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Hemocyanin is a major allergen in local mud crab *Scylla paramamosain* in Viet Nam

Kieu-Minh Le^{1®}, Hoang Kim Tu Trinh^{1,*®}, Thanh Niem Vo Van^{1®}, Duy Le Pham^{2,3®}, Hieu Thao Nguyen^{4®}, Thien-Tai Tran^{3®}, Minh Nguyet Tran Thi⁵, Bao Yen Pham⁵, Dinh Minh Pham⁶

ABSTRACT

Introduction: Shellfish allergy is prevalent in coastal countries across Asia. Despite crab being a popular dish, there is limited knowledge about the features of crab allergy in Viet Nam. This study aims to identify the IgE-binding allergens in the local crab species, Scylla paramamosain (S. paramamosain). Methods: The study involved 12 patients with crab allergy and 5 subjects with crab tolerance. Each participant underwent a skin prick test (SPT) with crab extracts and was measured for specific IgE levels to crab, as well as house dust mite species Dermatophagoides farinae (DF) and Dermatophagoides pteronyssinus (DP). An immunoblot was used to identify IgE-binding proteins in S. paramamosain, followed by identification using nanoscale liquid chromatography coupled with tandem mass spectrometry analysis. **Results**: Patients with crab allergy exhibited higher total IgE levels (2432 \pm 772.26 vs 886.15 \pm 3056.74), 100% positive SPT to crab, and higher specific IgE levels to crab (3.40 \pm 6.19 vs 0.48 \pm 8.86) compared to subjects with crab tolerance. The level of specific lgE to crab showed significant correlation with SPT results to DP and DF (r = 0.52, P = 0.033; r = 0.82, P < 0.001, respectively). The frequencies of IgE-binding proteins in crab allergy patients were higher than in tolerant subjects. Several putative allergens were identified, including hemocyanin as a major allergen, with arginine kinase, sarcoplasmic-calcium binding protein, and triosephosphate isomerase as minor allergens. **Conclusion**: This study is the first to demonstrate potential molecular allergens in local mud crabs among Vietnamese individuals sensitized to house dust mites. These findings require further clinical evaluation in future studies for their application. Key words: Allergen, crab allergy, Viet Nam, hemocyanin

¹Center for Molecular Biomedicine, University of Medicine and Pharmacy at Ho Chi Minh City, Ho Chi Minh City 700000, Viet Nam

²Faculty of Medicine, University of Medicine and Pharmacy at Ho Chi Minh City, Ho Chi Minh City 700000, Viet Nam

³University Medical Center Ho Chi Minh City, Ho Chi Minh City, 700000, Viet Nam

⁴*Tra Vinh University, Tra Vinh 87000, Viet Nam*

⁵The Key Laboratory of Enzyme and Protein Technology (KLEPT), VNU University of Science, Hanoi 100000, Viet Nam

⁶GENTIS JSC, 505 Minh Khai, Hai Ba Trung, Hanoi 100000, Viet Nam

Correspondence

Hoang Kim Tu Trinh,

Center for Molecular Biomedicine, University of Medicine and Pharmacy at Ho Chi Minh City, Ho Chi Minh City 700000, Viet Nam

Email: kim.tu.vn@ump.edu.vn

INTRODUCTION

Over the past few decades, there has been a global increase in the prevalence of food allergies 1,2 . As food allergy depends on eating habits, each country's burden and food allergens will differ³. In Asia, the population-based estimates of food allergy prevalence range from 1.1% to 7.7%². The spectrum of food allergens in Asia shows different features compared to Western countries, with the predominance of shell-fish and wheat³. In Viet Nam, where seafood plays a significant role in the daily diet, seafood allergy is most commonly observed among subjects with food allergies ^{4,5}. Surveys have shown that IgE-mediated crustacean allergy is 2.6% in adults and 2.8% to 4.79% in children, representing the highest prevalence compared to other allergens in Viet Nam⁵.

Among crustaceans, crabs are highly consumed. According to the literature, the genus Scylla consists of four species of swimming crabs: *Scylla serrata*, *Scylla olivacea*, *Scylla paramamosain*, and *Scylla tranquebarica*^{6,7}. *S. paramamosain* is a type of crab frequently

consumed in Southeast Asia. The increased consumption of crustacean products can be associated with a higher risk of allergy. Snow crab sensitization is common among workers and may contribute to occupational asthma^{8,9}. The prevalence of crab allergy is 15.2% in subjects with food allergies in Singapore [10] and 12.7% in Chinese asthmatic children^{10,11}. Until July 2022, Viet Nam was estimated to export 38 million U.S. dollars worth of crabs to the USA¹². Therefore, evaluating the characteristics of crab allergens among local fishery products is necessary.

The mechanisms of crab allergy can be roughly divided into IgE- and non-IgE-mediated pathways^{13,14}. In cases where allergic reactions are triggered by an immune response mediated through IgE, patients can experience various symptoms ranging from mild to severe, including potentially fatal anaphylactic shock. It is worth noting that, among allergens within the same type of food, some tend to bind to IgE more frequently, potentially causing more severe reactions than others, and this can vary from person to person^{13,15}. Moreover, different types of food may contain similar allergens, leading to the possibility of

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cross-reactivity. Consequently, a comprehensive understanding of allergen components is crucial for diagnosing food allergies. Several key allergen components in seafood have been identified, such as tropomyosin (33-39 kDa), arginine kinase (AK) (38-45 kDa), and hemocyanin (HM) (77 kDa), which are prevalent in various shellfish species^{14,16-18}. However, research on the molecular allergen components of crabs in Viet Nam still needs to be expanded. In this study, we aimed to identify the molecular allergens in patients with an allergy to mud crab (Scylla paramamosain). Briefly, raw allergen extracts were obtained from local mud crabs. These allergens were incubated with the patient's serum to examine the components typically bound to the IgE antibodies. Subsequently, IgE-bound proteins were digested and identified using nanoscale liquid chromatography coupled with tandem mass spectrometry (nano-LC/MS).

METHODS

Patient Recruitment

Patients with suspected crab allergy were enrolled from the Unit of Allergy and Clinical Immunology, University Medical Center (Ho Chi Minh City, Viet Nam). All subjects provided informed consent and underwent an allergy assessment. Participants were interviewed about their history of allergy and reactions after crab consumption. The diagnosis of crab allergy was confirmed if participants experienced either anaphylaxis after crab consumption or repetitive allergic events after crab consumption at home. Subsequently, participants were divided into two groups: crab allergic (CA) and crab tolerant (CT) groups. All participants underwent skin prick tests (SPT) with the following allergens: house dust mites, Dermatophagoides farinae (DF), Dermatophagoides pteronyssinus (DP), and extracts of raw crab S. paramamosain. The SPT was conducted according to the instructions of the European Academy of Allergy and Clinical Immunology¹⁹. Whole blood was collected from the participants in blood collection tubes without EDTA. The obtained blood samples were allowed to coagulate by being left undisturbed for 30 minutes at room temperature (RT) and then centrifuged at 3000 rpm for 10 minutes at 4°C. Patient sera were stored at -80°C for further use. The University of Medicine and Pharmacy Ethics Committee at Ho Chi Minh City approved this study.

Preparation of Crude Extracts

Mud crabs (Scylla paramamosain), approximately eight months old and female, were purchased from

a local market before tissue samples were collected. Crude extracts were prepared as previously described²⁰. Briefly, approximately 5 g of raw muscle tissues in the crab claws were homogenized in 20 mL of phosphate-buffered saline (PBS) extraction buffer and incubated overnight at 4°C on a constant temperature shaker. Subsequently, the extracts were centrifuged at 5000 \times g for 20 minutes at 4°C. The supernatant was transferred to 100K Omega Microsep Advance Centrifugal Devices (Pall, USA - MCP100C41) for protein purification and centrifuged at 5000 imes g for 20 minutes at 4°C. The filtered supernatant containing the retained bioactive compounds was preserved at -80°C. Protein concentrations were measured using PierceTM Bradford Plus Protein Assay Kits (Thermo Fisher Scientific, Massachusetts, USA).

Visualization of Allergens Using Sodium Dodecyl Sulfate-Polyacrylamide Gel Electrophoresis (SDS-PAGE)

Food allergens were subjected to 10-12% SDS-PAGE, followed by both Coomassie blue staining and electrotransfer into PVDF membranes. SDS-PAGE gels were then incubated in destain solution (10% methanol + 10% acetic acid). The SDS-PAGE gel containing the proteins was placed on a glass plate illuminated by a light box, and a razor blade was used to excise an individual band from the gel and place it into a microcentrifuge tube.

Detection of slgE-Binding Epitopes Using Immunoblot

Patient sera (1:10 dilution) were used for immunoblotting, followed by the addition of HRPconjugated anti-human IgG (ThermoFisher Scientific, USA). Immunocomplexes were visualized with 1-StepTM TMB-Blotting Substrate Solution (Thermo Fisher Scientific).

In-Gel Protein Digestion

The protein bands on the SDS-PAGE gel were sliced into small pieces and subjected to destain for 30 minutes until the gel particles became clear. Gel pieces were dehydrated by adding 100 μ L of 100% acetonitrile (ACN). Then, ACN was removed and the gel pieces were left to air-dry for 5-10 minutes. The gel pieces were then washed with a 50:50 mixture of 100 mM NH₄HCO₃ and 50% ACN for 10 minutes, which was then discarded. The samples were resuspended in 10 mM DTT in 100 mM NH₄HCO₃ for 45 minutes for reduction, followed by incubation with 50 μ L of 50 mM iodoacetamide in 100 mM NH₄HCO₃ and kept in the dark for 30 minutes. The gel pieces were washed as described above. Afterwards, gel pieces were rehydrated on ice with 10 ng/ μ L trypsin in 50 mM NH₄HCO₃ with 10% ACN at 4°C for 30 minutes. Fifty mM NH₄HCO₃ was added to cover the gel pieces, and the tubes were placed in a 30°C water bath overnight. To terminate the reaction, 30 μ L of 1% formic acid was added and centrifuged at 13,000 rpm for 5 minutes to collect the peptide-rich first supernatant in a labeled Eppendorf tube. The gel pieces were incubated with 50% ACN and 5% formic acid for 45 minutes. The tubes were subjected to sonication, followed by centrifugation at 13,000 rpm for another 5 minutes, and the second supernatant was collected in a labeled Eppendorf tube. Later, gel pieces were incubated with 90% ACN with 5% formic acid for 5 minutes. The final supernatant was collected in a labeled Eppendorf tube. Finally, all supernatants were dried via SpeedVac and desalted with a Ziptip C18 column. The pipette was set to $10 \,\mu$ L and attached to the Ziptip column. To prepare the ZipTip for peptide binding, the column was conditioned with 3 washes of 100% acetonitrile and 3 washes of 0.1% formic acid. The sample was loaded onto the column by pipetting the protein digest up and down 10 times. The ZipTip was washed with 6 washes of 0.1% formic acid. The peptides were extracted from the ZipTip by pipetting the elution buffer 10 times in a 4 μ L solution containing 60% acetonitrile/0.1% formic acid. The organic phase of the elution buffer eluted the peptides from the resin into the buffer. The sample was subsequently desalted and concentrated in a 4 μ L solution.

NanoLC-MS/MS Analysis of Tryptic Peptides

Tryptic peptides were separated and sequenced by nanoscale liquid chromatography on a nanoAcquity system (Waters, Milford, MA, USA) and MS/MS on an Orbitrap Elite hybrid mass spectrometer (Thermo Electron, Bremen, Germany). For peptide separation, a 25 cm long and 75 µm ID- C18 BEH column (Waters, Milford, MA) packed with 1.7 μ m particles with a pore size of 130 Å was used at a flow rate of 300 nL/min and a column temperature of 35°C. Peptides dissolved in Solvent A (0.1% formic acid in water) were loaded into the column and eluted by a 120minute segmented gradient from 5 to 35% solvent B (acetonitrile with 0.1% formic acid). The peptides were sequenced by the tandem MS/MS method. Initially, the mass spectrometer scanned precursor ions in orbitrap mode (m/z 350-1600, 120,000 resolution at m/z 400, and an automatic gain control (AGC) target at 10^6). The most intense ions were then isolated

for CID MS/MS fragmentation and detection in the linear ion trap (AGC target at 10,000). The mass spectrometer was operated using data-dependent scans of MS spectra, which excluded previously selected ions for 60 seconds. Single and unrecognized charge ions were also excluded for MS/MS fragmentation.

Protein Bioinformatic Analysis

Raw MS/MS data were first processed using PEAKS 6 software (Bioinformatics Solutions, Canada). The proteome database in PEAKS was taken from the UniProt Knowledgebase (UniProtKB), accessed in June 2023. Protein ID was identified by searching against the database with known parameters for protein modifications [variable oxidation (M) and fixed carbamidomethylation (C)]. Accepted peptides in protein ID were limited to two missed cleavages, and thresholds for parent mass inaccuracy and precursor mass tolerance were chosen at 10 ppm and 0.6 Da. The false detection rate (FDR) was set at 1%, and identified proteins must have had at least two unique peptides.

Statistical Analysis

The normality of the data was checked with the Shapiro–Wilk test. For continuous variables, data were presented as median \pm standard error (SE), and comparisons between the two groups were performed using the Mann–Whitney U test. For categorical variables, data were displayed as N (%), and Fisher's exact test was used to compare the two groups. Correlation was calculated by Spearman's rank correlation. Data analysis was performed using the statistical software packages IBM SPSS 20.0 (Armonk, NY, USA), and JASP (Version 0.18.3) [Computer software]. Graphs were prepared using GraphPad Prism 8 (San Diego, CA, USA). A P-value of less than 0.05 was considered significant.

RESULTS

Characteristics of subjects with mud crab allergy

In **Table 1**, 17 subjects were enrolled in this study, including 12 subjects with mud crab allergy (CA) and 5 with mud crab tolerance (CT). The median age for all participants was 29 years old, while the CT participants were relatively younger (27 years old). There was a predominance of females among the participants. Regarding allergic assessment, CA participants were more atopic than CT subjects, as indicated by the higher level of total IgE (2432 \pm 772.26 vs. 886.15 \pm 3056.74) and a higher percentage of positive SPT

Characteristics	All (N=17)	Crab allergic group	Crab tolerant group	P-values
		(N=12)	(N=5)	
Age (*)	29 ± 1 (*)	29 ± 1	27 ± 2	0.79
Sex (female, %)	11 (66.11%)	7 (58.33%)	4 (80%)	0.24
Comorbid fish/mollusk allergy	0	0	0	NA
Total IgE (*)	2411 ± 869.65	2432 ± 772.26	886.15 ± 3056.74	0.48
sIgE to seafood mix (*)	0.35 ± 3.11	0 ± 4.02	0.56 ± 3.02	0.48
sIgE to crab (*)	2.05 ± 5.05	3.40 ± 6.19	0.48 ± 8.86	0.55
Wheal diameter of Der P SPT (*)	7 ± 1.3	8 ± 1.5	5 ± 2.6	0.35
SPT to Der P (+) (*)	16 (88.89%)	12 (100%)	4 (80%)	1
Wheal diameter of Der F SPT	6 ± 1.7	7 ± 2	4 ± 3	0.20
SPT to Der F (+)	15 (83.33%)	12 (100%)	3 (60%)	0.43
Wheal diameter of crab SPT	4 ± 0.5	4 ± 0.5	3.5 ± 1.1	0.79
SPT to crab (+)	15 (83.33%)	12 (100%)	3 (60%)	0.43
Co-sensitization to Der P and crab	15 (83.33%)	12 (100%)	3 (60%)	0.43
Co-sensitization to Der F and crab	15 (83.33%)	12 (100%)	3 (60%)	0.43

Table 1: Demographic characteristics of study subjects

Categorical data were shown as N (%). P values between groups were calculated by Fisher's exact test. (*) Continuous data were shown as median \pm SE. P-values between groups were calculated by Mann-Whitney U test.

Table 2: Nano LC/MS search results summary of IgE-binding crab proteins

Band No.	MW (kDa) observed	MW (kDa) Predicted	Protein identification	Organism	Accession No.	Coverage of protein sequence
1	20-25	21.9	Sarcoplasmic calcium-binding protein	Scylla paramamosain	I2DDG2 SCP_SCYPA	59
2	25-35	26.7	Triosephosphate isomerase	Macaca mulatta	P15426 TPIS_MACMU	14
3	35-50	40	Arginine kinase	Scylla paramamosain	H6VGI3 KARG0_SCYPA	86
4	50-70	40	Arginine kinase	Scylla paramamosain	C9EIP1 KARG_SCYSE	32
5	70-100	75	Hemocyanin	Scylla serrata	C9EIP1 KARG_SCYSE	12

to DP (100% *vs.* 80%) and to DF (100% *vs.* 60%). However, no significant differences were observed between the groups.

Sensitization to mud crab and clinical symptoms after crab consumption

Sensitization to mud crab was assessed by SPT using the crude extracts and levels of sIgE to crab. All CA subjects had a positive crab SPT (100%) compared to CT subjects (60%) (**Table 1**). Moreover, the levels of specific IgE to crab among CA subjects were higher compared to the CT group, $3.40 \pm 6.19 vs. 0.48 \pm$ 8.86, despite no significant differences. These data advocate that the CA group was more sensitized to the crude crab allergens than the CT group.

In terms of clinical symptoms, non-specific rashes were frequently reported with the highest prevalence (76.47%), followed by urticaria in the majority (70.59%), upper airway reactions such as sore throat/itchy throat/coughing (58.82%), sneez-ing/itchy nose (41.18%), and itchy eyes/watery eyes/eye redness (35.29%). The spectrum of allergic symptoms for each subject was described in the **Supplementary Table**.

Correlation between SPT and specific IgE

There was no significant correlation between the level of specific IgE and the wheal size of SPT to crab extracts. Specific IgE to crab correlated with both specific IgE to house dust mites DP (r = 0.52, P = 0.033) and DF (r = 0.82, P < 0.001).

Immunoblotting analysis of IgE-binding proteins among subjects with mud crab allergy

We analyzed the major and minor proteins of raw mud crab S. paramamosain. SDS-PAGE of raw mud crab S. paramamosain extract revealed protein profiles with molecular masses ranging from 15 to 240 kDa (Figure 2A). Immunoblotting using 17 sera from CA patients and CT subjects demonstrated several IgE-binding proteins at various molecular masses between 15 and 240 kDa. Generally, more IgE-binding was detected in the immunoblotting of the CA group compared to CT groups (Figure 2B-C). Proteins at five different MW ranges: 20-25 kDa, 25-35 kDa, 35-50 kDa, 50-70 kDa, 70-100 kDa, and 100-140 kDa exhibited the highest frequency of IgE-binding protein bands in the tested sera of the crab-allergic patient group. Specifically, proteins with MW ranges from 70-100 kDa and 50-70 kDa were identified as the major allergens of S. paramamosain with frequencies of 50% and 41.67%, respectively, in the CA group.

Besides, more than 10% of the tested sera in the CT group retained IgE-binding to the protein bands from 20-25 kDa and 35-50 kDa MW ranges in the *S. paramamosain* raw extract.

Allergen identification

Table 2 summarizes the major allergens of S. paramamosain. The band at 20-25 kDa correlated with sarcoplasmic calcium-binding protein (SCP) from S. paramamosain, with peptide sequence coverage of 59%. The second band at 25-35 kDa was identical to triosephosphate isomerase (TIM). Meanwhile, the bands at 35-50 kDa and 50-70 kDa major allergens were similar to arginine kinase (AK), corresponding to 86% and 32% sequence coverages, respectively. Next, the band at 70-100 kDa minor allergen was found to be identical to hemocyanin from Scylla serrata, which corresponded to 12% sequence coverage. The identified proteins were illustrated in Figure 2D. We observed another band at the highest molecular weight > 100 kDa, which was identified as arginine kinase.

Association of IgE-binding proteins with clinical relevance

Patients A3, A5, A6, and A9 showed strong reactivity to the proteins at 70-100 kDa by immunoblotting. We found that these patients had higher levels of total IgE, and specific IgE levels to crab compared to the remaining subjects ($4807.80 \pm 1276.52 \text{ vs. } 2421.50 \pm 1039.98$; $12.44 \pm 18.02 \text{ vs. } 2.72 \pm 4.09$, respectively). However, no significant differences were observed between groups. Regarding clinical manifestations, there were no specific clinical symptoms (**Supplementary Table**), yet two patients, A5 and A6, displayed various symptoms.

DISCUSSION

Vietnam has an extensive coastline and abundant seafood resources, making it one of the leading seafood exporters in Asia²¹. However, little is known about the characteristics of shellfish allergens among local fishery products. This study identified the major mud allergens of the crab *S. paramamosain* among subjects with crab allergies, including HM, AK, TIM, and SCP. These allergens may play a role in clinical reactions and sensitization to arthropod allergens in Vietnamese crab-allergic subjects.

Patients with CA exhibited various symptoms after crab consumption, with rash and urticaria predominating. This finding was similar to a survey conducted among those with Vietnamese crustacean



Figure 1: Correlation between the levels of specific IgE to crab with the wheal size of SPT (A) to crab, (B) to house dust mite DP, and (C) to house dust mite DF. Data was analyzed by Spearman's rank correlation. A P < 0.05 was considered statistically significant. Abbreviations: DF: Dermatophagoides farinae; DP: Dermatophagoides pteronyssinus.





allergies, which showed that cutaneous symptoms were the most common. Cross-reactivity between house dust mites and crab was noted, as a portion of CT subjects did not have any reactions after crab consumption, yet their SPT remained positive for crab extracts. Cross-reactivity between arthropod allergens and crustacean allergens is well demonstrated. Tropomyosin (~38 kDa) is a pan-allergen in house dust mites and cockroaches, which have a high sequence homology to shellfish tropomyosin²². Tropomyosin is identified in shrimp and coral swimmer crab Charybdis feriatus, blue swimming crab Portunus pelagicus, and mangrove crab Scylla tranquebarica²³⁻²⁵. In countries with hot and humid conditions, where house dust mite sensitization is prominent, exposure to HDM tropomyosin can be the primary sensitizer for shellfish allergy²². Regarding shrimp allergy, tropomyosin is no longer an accurate marker allergen due to the fact that there are additional shrimp allergens, such as AK, myosin light chain, SCP, HM, etc.²⁶. However, the diagnostic role of tropomyosin remains vague for crab allergy.

Depending on IgE-binding frequency, an allergen can be classified as major (>50% IgE-binding) or minor (<50% IgE-binding)²⁷. In this study, sera from CA patients mostly reacted to protein bands from 70-100 kDa, suggesting another major allergen rather than tropomyosin, which is identical to HM from S. serrata. HM (60-80 kDa) is a sizeable copper-containing glycoprotein in the hemolymph of various arthropods and mollusks²⁸. HM was considered a heat-stable, highly conserved protein. However, its physicochemical properties remain unclear²⁸⁻³⁰. Evidence has accumulated on the critical role of HM as a crustacean allergen, especially for crabs. HM exerts antimicrobial properties and is a necessary respiratory protein for invertebrates²⁹. Interestingly, HM was considered a minor allergen in the Chinese mitten crab Eriocheir sinensis²⁸ and a major non-cross-reactive allergen in the giant freshwater prawn Macrobrachium rosenbergii^{30,31}. A study from Malaysia by Jasim et al. also discovered the presence of HM in Scylla tranquebarica as an allergenic protein among crab-allergic subjects²³. Sensitization to HM contributed to occupational crab allergy among crab processing workers³². HM can be a potential allergenic protein for crab allergy in Viet Nam.

Other minor allergens of *S. paramamosain* were revealed, including AK, SCP, and TIM. These proteins were found in various shrimps and crabs^{17,23–25,31–33}. AK is a minor allergen detected in raw shrimp but a major allergen in the blue swimming crab Portunus pelagicus²⁴. Its function is

to regulate the ATP level in the muscles of invertebrates 34. Besides tropomyosin, AK is a cross-reactive allergen in crustaceans and arthropods ^{24,26,31}. Notably, AK was detected in bands of higher molecular weights, indicating the binding of IgE to oligomers. A tetrameric AK of 150 kDa was found exceptionally in some marine worms³⁵. SCP is an EF-handtype Ca²⁺ binding protein, previously identified as an allergen in Penaeus monodon (Pen m 4), Crangon crangon (Cra c 4), Litopenaeus vannamei (Lit v 4) and also in mud crab S. paramamosain³⁶⁻³⁸. Lastly, current evidence showed that TIM is a novel allergen of crab and exerts a Th₂ skewed reaction in Balb/c mice³⁹. TIM is discovered in the octopus Octopus fangsiao, which may contribute to cross-reactions between crustaceans and mollusks⁴⁰. While AK was suggestive of occupational crab allergy among workers, the clinical role of SCP and TIM remains limited and needs further evaluation.

Compared to commercial IgE tests, such as specific IgE and SPT, commercial allergens may contain crossreactive allergens or exclude some components depending on the purification method. In our study, the specific IgE to crab strongly correlated with SPT to house dust mites (DP, DF), while there was no correlation between SPT to crab and specific IgE to crab. These findings reinforced the role of molecular allergen and component-resolved diagnosis in the precise diagnosis of crab allergy. Thus, understanding the allergen components of local crab species is mandatory for component-resolved diagnosis.

A major limitation of this study was the small sample size. Therefore, the findings would be applicable to patients in our center and cannot be generalized to the Vietnamese population. Secondly, several phenotypes in food allergy have been described⁴¹. In this study, we investigated the allergens involved in IgE-mediated seafood allergy with immediate reactions after food exposure. With regard to the mixed IgE-and non-IgE mediated seafood allergy, further studies are necessary to evaluate the profile of associated allergens.

CONCLUSIONS

In conclusion, we illustrated the Vietnamese local crab's major (HM) and minor allergens (AK, SCP, TIM) among Vietnamese individuals sensitized to house dust mites. These findings contribute to the characterization of molecular allergens in seafood in Viet Nam, creating a platform for componentresolved diagnosis and precision medicine. Future studies are necessary to evaluate these proteins' clinical reactivity and diagnostic applications.

ABBREVIATIONS

ACN: Acetonitrile, AK: Arginine Kinase, ATP: Adenosine Triphosphate, CA: Crab Allergic, CT: Crab Tolerant, DF: Dermatophagoides farinae, DP: Dermatophagoides pteronyssinus, HM: Hemocyanin, HRP: Horseradish Peroxidase, IgE: Immunoglobulin, MS: Mass Spectrometry, MS/MS: Tandem Mass Spectrometry, MW: Molecular Weight, nano-LC/MS: Nanoscale Liquid Chromatography/Tandem Mass Spectrometry, SCP: Sarcoplasmic Calcium-Binding Protein, SE: Standard Error, SPT: Skin Prick Test, TIM: Triosephosphate Isomerase

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None.

AUTHOR'S CONTRIBUTIONS

Hoang Kim Tu Trinh drafted the study protocol, contributed to writing and supervised the whole process. Kieu-Minh Le performed experiments, data analysis and drafted the manuscript. Thanh Niem Vo Van performed the experiment and contributed to writing. Duy Le Pham, and Hieu Thao Nguyen recruited patients and contributed to writing. Minh Nguyet Tran Thi, Bao Yen Pham, Dinh Minh Pham conducted experiments for protein identification and contributed to writing. All authors read and approved the final manuscript.

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AVAILABILITY OF DATA AND MATERIALS

Data and materials used and/or analyzed during the current study are available from the corresponding author on reasonable request.

ETHICS APPROVAL AND CONSENT TO PARTICIPATE

The University of Medicine and Pharmacy Ethics Committee at Ho Chi Minh City approved this study (no: 1105/HĐĐĐ-ĐHYD, on 9th November 2023).

CONSENT FOR PUBLICATION

Not applicable.

COMPETING INTERESTS

The authors declare that they have no competing interests.

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