated samples, indicating that the treatment can effectively inhibit the growth of microorganisms, especially spoilage microorganisms, and reduce the quality deterioration caused by bacteria.

# 1. Introduction

Pompano (*Trachinotus ovatus*), belongs to the Carangidea family, which is one of the important commercial fish species in China, Japan, Australia and other countries [1]. It has the characteristics of delicious meat, excellent quality, fast growth, and the fact its fillet has few small bones. It is an important variety of aquaculture in China [2]. However, due to a series of microbial and endogenous biochemical reactions in fish after death, it is vulnerable to microbial corruption, which could easily lead to the deterioration of quality and shorten the shelf-life of fish [3]. The challenge of fillets during storage is how to delay and evaluate their quality change.

At present, bio-based edible coating technology is considered to be an effective and eco-friendly protection method for food preservation, and has been widely used [4]. Chitosan (CS) is an amino-polysaccharide that could be obtained by partial deacetylation of chitin [5]. Due to its natural non-toxic properties, biodegradable, bio-compatible and nontoxic, as a bio-based coating, chitosan has great application prospects in food industry [6]. However, CS coating could not be widely used because of its low antioxidant activity because there is no phenol-like functional group or conjugated structure in the molecule [7]. Caffeic acid (CA) is one of the most commonly used phenolic acids, which has been reported as a kind of phenolic acid with a high antioxidant capacity [8]. Recently, the combination of phenolic acid and CS effectively enhanced the antioxidant and antibacterial activities of CS coatings, and further used CS graft copolymer for food preservation [9,10]. In our previous study, CS-g-CA was attained by 1-(3-dimethylaminopropyl)-3ethylcarbodiimide hydro (EDC)/N-hydroxy succinimide (NHS) coupling reaction, which displayed superior antioxidant and antimicrobial activities [11].

https://doi.org/10.1016/j.ultsonch.2022.106032

Received 23 March 2022; Received in revised form 25 April 2022; Accepted 6 May 2022 Available online 11 May 2022

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Fig. 1. Schematic diagram showing the experimental design.

At present, static coating method of immersion or impregnation is mostly used to preserve fish, but the coating efficiency is low. Ultrasound (US) is a eco-friendly and non-thermal technology to reach a variety of uses, such as defoaming, crystallization, thawing and so on [4]. Meanwhile, US technology has broad application prospects in food processing, preservation and safety [12]. US could inhibit some enzymes and inactivate microorganisms in food preservation. Compared to conventional method, ultrasound induces mechanical, physical and biochemical changes through cavitation, so as to improve the quality of aquatic products, destroy microbial cells and alter the structural and functional properties of proteins [13]. At present, the application of US to prolong the shelf-life of food has received extensive attention. Wang et al. [14] reported that US treatment could significantly promote the mass transfer of water and improved the salted-dried grass carp's appeal for customers. Lan et al. [15] found that US incorporated with slightly acidic electrolyzed water could prolong the shelf life of sea bass for another 4 days during cold storage. Sun et al. [16] showed that chitosan nanocomposite water retaining agent combined with ultrasonic treatment can significantly reduce the quality deterioration of crayfish during frozen storage. There is little information on united technologies to extend the shelf-life of aquatic products. Therefore, the aim of this research was to research the effects of ultrasound-assisted chitosan grafted caffeic acid coating on the quality attributes and microbial composition of pompano (Trachinotus ovatus) in terms of TVB-N, TBA, nucleotides, biogenic amines, microbial enumeration and highthroughput sequencing.

# 2. Materials and methods

#### 2.1. Sample preparation

The synthesis of G was performed by EDC/NHS coupling according to the previous study [11]. The grafting ratios of CS-g-CA were 93.68 mg/g.

Lived pompano were bought from the local market (Shanghai, China) and transported to the laboratory alive within 30 min. The mean  $\pm$  SD weight and body length of fish was 500  $\pm$  50 g and 30  $\pm$  5 cm,

respectively. The experimental design was shown in Fig. 1. Next, the head, tail and guts were removed. Four groups of samples were prepared and the treatments were applied to the fillets with a fillet/solution ration of nearly 1:5 (m/v): (1) Distilled water (CK); (2) 20 kHz 600 W ultrasound (US); (3) 1% CS-g-CA in the 1% acetic acid solution (G); (4) 1% CS-g-CA in the 1% acetic acid solution with 20 kHz 600 W ultrasound (USG).

# 2.2. Total volatile basic nitrogen (TVB-N)

According to Lan et al. [17], TVB-N was measured by micro Kjeldahl nitrogen meter (FOSS, Shanghai, China). The value of TVB-N was expressed as mgN/100 g muscle.

# 2.3. Thiobarbituric acid reactive substances (TBA)

According to Feng et al. [18], the TBA of samples was done. 5.0 g of samples were homogenized with 10 mL of 5% trichloroacetic acid (TCA) for 2 min. The homogenate was centrifuged twice at  $8500 \times g$  at 4 °C for 10 min and the filtrates were gathered and then transferred into a volumetric flask and diluted with TCA to a final volume of 50 mL. Then, 5 mL of filtrate and 0.02 mol/L TBA solution were mixed, followed by boiling in a water bath for 20 min. After cooling to room temperature, the absorbance of the final colored solution was measured at 532 nm. The results were represented by mg of malonaldehyde (MDA)/g of sample.

# 2.4. Adenosine triphosphate (ATP)-related compounds

ATP and its related compounds were extracted according to the previous study[19], which were measured by HPLC (LC-2010C HT, Shimadzu Corporation). The pH value of phosphate buffer as mobile phase was 6.5 and the flow rate was 1.0 mL/min. The injection volume was 10  $\mu$ L and selects 254 nm as the detection wavelength. The amounts of ATP-related compounds were defined and analyzed on the basis of adenosine triphosphate (ATP), adenosine diphosphate (ADP), adenosine monophosphate (AMP), inosine monophosphate (IMP), hypoxanthine



Fig. 2. Effects of different treatments on the variation of TVB-N (A) and TBA (B) in *Trachinotus ovatus* during ice storage. Within the same storage time, different lowercase superscripts indicate significant differences (P < 0.05).

riboside (HxR), and hypoxanthine (Hx) standards (Sigma-Aldrich (Shanghai) Trading Co., Ltd., Shanghai, China). Ki, K, and H values were determined as follow:

$$K \text{ value } (\%) = \frac{HxR + Hx}{ATP + ADP + AMP + IMP + HxR + Hx} \times 100$$
  
Ki value  $(\%) = \frac{HxR + Hx}{IMP + HxR + Hx} \times 100$   
H value  $(\%) = \frac{Hx}{IMP + HxR + Hx} \times 100$ 

# 2.5. Biogenic amines (BAs)

BAs were determined according to the method of Zhao et al. [20]. 5 g of minced flesh were homogenized with 10 mL perchloric acid (0.6 M), and the homogenate was centrifuged at  $10,000 \times g$  for 5 min. The sediment was washed with 10 mL perchloric acid and centrifuged as before. The combined supernatants were adjusted to 25 mL and used for further analysis. BAs were derived with dansyl chloride (Sigma-Aldrich Trading Co., Ltd., Shanghai, China) and analyzed by HPLC. Putrescine and cadaverine were identified by comparing to external standards (Sigma-Aldrich Trading Co., Ltd., St. Louis, USA).

#### 2.6. Microbial enumeration

All microbiological enumeration were measured by previous study [11]. Fish flesh (5 g) was put in a sterile bag and homogenized 5 min in 45 mL of 0.85% (w/v) sterilized normal saline with an Interscience Bag Mixer (HaiBo Biological Technology Co., Ltd, Qingdao, China) and then the homogenate was serially diluted. Total viable count (TVC) and *Pseudomonas* bacteria count were determined using plate count agar and *Pseudomonas* CFC Selective Agar (HaiBo Biological Technology Co., Ltd, Qingdao, China) at  $30 \pm 1$  °C for  $72 \pm 3$  h, and then enumerated. Psychrophilic bacteria count (PBC) was determined using the plate count agar and then enumerated after 10 days of incubation at  $4 \pm 1$  °C. The H<sub>2</sub>S-producing bacteria count was determined using Iron Agar (HaiBo Biological Technology Co., Ltd) at  $30 \pm 1$  °C for  $72 \pm 3$  h. All counts were performed in triplicate and are expressed as log CFU/g.

# 2.7. High-throughput sequencing

Bacterial DNA was extracted using the bacterial genome extraction kit (Nanjing Jiancheng Bioengineering Institute, Jiangsu) and highthroughput sequencing was done according to the previous method[17].

#### 2.8. Statistical analysis

Results were analyzed with SPSS software (Version 18.0, Inc., Chicago, IL) and done with triplicate. All analysis used the least significant difference (LSD) method and Duncan's test at a significance level of 0.05. All the results were showed as "average value  $\pm$  standard deviation".

3. Results and discussion.

#### 2.9. TVB-N and TBA analysis

TVB-N could represent the quality of fishery products. Fig. 2 showed that the TVB-N values of fresh samples were 10.26-11.92 mg N/100 g at day 0. The increase of TVB-N values in all the samples was less visible at the first 16 days of storage. Since then, TVB-N values increased rapidly, which was consistent with the results of other studies<sup>[21]</sup>. Significant decrease of TVB-N was found after 20 days of storage in G and USG groups compared to the CK group. According to the research of Liu et al. [19], the TVB-N value of 30 mg N/100 g could be used as an acceptable threshold for the spoilage of marine fish. At day 20, the TVB-N value of CK group came up to 21.14 mg N/100 g. The TVB-N values of the US, G, and USG groups were only 19.1, 12.2 and 10.26 mg N/100 g, respectively. However, TVB-N values showed a rapid upward trend in the later storage, which was similar to the results of other studies[22]. The increase of TVB-N value is due to the increase of endogenous enzymes, spoilage bacteria and the subsequent biochemical reactions leading to the increase of microbial degradation substances. The lower content of the treated samples was mainly due to the presence of bioactive phenolic compounds in the coating, which significantly inhibited the ability of bacteria to oxidize and deaminate non-protein nitrogen compounds. It could be inferred, as an antibacterial coating, chitosan grafted caffeic acid coating could more effectively inhibit the growth of TVB-N value, suggesting that ultrasound could be used as an auxiliary means to enhance the inhibitory effect of grafts on microbial growth. Similar results showed that chitosan based coating could inhibit the production of TVB-N, and ultrasonic assisted treatment could further enhance this inhibition[4].

As a vital index to evaluate lipid oxidation, Changes in TBA value of all treatments were shown in Fig. 3. The initial TBA value of fresh meat was 0.22–0.39 mg MDA/kg, and there was no significant change in all treated group during the first 16 days of storage. Afterward, the TBA value of CK group sharply increased from 0.64 mg MDA/kg to final value of 1.25 mg MDA/kg. This may be due to the accumulation of lipid peroxide and peroxide degradation products during the oxidation of unsaturated fatty acids[22]. However, it could be found that the growth rate of TBA value in the CK sample was significantly higher than that in the treatments. Additionally, the USG group had the lowest growth rate



Fig. 3. Effects of different treatments on the variation of IMP (A), HxR (B), Hx (C), Ki (D), H (E), K value (F) in *Trachinotus ovatus* during ice storage. Within the same storage time, different lowercase superscripts indicate significant differences (P < 0.05).

with being recorded at 0.45 mg MDA/kg, it may be due to its inhibitory effect on lipid oxidation, its ability to scavenge free radicals and improve the activity of antioxidant enzymes in vitro. In addition, no statistic difference ( $p \geq 0.05$ ) of TBA value between G and USG indicated ultrasonic treatment has little positive effect on delaying lipid oxidation of fish fillets. This was consistent with the report of Shokri et al.[23], who reported that EO-chitosan coating could prevent lipid oxidation in fish samples.

# 2.10. ATP-related compounds, H, Ki and K value

Generally, the important variation happened in IMP,  $H_XR$  and  $H_X$ , while the contents of ATP, ADP and AMP were low during the whole storage process[24]. ATP degradation could produce IMP, which is the vital flavor substance of fish and usually has a high content in early

storage. With the extension of storage time, IMP was gradually degraded into bitter substances  $H_XR$  and  $H_X$ . As shown in Fig. 3, the initial concentrations of IMP were 0.04–0.47 µmol/g, and their concentration of all groups presented an increasing trend first and then a downward trend. IMP contents in the CK, US, and USG obtained maximum values of 6.16, 6.17, and 8.37 µmol/g at day 8, and G reached maximum value of 6.60 µmol/g at day 4. The higher IMP content in USG indicated that the active components in ultrasonic assisted chitosan grafted caffeic acid coating may reduce the activity of enzymes involved in IMP changes[24]. The remaining IMP content in the CK, US, G, and USG groups after 24 days of storage decreased to 1.87, 2.21, 3.37, and 3.75 µmol/g. The initial content of HxR was 0.13–0.16 µmol/g in fresh samples. For the CK group, the HxR content had a steep rise to 2.13 µmol/g at day 20. Meanwhile, the content of HxR decreased significantly (0.86 µmol/g) for USG compared to the CK group. The concentration of HxR decreased in

#### Table 1

Changes in biogenic amines concentration of *Trachinotus ovatus* during ice storage.

Biogenic amines (mg/kg)	Storage time(d)	СК	US	G	USG
Putrescine	0	0.02 ±	0.01 ±	0.04 ±	0.01 ±
		0.02ª	0.01	0.035	0.015
	4	$1.81 \pm$	$1.57 \pm$	$1.57 \pm$	1.54 ±
	_	0.09ª	0.03	0.06	0.08
	8	$1.89 \pm$	$1.93 \pm$	$1.88 \pm$	$1.88 \pm$
		$0.01^{a}$	$0.08^{a}$	0.03 <sup>a</sup>	$0.01^{a}$
	12	$2.02 \pm$	$1.80 \pm$	$1.95 \pm$	$1.93 \pm$
		$0.11^{a}$	$0.10^{a}$	0.06 <sup>a</sup>	$0.06^{a}$
	16	$2.03 \pm$	1.88 $\pm$	$1.93 \pm$	$1.88 \pm$
		$0.14^{a}$	$0.03^{a}$	$0.07^{a}$	$0.06^{a}$
	20	$2.72 \pm$	$2.51 \pm$	$2.04 \pm$	$2.04 \pm$
		0.09 <sup>a</sup>	0.23 <sup>a</sup>	0.20 <sup>b</sup>	0.09 <sup>b</sup>
	24	$3.99 \pm$	$2.82 \pm$	$2.29 \pm$	$2.03~\pm$
		0.14 <sup>a</sup>	0.23 <sup>b</sup>	0.01 <sup>c</sup>	0.06 <sup>c</sup>
Cadaverine	0	$0.13~\pm$	0.13 $\pm$	0.14 $\pm$	0.11 $\pm$
		$0.01^{a}$	$0.02^{a}$	$0.01^{a}$	$0.02^{a}$
	4	0.17 $\pm$	0.14 $\pm$	0.16 $\pm$	0.14 $\pm$
		0.04 <sup>a</sup>	$0.01^{a}$	0.04 <sup>a</sup>	$0.02^{a}$
	8	$0.21~\pm$	0.20 $\pm$	$0.16 \pm$	0.15 $\pm$
		0.03 <sup>a</sup>	$0.01^{ab}$	0.01 <sup>bc</sup>	0.01 <sup>c</sup>
	12	0.26 $\pm$	0.23 $\pm$	0.25 $\pm$	0.23 $\pm$
		$0.02^{a}$	$0.01^{a}$	$0.03^{a}$	0.04 <sup>a</sup>
	16	0.48 $\pm$	$0.36 \pm$	0.28 $\pm$	0.26 $\pm$
		0.06 <sup>a</sup>	$0.05^{b}$	$0.01^{b}$	$0.02^{b}$
	20	1.87 $\pm$	1.68 $\pm$	$0.60 \pm$	0.46 $\pm$
		0.07 <sup>a</sup>	$0.06^{b}$	0.06 <sup>c</sup>	0.04 <sup>c</sup>
	24	$2.10~\pm$	1.78 $\pm$	1.01 $\pm$	0.87 $\pm$
		$0.12^{a}$	$0.06^{b}$	0.04 <sup>c</sup>	0.07 <sup>c</sup>

the CK group at the end, because HxR decomposed into Hx. Hx of all samples presented an increasing trend during storage. Hx in the group was 0.78 and 0.64  $\mu$ mol/g at day 0, which exceed 1.00  $\mu$ mol/g in the CK and US groups at the end of storage.

Gao et al.[25] suggested that K value <20% indicates that the fish is very fresh, while K value >70% indicates that the fish is unacceptable. The CK showed a sharp increase of K value, exceeding the threshold at day 20, when treated groups were less than it. Besides, Ki and H value showed a similar slow increase trend in treatments. The accumulation of Hx and HxR in fish fillets would lead to a sharp increase in Ki and H values. Gao et al.[25] indicated that rosemary extract combined with nisin treatment could effectively reduce ATP loss and prolong the shelf-life of fish.

# 2.11. Biogenic amines

Within the same storage time, different lowercase superscripts indicate significant differences (P < 0.05).

Two vital BAs (putrescine and cadaverine) were found in fish that could test the safety and characteristics of fish. The formation of BAs in aquatic products depends on various factors, such as the free amino acid contents in fish, the presence of bacterial biogenic amine decarboxylases and favorable environmental conditions[26]. The initial putrescine and cadaverine in fresh fillets was 0.01-0.04 and 0.11-0.14 mg/kg (Table 1). With the extension of storage time, the concentration of putrescine and cadaverine rose gradually. The concentration of putrescine in the CK sample was significantly higher than those in the other samples (P < 0.05). Putrescine in the CK group increased to 3.99 mg/kg at day 24, although the concentration of USG was only 2.03 mg/kg, it was still less than that in the other samples (P < 0.05). At the beginning of storage, there was no significant difference between the treated samples and the



**Fig. 4.** Effects of different treatments on the variation of total viable count (A), *Pseudomonas* bacteria count (B), psychrophilic bacteria count(C), and H<sub>2</sub>S-producing bacteria count (D) in *Trachinotus ovatus* during ice storage. Within the same storage time, different lowercase superscripts indicate significant differences (P < 0.05).

Table 2

Phylotype coverage and alpha diversity estimation of bacterial communities in pompano (Trachinotus ovatus) during ice storage.

Storage time (d)	Sample	Number	OTUs	Shannon	Chao	Ace	Simpson	Coverage
0	CK	36,516	540	4.22	556.50	563.43	0.04	0.9998
	US	11,491	346	3.96	398.25	407.10	0.06	0.9997
	G	36,287	584	3.94	613.71	615.91	0.07	0.9999
	USG	39,975	568	3.99	613.31	615.54	0.05	0.9999
12	CK	34,585	411	3.67	461.83	446.11	0.07	0.9999
	US	34,474	340	3.86	353.64	352.57	0.05	0.9999
	G	42,683	538	1.53	576.59	581.19	0.06	0.9999
	USG	39,821	539	4.37	597.60	605.04	0.02	0.9998
24	CK	15,074	33	1.53	48.000	87.380	0.38	0.9999
	US	18,536	28	1.18	83.000	78.770	0.48	0.9999
	G	21,592	124	1.82	260.15	323.62	0.23	0.9999
	USG	25,919	352	3.3	420.02	427.23	0.09	0.9999

CK sample (P > 0.05), and then the values of putrescine in the treatments was significantly lower than that in the CK sample (P < 0.05) except for day 12, which indicated that USG was more effective in the preservation of fish fillets. At the end of storage, the content of cadaverine in CK group was 2.10 mg/kg, which was significantly higher than those in treated groups (P > 0.05). However, cadaverine content in the USG group was only 0.87 mg/kg. Similarly, Zhang et al. [27] reported that putrescine and cadaverine concentrations in common carp fillets with cinnamon essential oil was lower than control group.

#### 2.12. Microbial enumeration

The changes in microbial enumeration of *Trachinotus ovatus* fillets were presented in Fig. 4. During storage, the number of microorganisms in all groups showed an increasing trend. The initial TVC was 2.80–3.17 log CFU/g. TVC reached approximately 7.60 log CFU/g in the CK group at day 20, whereas lower TVC were observed in the US, G and USG samples at the same interval (7.10, 5.35, and 5.08 log CFU/g, respectively), indicating that G was effective in delaying microbial growth, and the effects could be further enhanced using US assisted. According to TVC of 7 log CFU/g as the standard of corruption[20]. The TVC in the CK, US, and G groups reached the acceptance value at day 20 20, and 28, respectively, whereas USG was always acceptable during storage, which was showing no difference with the TVB-N and TBA value results. Therefore, the USG group had longer shelf-life (28 days) than the CK group (20 days).

The production of off-odor and harmful metabolites was related to Pseudomonas and H<sub>2</sub>S-producing bacteria[24]. The growth trend for the counts of psychrophilic bacteria, Pseudomonas bacteria and H2S-producing bacteria was agreement with TVC. The initial counts of psychrophilic bacteria and Pseudomonas bacteria were 1.76-2.04, and 2.02-2.24 log CFU/g. H<sub>2</sub>S-producing bacteria were not found in all groups at day 0. H<sub>2</sub>S-producing bacteria were first detected in all groups at day 4. The psychrophilic bacteria counts of CK group was more than 7.0 log CFU/g at day 20, but the psychrophilic bacteria counts of US, G, and USG samples were all significantly lower than the threshold of seafood. The final bacterial number in USG reduced by 0.72 log CFU/g (Pseudomonas) and 1.08 log CFU/g (H2S-producing bacteria) when compared to the CK group. All treatments evidently reduced microbial levels. Similar results were also found for the samples treated with chitosan-based coating[24]. A number of previous studies have assessed the ability of US to inhibit the growth of microorganisms in food. Garrido et al.[28] reported that samples with US treatment were significant reduction in microbial counts for both salmon and mackerel when compared to untreated groups.

# 2.13. Microbial composition analysis using high-throughput sequencing

The microbiota composition of different treatments was determined by high-throughput sequencing. Table 2 showed that the coverage of all groups was >0.99, meaning that almost all microbial system types were detected in the samples. The Shannon, Simpson, ACE and Chao indicators were determined. 356,953 valid sequences were detected. The number of OTUs in all treatments was lower than Chao1 and ACE index, which represented that there may be some other bacterial system types in these samples. The results of Shannon and Simpson showed that the bacterial diversity of samples at the end was lower than that at the beginning of storage, which was agreement with other study of Jia et al. [29].

The relative abundances of the top 46 genera at the genus levels of all samples were represented by histogram. As shown in Fig. 5A, on day 0, unclassified Bacteria, *Acinetobacter*, and *Bradyrhizobium* were the dominant microbiota in the CK group, which accounted for 41.78%, 14.36%, and 6.93% of all OTU numbers, respectively. Because the ability of *Acinetobacter* not produce some corrupt substances, the factors leading to corruption were relatively weak[30]. However, at the initial stage, the relative abundance of microbiota in sample with different treatments were changed and then *Pseudomonas, Chryseobacterium*, and *Phenylobacterium* became the dominant microbiota in the US, G, and USG samples, respectively.

The microbiota composition at the beginning of storage was detected by high-throughput sequencing, and the results were consistent with the results of microbial count. H<sub>2</sub>S-producing bacteria were not detected by the microbial enumeration method in all groups at the initial stage of storage. The relative abundance of *Shewanella* was also less than 0.5% of bacterial populations in the all samples at day 0, which revealed by high throughput sequencing. In addition, *Pseudomonas* which had been identified as the dominant microorganism in all four groups at day 24 by microbial count was detected at 8.97%, 78.79%, 68.95%, and 41.55% in the CK, US, G, and USG groups, respectively.

As for the CK group, *Lactococcus* was found to be the prevalent bacteria at day 12. After 24 days, *Brochothrix, Pseudomonas*, and *Shewanella* sharply increased to 59.78%, 8.97% and 11.44% became the predominant bacteria, whereas, *Lactococcus* were decreased to negligible level. For US and G, *Pseudomonas* accounted for 0.14%, and 1.49% at day 0. Then *Pseudomonas* rose with the increase of storage time. At the end of the shelf-life, *Pseudomonas* (78.79% and 68.95%) was the dominant bacteria. *Pseudomonas* was the one of the main TVB-N producers [31], correspondingly, a sudden increase in TVB-N was showed in all groups at the end of storage. The abundances of *Shewanella* in treated groups were reduced compared with the CK group. For USG, microbial diversity was still high at the end of storage. *Pseudomonas, Shewanella*, and *Janthinobacterium* accounted for 41.55%, 12.95%, and 9.72%, respectively.

Α

С



Fig. 5. Community treebarplot analysis (A), Community heatmap (B), and PCA (C) on genus level of microbiota at the genus level of microbiota in *Trachinotus ovatus* during ice storage.

#### Table 3

Regression correlation coefficients between microbial enumerations and biochemical indicators (TVB-N, TBA, Putrescine, Cadaverine, K value, and TVC) in pompano (*Trachinotus ovatus*) during ice storage.

Parameters	Shewanella	Vagococcus	Pseudomonas	Brochothrix	Acinetobacter	Carnobacterium
К	0.489	0.492	0.652*	0.630*	-0.792**	0.772**
TVBN	0.202	0.715**	0.374	0.846**	-0.453	0.950**
Putrescine	0.596*	0.325	0.403	0.657*	-0.806**	0.755**
Cadaverine	0.358	0.655*	0.547	0.808**	-0.645*	0.929**
TBA	0.538	0.395	0.121	0.924**	-0.493	0.897**
TVC	0.564	0.435	0.54	0.662*	$-0.782^{**}$	0.801**

Note. Levels of significance and defined as \*p < 0.05 and \*\*p < 0.01.

As shown in Fig. 5B, the composition and dynamics of bacterial community in all samples were analyzed using the heat map of common microorganisms at the genus level. The relative abundance of the genus was represented by the type and depth of color, the heat map showed that the diverse of microbiota in the fresh fish was the highest at day 0. Burkholderia, Streptophyta, Chryseobacterium, and Brevundimonas occupied a large proportion in fresh samples, but almost disappeared when samples deteriorated. After storage for a period, relative abundance of most microbiota decreased. Similarly, Zhang et al.[32]also identified that the bacterial diversity of grass carp fillets declined during storage. Brochothrix, Pseudomonas, and Lactococcus were abundant in spoiled fish. Pseudomonas has been reported to be a spoilage inducer and contributed to the formation of putrescine and cadaverine in various fish species from fresh and marine waters under aerobic chilled storage[3]. It is reported that Brochothrix will also affect the character of aquatic products related to corruption. Corruption will mainly produce lactic acid, ethanol and a small amount of short chain fatty acids, resulting in peculiar smell[33]. Some low abundance microbial communities are not well showed in the bar chart, but their variations are clearly reflected in the heat map.

In order to compare the microbial composition of each group, the analysis of PCA were used (Fig. 5C). The clustering results showed that the microbial community changed dynamically during storage. The similarity of microbial community structure was high in all groups at the initial stage. Meanwhile, the CK, US G, and USG groups represented two clusters at day 24. The results showed that the preservative treatment changed the microbial community structure of fish during storage. Yu et al. [24]showed that compound preservative treatment could alter the microbial community composition of fish fillets during storage.

# 2.14. Relationship of microflora and quality indicators

The accumulation of harmful components such as TVB-N, TBA,  $H_X$  and  $H_XR$  is the vital reason for the deterioration of fish quality. The correlation coefficients between 6 kinds of bacteria, and biochemical indexes for fish fillets were presented in Table 3. All quality index showed significant (P < 0.05) correlations with *Carnobacterium*. Most quality indicators associated with *Brochothrix* and *Acinetobacter* values significantly. *Acinetobacter* showed negative correlation with all quality indexes. This may be due to the relatively weak ability of *Acinetobacter* to cause corruption, because it would not produce some corrupt substances[30]. *Shewanella* was significantly correlated with putrescine value, which could cause the formation of putrescine.

#### 3. Conclusions

The current research indicated that ultrasound-assisted chitosan grafted caffeic acid coating effectively inhibit the increase of components including TVB-N, TBA, HxR and Hx, and decreased K value, microbial reproduction and the reduction of IMP in pompano (*Trachinotus ovatus*) during ice storage. Meanwhile, the results of high throughput sequencing analysis presented that the microbial diversity of all samples at the end of storage was lower than that of fresh sample. *Acinetobacter* was the major microorganism in fresh pompano. *Shewanella*, and

*Brochothrix* became the major microbiota in the CK samples after 24 days of storage. In addition, *Brochothrix* in treatments were effectively controlled. Treatment changed the microbial communities of *Trachinotus ovatus* fillets during storage. All of the indicators represented that the ultrasound-assisted chitosan-based coating retarded the bacterial spoilage and extend the shelf-life of *Trachinotus ovatus*.

# CRediT authorship contribution statement

Weiqing Lan: Conceptualization, Writing - review & editing, Supervision, Visualization. Yuqing Sun: Investigation, Methodology, Formal analysis, Data curation, Writing - original draft, Writing - review & editing, Software. Shucheng Liu: Project administration, Resources. Yuan Guan: Investigation, Methodology. Shengyun Zhu: Investigation, Methodology. Jing Xie: Project administration, Resources.

# **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.

# Acknowledgements

The work was financially supported by China Agriculture Research System of MOF and MARA (CARS-47), Guangdong Provincial Key Laboratory of Aquatic Product Processing and Safety (GDPKLAPPS1802), Southern Marine Science and Engineering Guangdong Laboratory (Zhanjiang) (ZJW-2019-06), Guangdong Innovation Team of Seafood Green Processing Technology (2019KCXTD011).

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