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# Climate-driven changes to taste and aroma determining metabolites in an economically valuable portunid (*Portunus armatus*) have implications for future harvesting

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The effects of climate change on the distribution and biology of fisheries species have received substantial attention, but quantitative assessments of changes to taste and aroma determining compounds remain limited—despite sensory quality being a key driver of demand for most harvested species. Utilising the economically important blue swimmer crab (*Portunus armatus*), we tested the effects of temperature and salinity treatments aligned with near-future climate change projections on volatile and non-volatile sensory compounds that determine seafood flavour. Volatile compounds were analysed using solid phase microextraction and gas chromatography-mass spectrometry and non-volatiles were identified using liquid chromatography (mass spectrometry analysis). Multivariate analyses revealed that temperature, but not salinity, significantly affected the compositions of both volatile and non-volatile compounds in crab meat following a 30-day exposure period. Univariate analyses highlighted significant reductions in amino acids and amines associated with bitter and sweet organoleptic properties following exposure to elevated temperature. These results imply the potential for climate change to alter taste and aroma determining compounds in seafood, which could affect future harvesting priorities. Assessments of seafood sensory quality under climate change can produce valuable information to help predict shifts in fishing effort for harvested species that form the basis of important global fisheries.

## KEYWORDS

climate change, non-volatile, *Portunus armatus*, seafood flavour, volatile

## Introduction

The impacts of climate change on harvested marine resources are evident throughout the global ocean (Weatherdon et al., 2016; Free et al., 2019). Climate-driven changes to species geographic distributions (Pinsky et al., 2019; Gervais et al., 2021), phenology (Poloczanska et al., 2013) and temporal persistence within fishing regions (Champion et al., 2019; Rogers et al., 2019) presently challenge the sustainable harvest of seafood and the economic viability of fisheries (Sumaila et al., 2011). In addition to impacting the sustainable and profitable harvest of marine resources, studies suggest the environmental effects of climate change can also alter the quality of seafood, including nutritional (e.g. protein and fatty acids) and sensory properties (e.g. metabolites that contribute to flavour) (Dupont et al., 2014; Tate et al., 2017; Ab Lah et al., 2018; Maire et al., 2021). Given the large and growing global dependence on seafood (Gerland et al., 2014; FAO, 2020), understanding the effects of climate change on seafood quality is a priority for climate-impact research that intersects biological, economic and social issues.

The health benefits that humans derive from consuming bioavailable nutrients contained within seafood are immense (Lund, 2013). For example, seafood accounts for approximately 17% of the total animal protein consumed globally (FAO, 2020). Further, micronutrients (e.g. vitamin A, iron, calcium and zinc) essential for cognitive function and hormone regulation are highly concentrated in seafood, which is particularly important for many developing coastal regions that rely on ocean harvests (Hicks et al., 2019). Given the provision of nutrients through seafood is integral to the future of human health globally (Golden et al., 2021), it is concerning that some studies have noted the nutritional quality of seafood may be vulnerable to climate change (Valles-Regino et al., 2015; Lemasson et al., 2019; Alma et al., 2020; Lan et al., 2020). Also concerning is the potential for seafood sensory quality (determined by attributes such as taste, smell and texture) to be affected by climate-driven environmental change, considering sensory attributes contribute to consumer choices regarding seafood consumption and, therefore, the derived health benefits.

Despite the crucial role that sensory quality plays in seafood consumer appeal, studies testing the effects of climate-driven environmental change on seafood flavour remain limited. Of those that have, results vary among the species studied, environmental factors tested and exposure durations assessed. For example, simulated ocean acidification significantly reduced the taste and appearance of northern shrimp (*Pandalus borealis*), with individuals exposed to lowered pH more likely to score lower in sensory quality assessments than those exposed to higher pH (Dupont et al., 2014). In the context of climate-driven extreme events, simulated marine heatwaves were found to significantly reduce the moisture content of the harvested

gastropod *Turbo militaris*, which could compromise the desirable meat texture (Mamo et al., 2019). Conversely, exposure to climate-driven environmental change can improve the sensory quality of seafood. For example, Ran et al. (2019) found that volatile compounds pertaining to the aroma of both raw and cooked Chinese razor clams (*Simonovacula constricta*) significantly improved following exposure to elevated salinity. There is also evidence for no detected effects of simulated climate change on seafood sensory properties, with the appearance, aroma and taste of the Pacific oyster (*Magallana gigas*) not significantly affected following exposure to ocean acidification and warming treatments (Lemasson et al., 2017). Collectively, these examples demonstrate that, for some seafood species, climate-driven environmental change has been found to have a various of effects on the sensory properties of seafood.

Volatile and non-volatile compounds underpin food sensory quality and represent valuable responses to assess when testing for environmentally induced changes to seafood aroma and taste (Wang et al., 2004; Sarower et al., 2012; Liu et al., 2018; Ran et al., 2019; Duan et al., 2021). Volatile organic compounds, comprised mainly of aldehydes, ketones, and alcohols, occur in gaseous form at room temperature and are therefore key determinates of seafood aroma (Frank et al., 2009). Non-volatile compounds are associated with relatively low vapour pressures and do not similarly evaporate. Therefore, non-volatile metabolites, such as free amino acids (FAAs), primarily determine the taste of seafood (Chen and Zhang, 2007; Liu et al., 2018; Zhang et al., 2019). When taste receptors sense non-volatile compounds, the combination and concentration of these metabolites determine the tastes associated with seafood, such as umami, sweetness, saltiness and bitterness (Chen and Zhang, 2007; Fuentes et al., 2010; Sarower et al., 2012). For example, FAAs play an important role in the umami taste of seafood, while sweetness (a desired trait of many harvested crustaceans) is determined by soluble sugars, including betaine and multiple AAs (e.g. threonine, serine, glycine, alanine, arginine and proline). In addition to being reliable indicators of seafood aroma and taste for quantifying any effects of climate change on seafood sensory quality, the analyses of volatile and non-volatile compounds also reduces limitations associated with sensory panel analyses, including access to adequately trained and calibrated panellists (Mabuchi et al., 2019).

Estuarine systems are vulnerable to the environmental effects of climate change (Scanes et al., 2020) due to their interconnectedness with regional oceanographic processes and terrestrial climate dynamics (Gillanders et al., 2011). Estuaries are crucial habitats that support the recruitment, growth and maturation of harvested marine resources (Beck et al., 2001). With >500 species globally, portunids are a key group of economically important species that recruit and mature within tropical and temperate estuaries (Kailola et al., 1993). In Australia, blue swimmer crabs (*Portunus armatus*) are the

most economically important portunid, supporting a national harvest (commercial and recreational) of 2500 t per annum in the lower reaches of saline estuaries throughout subtropical and temperate latitudes. Given the Australian distribution of *P. armatus* encompasses two marine climate change hotspots that are warming at rates within the top 10% of coastal ocean regions globally (Hobday and Pecl, 2014), this species is ideal for investigating any potential effects of climate-induced changes to sensory compounds responsible for seafood aroma and taste.

We aimed to test if the concentration of volatile and non-volatile compounds pertaining to seafood aroma and taste significantly differed in *P. armatus* exposed to temperature and salinity treatments that align with near-future climate change projections, relative to individuals exposed to present-day environmental conditions. By testing the responses of compounds that determine decapod sensory quality to environmental treatments aligned with projected environmental change, we contribute to an emerging understanding of the likely impacts of climate change on seafood consumer appeal.

## Material and methods

### *Portunus armatus* collection

*Portunus armatus* were collected from Wallis Lake, New South Wales (NSW; 32.271°S, 152.471°E) using commercial round traps following Broadhurst et al. (2017). One-hundred live individuals were transported in an aerated fish transporter from Wallis Lake to the National Marine Science Centre (NMSC), Coffs Harbour, Australia. Intermoult crabs (determined following Hay et al. [2005]) with no physical damage were randomly selected as experimental replicates, with carapace length (CL; nearest mm) and mass (nearest 0.01 g) measured immediately prior to their acclimation and experimental exposure period.

### Experimental design

Fifty-six polyvinyl chloride experimental tanks (15 l) were partitioned into an orthogonal combination of salinity (~35 and ~40 practical salinity units [PSU]) and temperature treatments (~21 and ~27°C;  $n = 14$  tanks per treatment; Figure 1). Each tank contained one crab (to avoid competitive interactions) acclimated to experimental treatment levels over four days prior to a 30-day experimental exposure period. Treatment levels were strategically selected to reflect present-day environmental conditions experienced by *P. armatus* throughout their temperate Australian distribution and conditions predicted under near-future climate change (Gillanders et al., 2011; Hobday and Lough, 2011). Each

experimental treatment contained an approximate 1:1 sex ratio. This experimental design was previously utilised to test for the effects of climate change on the nutritional quality (e.g. protein, lipids, and fatty acids) of *P. armatus* (Champion et al., 2020).

### Experimental maintenance

Flow-through seawater from the adjacent ocean was supplied directly to the experimental system and adjusted to treatment levels in four separate 400 l header tanks prior to entering experimental tanks at a rate of 0.1 l min<sup>-1</sup>. Water temperature was controlled by heater-chiller units (Aquahort Ltd., Omana Beach, New Zealand) in header tanks and temperature stabilising water baths that housed individual experimental tanks. Ambient salinity from incoming seawater was stable during the experiment (i.e. ~35 PSU; Figure 1). Elevated salinity representative of future estuarine conditions was achieved by supplying hypersaline seawater (created by dissolving aquarium-grade crystallised sea salt at 150 g l<sup>-1</sup> [Cheetham Salt, South Australia]) from a sub-header tank into the primary header tanks that serviced relevant treatments.

During a one-week trial of the experimental system without live animals, heater-chiller units and the flow of hypersaline seawater into primary header tanks were adjusted and monitored to confirm stable treatment levels. Temperature and salinity within each experimental replicate were measured once every two days during the experiment using a TP20 digital thermometer (ThermPro, Toronto, Canada) and Vital Sine<sup>TM</sup> SR6 refractometer (Apopka, Florida, USA), respectively (Figure 1). The experiment took place in an enclosed room with a 12:12 h photoperiod. Throughout the experiment, *P. armatus* were fed school prawns (*Metapenaeus macleayi*) as per Uhlmann et al. (2009), with faeces routinely sieved from experimental tanks 14 h after feeding. The status of each crab (i.e. dead or alive) was monitored daily, after which individuals were euthanised by cutting along the ventral longitudinal midline, and the edible abdomen tissue (~20 g) was snap-frozen prior to volatile and non-volatile metabolite analyses.

### Volatile metabolite analyses

Solid phase microextraction (SPME) and gas chromatography-mass spectrometry (GC-MS) were used to analyse volatile compounds from experimental replicates. Crab meat samples (~10 g) were thawed and homogenised using an Ultra-Turrax (IKA-works, Malaysia). A subsample (2 g) was homogenised with water at a ratio of 1:2 in a 20 ml glass headspace vial, and the slurry (3 g) was transferred into headspace glass vials with an internal standard (4-methyl-1-pentanol; 0.5 µg g<sup>-1</sup>). Headspace extraction of volatiles was conducted at 40°C for 60 min and desorbed into the GC

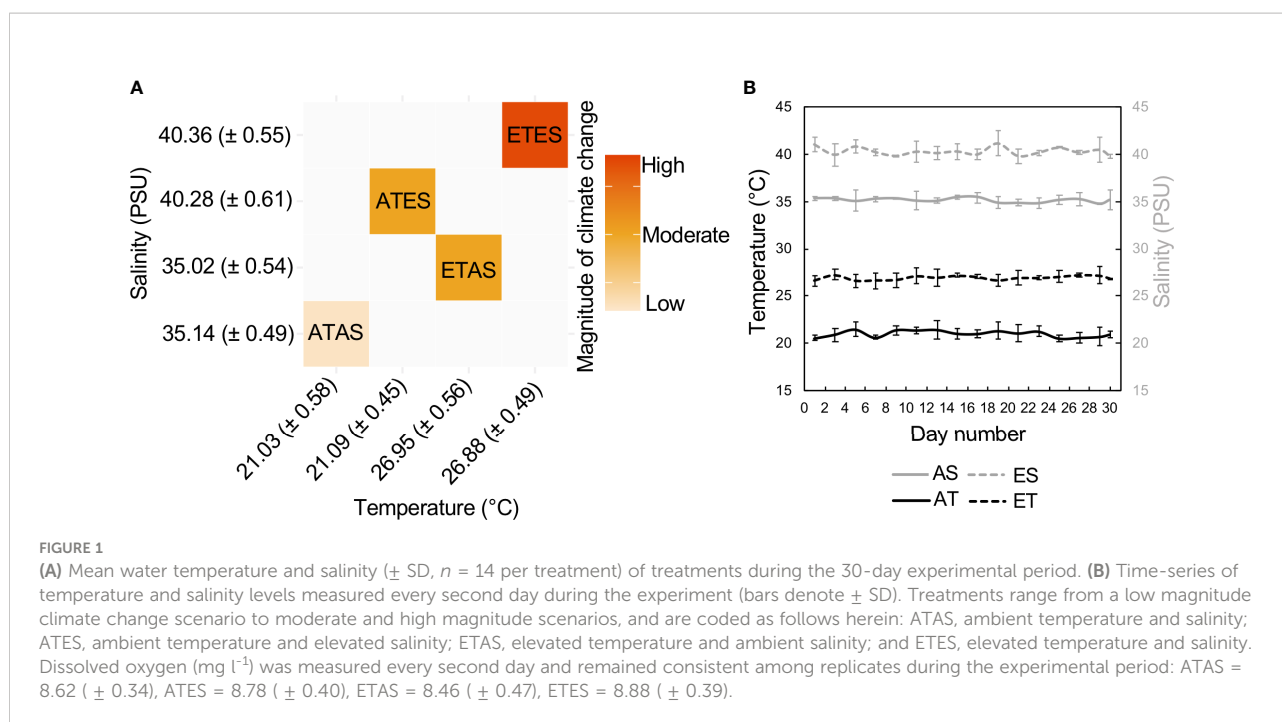


FIGURE 1

(A) Mean water temperature and salinity ( $\pm$  SD,  $n = 14$  per treatment) of treatments during the 30-day experimental period. (B) Time-series of temperature and salinity levels measured every second day during the experiment (bars denote  $\pm$  SD). Treatments range from a low magnitude climate change scenario to moderate and high magnitude scenarios, and are coded as follows herein: ATAS, ambient temperature and salinity; ATEs, ambient temperature and elevated salinity; ETAS, elevated temperature and ambient salinity; and ETES, elevated temperature and salinity. Dissolved oxygen ( $\text{mg l}^{-1}$ ) was measured every second day and remained consistent among replicates during the experimental period: ATAS =  $8.62 (\pm 0.34)$ , ATEs =  $8.78 (\pm 0.40)$ , ETAS =  $8.46 (\pm 0.47)$ , ETES =  $8.88 (\pm 0.39)$ .

injector in splitless mode at  $240^{\circ}\text{C}$  for 5 min. Divinylbenzene/carboxen/polydimethylsiloxane solid phase microextraction (SPME) fibres (23-gauge, 2 cm; Agilent Technologies, Bellefonte, USA) were used for the extraction.

Headspace analysis of samples was undertaken using SPME and GC-MS (Schimadzu QP-2010 Plus, Tokyo, Japan) and an auto-sampler (AOC-5000, Schimadzu). Compounds were separated on a Zebtron-WAX column (length 30 m; I.D. 0.25 mm and thickness 0.5  $\mu\text{m}$ ; Phenomenex, Lane Cove West, Australia). The carrier gas was helium (flow rate =  $1.04 \text{ ml min}^{-1}$ ). The initial column temperature was held at  $35^{\circ}\text{C}$  for 5 min, then increased to  $250^{\circ}\text{C}$  at  $5^{\circ}\text{C min}^{-1}$  and held for 5 min. Detection of volatiles was performed in electron ionisation mode (EI), 70 eV over a mass range  $m/z$  40–250. Kovats retention indices (RI), EI mass spectral library matches and, in most cases reference chemicals (RI), were used to identify volatile metabolites. Volatile data were semi-quantitated using the IS and expressed as  $\mu\text{g g}^{-1}$ .

## Non-volatile metabolite analyses

A sample of thawed homogenised crab meat (500 mg) was combined with 1 ml of 1:2 methanol:water solution and further homogenised for 10 min using a TissueLyser (Qiagen Retsch MM300, Haan, Germany). Samples were then centrifuged (Model 1-15, Sigma Laborzentrifugen, Germany) at 18,000 relative centrifugal force for 15 min. The supernatant was collected, and the residue re-extracted under the same

conditions. The supernatants were pooled and filtered using a nylon filter (Phenex, 0.2  $\mu\text{m}$ , Phenomenex, Lane Cove West, NSW, Australia).

Identification of non-volatile metabolites was achieved using liquid chromatography – mass spectrometry (LC-MS) analysis undertaken using a Dionex liquid chromatograph equipped with a pump, autosampler and heated column compartment (Ultimate 3000RS, Thermo Scientific, San Jose, CA, USA). Chromatographic separation of compounds was achieved using an Intrada Amino Acid column (length 150 mm x I.D. 3 mm; particle size 3  $\mu\text{m}$ ; Imtakt, Portland, OR, USA). The mobile phase consisted of acetonitrile/0.1% formic acid (solvent A) and 100 mM ammonium formate (solvent B). A constant flow rate of  $0.6 \text{ ml min}^{-1}$  was used with a gradient elution program, with an initial gradient of 14% B held constant for 3 minutes before being increased to 100% B for 10 minutes. At 12.5 minutes the gradient was returned to 14% B. An injection volume of 5  $\mu\text{l}$  was used.

Accurate mass measurement of non-volatile metabolites was conducted on a Q-Exactive Orbitrap LC-MS (Thermo Scientific, Scoresby, VIC, Australia) equipped with a heated electrospray ionization (H-ESI) source. The source conditions were as follows: spray voltage (positive ion 3.9 kV), sheath gas 60 (arbitrary units), auxiliary gas 10 (arbitrary units) and sweep gas 1 (arbitrary units), capillary temperature of  $350^{\circ}\text{C}$  and auxiliary gas heating temperature of  $400^{\circ}\text{C}$ . Mass spectra were acquired in positive ionisation mode and recorded over the mass range of 70 – 500  $m/z$  at 70 000 resolution (Xcalibur 4.3, Thermo Fisher Scientific, USA). A data-dependant workflow was used to

acquire MS/MS spectra using a stepped collision energy of 15, 30 and 100 eV in the HCD collision cell. The top five most abundant ions were selected for MS/MS scans with a dynamic exclusion of 10.0 seconds to prevent re-analysis. Compound Discoverer Ver 3.1 (Thermo Fisher Scientific, San Jose, CA, USA) and m/zCloud™ (<https://www.mzcloud.org/>) were used for identification of untargeted metabolites using a standard workflow template for food science. Aggregated computational toxicology resource, (ACToR, USA), FDA UNII – NLM (Maryland, USA), FooDB (The Food Database, Canada) and peptides databases were utilised for identification of metabolites. Only compounds with a high m/zCloud match (>60 with majority >80) were assigned an identity and subjected to statistical analyses.

## Statistical analyses

Curation of volatile and non-volatile compound data was conducted using Microsoft Excel and the ‘tidyverse’ package (Wickham et al., 2019) in R version 4.0.1 (R Core Team, 2020). The non-volatile dataset was further filtered for compounds that were in >75% of samples. Semi-quantitative volatile data were calculated using the IS and normalised prior to analysis. Non-volatile compound data were normalised prior to analysis using partial quotient normalisation (Dieterle et al., 2006). Permutational multivariate analysis of variance (PERMANOVA) testing for effects of temperature, salinity and their interaction on the composition of compounds was applied to volatile and non-volatile datasets in R using the ‘vegan’ package (Oksanen et al., 2020). These analyses utilised 999 permutations and similarity matrices based on Euclidean distance. A permutational distance-based analysis was selected over parametric analyses (e.g. multivariate analysis of variance) to ensure rank deficiency of models fitted would not cause unequal residuals between groups within variance-covariance matrices. Principal components analysis (PCA) was also applied to volatile and non-volatile multivariate data to assess for similarity in the composition of compounds among experimental treatments and identify compounds most correlated with principal components explaining the greatest variation in compounds among treatments.

Two-factor analyses of variance (ANOVA) were applied to test for effects of temperature, salinity and their interaction on the univariate responses of volatile and non-volatile compounds. All ANOVAs utilised type III sums of squares as appropriate for unbalanced data (Shaw and Mitchell-Olds, 1993; Queen et al., 2002), which resulted from crab mortalities during the experiment. Tukey’s HSD *post-hoc* pairwise tests were used to separate significant treatment-level effects. Shapiro-Wilk and Levene tests were applied to confirm that normalised volatile and non-volatile response variables satisfied assumptions of normality and homogeneity of variance. Finally, a binomial generalised linear model (GLM) using a logit link function was

applied in R using the ‘MASS’ package (Ripley and Venables, 2002) to test if the proportion of crab mortalities recorded throughout the experiment was significantly related to treatment.

## Results

### *Portunus armatus* mortality

Mortalities were between 21.4 and 28.6% among treatments during the experiment (mortalities per treatment = ATAS: 3, ATEs: 3, ETAS: 4, and ETES: 3). Neither temperature, salinity nor their interaction significantly affected mortalities (GLM,  $p > 0.05$ ).

### Volatile compounds

A total of 48 volatile compounds were identified and semi-quantified by the SPME method (Table 1; Figure 2A). Permutational multivariate analysis revealed the composition of volatile compounds in *P. armatus* meat significantly differed between crabs exposed to 21 and 27°C for 30 days (pseudo- $F = 1.97$ ,  $p = 0.02$ ; Figure 2B). However, no significant differences in the composition of volatile compounds were identified between crabs exposed to ambient (35 PSU) or elevated (40 PSU) salinities (pseudo- $F = 0.67$ ,  $p = 0.86$ ; Figure 2C), nor was there a significant interaction between temperature and salinity (at the treatment levels investigated) on the composition of volatile compounds in *P. armatus* meat (pseudo- $F = 1.05$ ,  $p = 0.37$ ).

Principal components analysis identified that 24.4 and 13.2% of variation in volatile compounds among treatments was explained by principal components 1 (PC1) and 2 (PC2), respectively, with PC1 and PC2 separating the composition of volatile compounds in crabs exposed to ambient and elevated temperatures (Figure 2B). The volatile compounds most positively correlated with PC1 were 1-hexanol (loading = 0.23) and 1-octanol (loading = 0.23). Dimethyl disulfide (loading = 0.20) and benzaldehyde (loading = -0.22) were most positively and negatively correlated with PC2, highlighting that differences in the concentration of these compounds among treatments primarily drove the overarching responses in the multivariate data.

Of the 48 volatile compounds identified and semi-quantified, 12 (25%) were significantly affected by temperature ( $p < 0.05$ ; Table 1). Trimethylamine, methanethiol, 2-butanone, methyl butanoate and 2-octanone were significantly lower following exposure to elevated temperature (Table 1). Conversely, hexanal, 6-methyl-5-hepten-2-one, 2-acetyl-1-pyrroline, 2-decanone, benzaldehyde, benzyl alcohol, benzothiazole were significantly higher in the meat of crabs exposed to elevated temperature. Elevated salinity did not significantly affect the concentration of

TABLE 1 Summary of volatile compounds ( $n = 48$ ) identified in the meat of blue swimmer crabs (*Portunus armatus*) exposed to elevated temperature and salinity treatments for 30 days.

Compound Identity	LRI <sup>a</sup>	m/z <sup>b</sup>	Mean ( $\pm$ SD) concentration ( $\mu\text{g g}^{-1}$ )				Temperature		Salinity		Temperature $\times$ Salinity	
			ATAS	ATES	ETAS	ETES	F-ratio (d.f.)	p value	F-ratio (d.f.)	p value	F-ratio (d.f.)	p value
Trimethylamine	<700	59	89.0 (56.9)	97.4 (22.3)	81.8 (28.7)	56.6 (30.3)	4.70 (1, 39)	<b>0.036</b>	0.50 (1,39)	0.487	2.18 (1,39)	0.147
Methanethiol	<700	48	1.9 (1.8)	5.5 (5.4)	1.5 (1.2)	1.4 (1.3)	5.76 (1, 39)	<b>0.021</b>	3.70 (1, 39)	0.062	3.62 (1, 39)	0.065
Carbon disulfide	704	76	168.9 (116.9)	144.5 (117.6)	187.5 (122.0)	160.3 (206.5)	0.14 (1, 39)	0.714	0.36 (1, 39)	0.567	0.01 (1, 39)	0.973
Dimethyl sulfide	720	62	1.2 (1.2)	0.8 (1.3)	0.7 (1.4)	0.3 (0.7)	2.08 (1, 39)	0.157	1.00 (1, 39)	0.324	0.01 (1, 39)	0.979
Acetone	816	43	77.9 (38.1)	90.7 (35.8)	89.0 (27.8)	83.9 (61.9)	0.02 (1, 39)	0.879	0.09 (1, 39)	0.763	0.47 (1, 39)	0.501
2-Butanone	886	43	48.7 (14.3)	56.1 (13.0)	39.7 (5.5)	41.1 (9.7)	12.14 (1, 39)	<b>0.001</b>	1.72 (1, 39)	0.197	0.77 (1, 39)	0.386
Ethanol	944	45	5.0 (7.1)	16.0 (31.3)	6.2 (9.1)	15.1 (23.1)	0.01 (1, 39)	0.951	2.53 (1, 39)	0.12	0.03 (1, 39)	0.862
Methyl butanoate	980	74	29.8 (17.1)	43.7 (23.4)	21.2 (15.5)	29.1 (14.7)	4.28 (1, 39)	<b>0.045</b>	3.93 (1, 39)	0.054	0.29 (1, 39)	0.590
Hexanal	1078	56	15.2 (9.4)	12.7 (7.8)	34.2 (28.1)	30.1 (21.1)	10.59 (1, 39)	<b>0.002</b>	0.38 (1, 39)	0.56	0.02 (1, 39)	0.888
Methyl pentanoate	1080	74	61.4 (87.0)	53.9 (37.4)	21.5 (18.7)	42.1 (46.9)	2.35 (1, 39)	0.134	0.14 (1, 39)	0.712	0.72 (1, 39)	0.401
Dimethyl disulfide	1084	94	2.8 (2.0)	6.7 (12.5)	1.6 (1.8)	0.4 (1.0)	3.65 (1, 39)	0.063	0.54 (1, 39)	0.467	1.72 (1, 39)	0.198
1-Butanol	1162	56	179.2 (322.8)	30.1 (18.4)	51.1 (28.3)	29.1 (29.2)	1.66 (1, 39)	0.203	3.03 (1, 39)	0.092	1.60 (1, 39)	0.214
Heptanal	1194	70	7.2 (17.5)	1.2 (4.0)	3.5 (5.8)	4.6 (7.2)	0.01 (1, 39)	0.962	0.67 (1, 39)	0.42	1.34 (1, 39)	0.254
Pyridine	1200	79	1.2 (3.0)	1.3 (3.1)	1.4 (3.5)	11.9 (39.5)	0.84 (1, 39)	0.365	0.70 (1, 39)	0.409	0.71 (1, 39)	0.404
2-Heptanone	1204	58	60.3 (136.7)	30.6 (40.4)	19.5 (10.6)	40.3 (47.6)	0.41 (1, 39)	0.523	0.05 (1, 39)	0.829	1.18 (1, 39)	0.285
2-Octanone	1238	58	98.6 (165.8)	44.5 (63.9)	0.2 (0.5)	9.7 (24.0)	5.56 (1, 39)	<b>0.021</b>	0.60 (1, 39)	0.41	1.43 (1, 39)	0.258
1-Pentanol	1270	57	28.3 (25.0)	26.8 (11.5)	28.5 (14.0)	24.4 (9.6)	0.06 (1, 39)	0.811	0.32 (1, 39)	0.575	0.08 (1, 39)	0.788
p-Cymene	1280	119	11 (10.7)	8.6 (10.0)	8.6 (10.0)	11.1 (10.4)	0.01 (1, 39)	0.977	0.01 (1, 39)	0.991	0.63 (1, 39)	0.435
6-Methyl-5-hepten-2-one	1304	108	6.2 (3.1)	8.5 (2.6)	11.5 (6.7)	8.8 (3.8)	4.40 (1, 39)	<b>0.042</b>	0.92 (1, 39)	0.918	3.71 (1, 39)	0.062
3-Hydroxy-2-butanone	1304	88	162.2 (290.2)	52.4 (79.8)	10.3 (32.7)	112.9 (227.8)	0.55 (1, 39)	0.464	0.01 (1, 39)	0.916	3.30 (1, 39)	0.077
(E)-2-heptenal	1320	83	117.9 (67.4)	150.9 (67.0)	94.7 (63.1)	99.0 (52.6)	3.82 (1, 39)	0.058	0.99 (1, 39)	0.327	0.56 (1, 39)	0.457
Octanal	1328	56	7.7 (9.5)	3.8 (8.5)	8.3 (4.5)	8.4 (6.5)	1.33 (1, 39)	0.257	0.71 (1, 39)	0.406	0.77 (1, 39)	0.389
2,3-Octanedione	1340	99	11.2 (17.6)	8.0 (13.5)	6.8 (13.5)	6.0 (6.2)	0.69 (1, 39)	0.411	0.27 (1, 39)	0.603	0.10 (1, 39)	0.752
1-Hexanol	1364	56	76.4 (76.4)	60.8 (43.6)	51.9 (34.4)	58.4 (28.6)	0.79 (1, 39)	0.384	0.10 (1, 39)	0.751	0.54 (1, 39)	0.470
2-Acetyl-1-pyrroline	1366	111	4.1 (10.6)	13.7 (17.4)	32.2 (27.3)	10.5 (13.1)	4.80 (1, 39)	<b>0.035</b>	1.06 (1, 39)	0.311	8.17 (1, 39)	<b>0.007</b>
Nonanal	1378	56	47.5 (43.9)	37.5 (16.0)	61.7 (83.1)	41.6 (33.2)	0.33 (1, 39)	0.568	0.99 (1, 39)	0.327	0.11 (1, 39)	0.741
2-Nonanone	1405	58	76.8 (59.4)	135.3 (219.5)	126.9 (120.7)	153.3 (120.3)	0.64 (1, 39)	0.43	0.98 (1, 39)	0.331	0.13 (1, 39)	0.714
(E)-2-octenal	1432	83	2.9 (5.3)	1.4 (2.0)	1.7 (2.3)	2.4 (2.6)	0.01 (1, 39)	0.965	0.21 (1, 39)	0.650	1.61 (1, 39)	0.289
Methional	1448	104	0.7 (1.6)	0.6 (1.1)	0.8 (1.1)	0.8 (1.3)	0.13 (1, 39)	0.722	0.01 (1, 39)	0.997	0.03 (1, 39)	0.870
1-Octen-3-ol	1450	57	352.0 (436.5)	241.9 (265.0)	140.6 (87.6)	204.1 (137.7)	2.03 (1, 39)	0.161	0.09 (1, 39)	0.77	1.01 (1, 39)	0.320
1-Heptanol	1460	70	37.4 (53.9)	25.2 (14.8)	23.0 (16.2)	18.4 (18.6)	1.32 (1, 39)	0.26	0.81 (1, 39)	0.375	0.16 (1, 39)	0.690
Acetic acid	1460	60	58.1 (144.0)	21.9 (44.9)	7.3 (12.3)	90.2 (188.9)	0.08 (1, 39)	0.776	0.34 (1, 39)	0.562	2.54 (1, 39)	0.119
Furfural	1472	95	1.2 (1.5)	0.9 (1.5)	1.7 (2.3)	1.4 (1.8)	0.87 (1, 39)	0.357	0.26 (1, 39)	0.617	0.01 (1, 39)	0.998
2-Ethylhexanol	1480	57	148.6 (60.9)	159.3 (40.2)	245.5 (259.9)	213.5 (136.5)	2.75 (1, 39)	0.104	0.05 (1, 39)	0.823	0.22 (1, 39)	0.637
2-Decanone	1490	58	9.8 (9.1)	12.1 (11.6)	17.4 (16.2)	23.2 (20.3)	4.32 (1, 39)	<b>0.044</b>	0.79 (1, 39)	0.381	0.14 (1, 39)	0.709
Benzaldehyde	1534	105	25.3 (4.7)	29.6 (8.5)	48.7 (36.3)	43.5 (30.8)	6.47 (1, 39)	<b>0.015</b>	0.01 (1, 39)	0.968	0.42 (1, 39)	0.519

(Continued)

TABLE 1 Continued

Compound Identity	LRI <sup>a</sup>	m/z <sup>b</sup>	Mean ( $\pm$ SD) concentration ( $\mu\text{g g}^{-1}$ )				Temperature		Salinity		Temperature $\times$ Salinity	
			ATAS	ATES	ETAS	ETES	F-ratio (d.f.)	p value	F-ratio (d.f.)	p value	F-ratio (d.f.)	p value
Methyl decanoate	1590	74	32.5 (37.3)	32.5 (16.16)	19.9 (13.0)	28.5 (25.1)	1.14 (1, 39)	0.293	0.30 (1, 39)	0.587	0.31 (1, 39)	0.580
Butyrolactone	1610	86	90.0 (35.4)	93.0 (47.0)	108.3 (37.8)	116.2 (33.8)	3.11 (1, 39)	0.086	0.21 (1, 39)	0.652	0.05 (1, 39)	0.835
1-Penten-3-ol	1658	57	33.9 (43.3)	10.6 (7.6)	14.7 (17.1)	14.3 (8.1)	1.12 (1, 39)	0.298	2.72 (1, 39)	0.107	2.27 (1, 39)	0.128
1-Octanol	1844	60	48.7 (55.7)	40.8 (19.2)	36.7 (35.1)	35.9 (24.0)	0.59 (1, 39)	0.448	0.16 (1, 39)	0.688	0.10 (1, 39)	0.750
Benzyl alcohol	1864	79	38.1 (29.4)	41.4 (33.3)	60.0 (41.1)	83.0 (58.6)	6.27 (1, 39)	<b>0.016</b>	1.01 (1, 39)	0.323	0.58 (1, 39)	0.452
Benzothiazole	1988	135	39.3 (5.6)	43.9 (10.4)	48.6 (11.3)	46.2 (7.9)	4.29 (1, 39)	<b>0.045</b>	0.17 (1, 39)	0.68	1.65 (1, 39)	0.207
Pantolactone	2030	71	1.5 (3.5)	0.4 (1.4)	2.9 (4.0)	2.6 (4.5)	1.81 (1, 39)	0.188	0.49 (1, 39)	0.484	0.05 (1, 39)	0.827
Octanoic acid	2044	60	74.6 (65.5)	55.7 (25.1)	55.3 (34.7)	70.5 (55.6)	0.04 (1, 39)	0.849	0.01 (1, 39)	0.932	1.51 (1, 39)	0.226
p-Cresol	2098	107	1.9 (1.8)	7.0 (9.0)	11.9 (22.5)	3.2 (2.3)	0.64 (1, 39)	0.427	0.21 (1, 39)	0.647	3.69 (1, 39)	0.062
Nonanoic acid	2178	60	16.1 (12.5)	21.8 (14.1)	32.6 (23.7)	19.5 (18.5)	1.63 (1, 39)	0.21	0.43 (1, 39)	0.517	3.11 (1, 39)	0.085
Decanoic acid	2270	60	12.0 (11.9)	11.9 (10.5)	13.6 (11.6)	19.0 (22.7)	0.92 (1, 39)	0.341	0.36 (1, 39)	0.575	0.36 (1, 39)	0.551
Indole	2470	117	132.0 (177.2)	266.9 (178.1)	179.8 (127.7)	105.4 (57.1)	1.77 (1, 39)	0.19	0.56 (1, 39)	0.460	5.65 (1, 39)	<b>0.022</b>

<sup>a</sup>Linear Retention Index.

<sup>b</sup>Mass to charge ratio.

Summary statistics for analyses testing for effects of temperature and salinity (and their interaction) on the suite of volatile compounds identified are presented, with significant (at  $\alpha = 0.05$ ) effects in bold. ATAS, ambient temperature and salinity; ATES, ambient temperature and elevated salinity; ETAS, elevated temperature and ambient salinity; and ETES, elevated temperature and salinity.

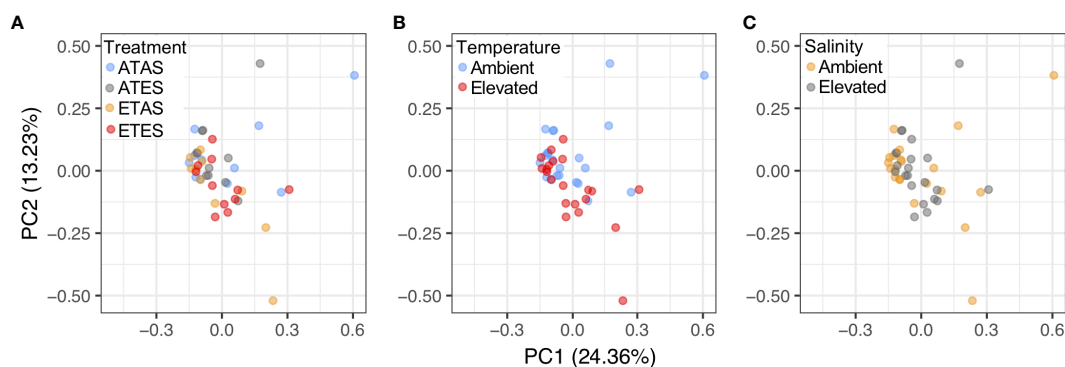


FIGURE 2

Principal components analysis biplots showing the similarity in composition of volatile compounds in blue swimmer crab (*Portunus armatus*) meat following experimentation (A) among all treatments, and between levels within the (B) temperature, and (C) salinity treatments. ATAS, ambient temperature and salinity; ATES, ambient temperature and elevated salinity; ETAS, elevated temperature and ambient salinity; and ETES, elevated temperature and salinity.

volatile compounds alone ( $p > 0.05$ ), however the interaction between temperature and salinity treatments significantly affected 2-acetyl-1-pyrroline and indole ( $p < 0.05$ ; Table 1). At ambient temperature, 2-acetyl-1-pyrroline and indole were significantly higher when salinity was elevated, with the opposite effect of elevated salinity observed at the elevated temperature level ( $p < 0.05$ ).

## Non-volatile compounds

Non-volatile compounds analysed using LC-MS/MS presented a total of 294 mass features in at least 75% of the samples and 104 of these were positively identified using ms/ms search of mzCloud (<https://doi.org/10.5281/zenodo.6775167>). Consistent with the results for volatile compounds, the

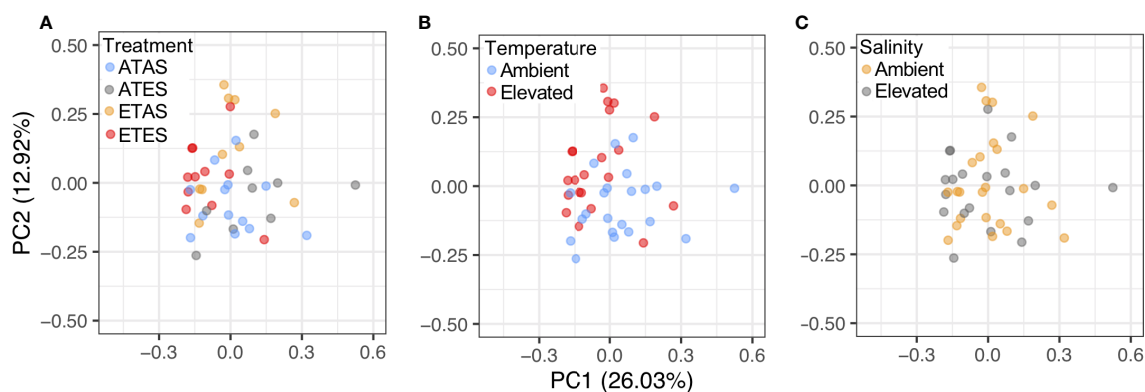


FIGURE 3

Principal components analysis biplots showing the similarity in composition of non-volatile compounds in blue swimmer crab (*Portunus armatus*) meat following experimentation (A) among all treatments, and between levels within the (B) temperature, and (C) salinity treatments. ATAS, ambient temperature and salinity; ATES, ambient temperature and elevated salinity; ETAS, elevated temperature and ambient salinity; and ETES, elevated temperature and salinity.

composition of non-volatile compounds in *P. armatus* meat significantly differed between crabs exposed to 21 and 27°C for a period of 30 days (pseudo- $F = 6.13$ ,  $p = 0.003$ ; Figure 3B). No significant differences in the composition of non-volatile compounds were found between crabs exposed to ambient or elevated salinity levels (pseudo- $F = 0.70$ ,  $p = 0.57$ ; Figure 2C) and there was no evidence of a significant interaction between temperature and salinity on the composition of non-volatile compounds (pseudo- $F = 1.02$ ,  $p = 0.38$ ). PC1 and PC2 explained 26.0 and 12.9% of variation in non-volatile compounds among treatments, respective (Figure 3). Non-volatile compounds most positively correlated with PC1 were lysine (loading = 0.15), histidine (loading = 0.15) and serine (loading = 0.15). Prolyl-glycine (loading = 0.18) was most positively correlated with PC2, while proline (loading =  $-0.08$ ) was most negatively correlated with this principal component.

Temperature, salinity or their interaction caused significant differences in the concentration of nineteen unique non-volatile compounds that were confidently identified following manual curation ( $p < 0.05$ ; Table 2). Of these, temperature alone was responsible for 83% of significant changes to the concentration of non-volatiles. The bitter tasting amino acids and amines tyrosine, iso-leucine, taurine and prolyl-leucine were present in significantly lower concentrations following exposure to the elevated temperature treatment, while the sweet tasting amino acid proline was also less abundant under the elevated temperature (Table 2).

A significant decrease in methionine, which is a major flavour precursor as well as an important amino acid, was found in the evaluated temperature treatment ( $F_{1,39} = 13.40$ ,  $p < 0.001$ ). A significant decrease in betaine was also identified under elevated temperature conditions ( $F_{1,39} = 16.62$ ,  $p < 0.001$ ). Betaine is involved in the methylation of cystathionin to form

methionine and in so doing forms dimethyl glycine, which was found to be significantly elevated under elevated temperature and salinity conditions (Table 2). Histidine was on average lower in the elevated temperature treatment and this coincided with a significant increase in its breakdown metabolite urocanic acid ( $F_{1,39} = 11.56$ ,  $p < 0.001$ ). Finally, amino acid degradation was also apparent in the elevated temperature treatment, as evidenced by a significant decrease in lysine and a significant increase in its metabolite pipercolinic acid (Table 2).

## Discussion

The effects of climate-driven environmental change on the sensory quality of seafood have considerable implications for consumer appeal and human health, but remain largely understudied. Here we tested the effects of simulated climate change on the composition of compounds that determine seafood aroma and flavour in the widely harvested *P. armatus*. We show the compositions of volatile and non-volatile compounds were significantly altered in crab meat following 30 days of exposure to elevated temperature. However, no overarching effects of elevated salinity on the compositions of volatile and non-volatile compounds were identified, nor was there a significant interaction between temperature and salinity on the composition of these compounds. This dominant effect of temperature, but not of salinity, on the composition of aroma- and taste-active compounds in crab meat suggests a relatively greater sensitivity of *P. armatus* flavour to changes in temperature than salinity. For example, significant reductions in amino acids and amines associated with bitter and sweet organoleptic properties (e.g. tyrosine, isoleucine, taurine and proline) were identified following exposure to elevated



temperature but not salinity. These results support the hypothesis that the warming of marine and estuarine environments expected under near-future climate change could alter the composition of compounds that determine the aroma and flavour of seafood. Given that seafood flavour is inextricably linked with consumer choices and market demand, climate-driven changes to aroma- and taste-active metabolites may manifest as changes in fishing effort for impacted species and, in extreme instances, the economic viability of fisheries.

## Changes to volatile and non-volatile compounds affect the flavour of *Portunus armatus* meat

The sensory attributes of numerous volatile compounds significantly affected by the experimental treatments applied here have been previously reported for some crustaceans (Matiella and Hsieh, 1990; Chung and Cadwallader, 1993; Chung, 1999; Sarnoski et al., 2010; Gu et al., 2013; Mall and Schieberle, 2016; Mall and Schieberle, 2017; Nanda et al., 2021). For example, trimethylamine and methanethiol are known to make important contributions to the aroma of crustaceans (Giri et al., 2010), with a decrease in these volatiles under elevated temperatures (as found here) likely to contribute to changes in the overall organoleptic quality of crab meat. Further, benzaldehyde produces a sweet odour through the degradation of unsaturated lipids, and a significant difference in the concentration of this compound is indicative of a different lipid composition and reduced meat sweetness in crabs exposed to ambient relative to elevated temperatures. 2-Acetyl-1-pyrroline has also been previously identified as an important odour-impacting compound in cooked crustaceans (Mall and Schieberle, 2016; Mall and Schieberle, 2017). The concentration of this compound within crab meat varied in response to the interaction between temperature and salinity treatments, with significantly higher concentrations found when salinity was elevated and the opposite effect observed at elevated temperature. Although we generally did not find significant interactive effects between temperature and salinity on odour-impacting compounds, this result demonstrates that interactive environmental effects can make important contributions to the aroma of seafood.

In addition to volatile odour-impacting compounds, the significant effect of temperature on the composition of non-volatile taste-active compounds further implies changes to the flavour of *P. armatus* meat under ocean warming. Betaine is a modified amino acid known to make important contributions to sweetness and umami flavours of crustacean meat (Sarower et al., 2012; Liu et al., 2018) and was significantly lower in *P. armatus* exposed to elevated temperatures for 30 days. Given this compound is commonly associated with high taste activity values (indicative of compounds that make substantial

contributions to food taste; Meyer et al., 2016), the reduction of betaine observed here may be a primary mechanism contributing to changes in seafood taste under climate change. Further, lysine is also known to influence sweet and bitter attributes associated with crustacean meat and products (e.g. sauces), and its reduction here could manifest as relatively bitter *P. armatus* meat under ocean warming. (Liu et al., 2019). Lysine is also a building block for protein and, unlike other amino acids, cannot be produced by the human body and must be consumed. Consequently, reductions in this amino acid under elevated temperature might also reduce the nutritional quality of *P. armatus* meat.

Linking changes in individual volatiles and non-volatiles to specific changes in sensory attributes (e.g. sweetness, bitterness, umami) is complicated by the multivariate nature of metabolite data. While our findings support the potential for near-future climate change to alter the composition of aroma- and taste-active compounds in seafood, confirmation of such changes requires sensory panel analyses. For example, Lemasson et al. (2017) utilised an expert panel to assess the appearance, aroma, taste and overall acceptability of *M. gigas* following exposure to elevated temperature and ocean acidification, finding non-significant trends towards improved sensory qualities under acidification and warming scenarios. Furthermore, Dupont et al. (2014) also used an expert panel to identify that the appearance and taste, but not texture, of *P. borealis* is likely to significantly decline in response to ocean acidification. However, studies integrating analyses of aroma- and taste-active compounds with formal assessments of seafood appearance, aroma and taste by trained assessors are required to provide a mechanistic understanding of the impacts of near-future climate change on the sensory quality of seafood. A recent study by Hsieh et al. (2021) found that exposing tiger shrimp (*Penaeus monodon*) to lowered pH reduced production of the amino acids glutamate and aspartic acid, which provide a umami flavour in seafood. These changes in amino acid production were subsequently supported by a blind sensory quality test, where *P. monodon* that had been exposed to lower pH received lower scores for flavour, appearance and texture. Further integrating sensory panel assessments with analyses of aroma- and taste-active compounds is an important future direction for fisheries climate-impact research and a logical next step for understanding how multivariate changes in *P. armatus* sensory metabolites manifest in changes to flavour attributes.

## Resilience of *Portunus armatus* flavour active compounds to elevated salinity

An important finding of this study was the apparent resilience of volatile and non-volatile compounds in *P. armatus* meat to altered salinity, which is consistent with

TABLE 2 Non-volatile compounds ( $n = 19$ ) positively identified in the meat of blue swimmer crabs (*Portunus armatus*) that were significantly ( $\alpha = 0.05$ ) affected by temperature or salinity treatments (or their interaction) after 30 days of experimental exposure.

Compound Identity	RT <sup>a</sup> (min)	ms/ms match	Mean ( $\pm$ SD) normalised peak area				Temperature		Salinity		Temperature $\times$ Salinity	
			ATAS	ATES	ETAS	ETES	F-ratio (d.f.)	p value	F-ratio (d.f.)	p value	F-ratio (d.f.)	p value
Taurine	7.53	99.8	242.5 (45.8)	222.4 (47.3)	183.3 (59.8)	199.0 (42.2)	8.49 (1, 39)	<b>0.006</b>	0.02 (1,39)	0.897	1.44 (1,39)	0.237
Proline	7.34	99.4	6297.9 (2066.0)	5879.4 (1511.7)	3335.8 (1831.8)	5153.7 (2372.8)	9.78 (1, 39)	<b>0.003</b>	1.40 (1, 39)	0.243	3.40 (1, 39)	0.072
Serine	8.22	99.3	19.9 (5.0)	26.1 (10.6)	23.2 (10.8)	17.1 (8.6)	0.73 (1, 39)	0.398	0.01 (1, 39)	0.981	5.21 (1, 39)	<b>0.028</b>
Carnitine	7.98	98.5	2725.8 (715.3)	2968.2 (1223.3)	3394.4 (695.0)	3758.0 (935.5)	7.83 (1, 39)	<b>0.001</b>	1.26 (1, 39)	0.268	0.05 (1, 39)	0.824
Iso-leucine	7.00	98.0	1143.4 (338.0)	1226.9 (258.6)	917.8 (308.2)	805.8 (401.6)	9.82 (1, 39)	<b>0.003</b>	0.02 (1, 39)	0.877	0.93 (1, 39)	0.342
Methionine	7.21	97.3	546.9 (201.4)	576.8 (192.7)	374.4 (139.8)	330.9 (205.2)	13.40 (1, 39)	<b>0.001</b>	0.02 (1, 39)	0.897	0.42 (1, 39)	0.522
Pipecolic acid	6.86	97.2	11.1 (2.3)	9.9 (2.9)	14.1 (2.0)	15.7 (5.2)	15.25 (1, 39)	<b>0.001</b>	0.04 (1, 39)	0.85	1.68 (1, 39)	0.204
Acetylcholine	7.79	95.4	992.9 (302.0)	1002.5 (354.6)	1525.3 (560.6)	1428.1 (516.5)	10.97 (1, 39)	<b>0.002</b>	0.12 (1, 39)	0.734	0.14 (1, 39)	0.716
Histidine	12.69	95.4	559.6 (94.7)	595.6 (162.2)	585.3 (225.8)	400.1 (146.0)	2.82 (1, 39)	0.101	2.43 (1, 39)	0.127	5.08 (1, 39)	<b>0.030</b>
Dimethyl glycine	7.22	91.8	25.0 (7.17)	20.4 (7.0)	36.6 (17.2)	25.6 (7.5)	6.26 (1, 39)	<b>0.017</b>	5.97 (1, 39)	<b>0.019</b>	1.00 (1, 39)	0.324
Betaine	6.76	91.1	41293.9 (8321.2)	35891.4 (6684.7)	30059.9 (5380.0)	32398.1 (3935.3)	16.63 (1, 39)	<b>0.001</b>	0.57 (1, 39)	0.455	3.97 (1, 39)	0.053
Citrulline	8.35	90.9	35.3 (27.8)	63.6 (88.4)	154.0 (171.6)	82.1 (48.7)	5.72 (1, 39)	<b>0.022</b>	0.57 (1, 39)	0.455	2.79 (1, 39)	0.103
Tryptophan	6.91	89.6	97.8 (42.0)	130.7 (57.4)	93.5 (58.4)	62.6 (38.6)	5.04 (1, 39)	<b>0.030</b>	0.01 (1, 39)	0.976	4.56 (1, 39)	<b>0.039</b>
Hypoxanthine	4.2	89.0	562.8 (118.9)	522.7 (96.1)	505.3 (88.6)	444.5 (92.8)	5.52 (1, 39)	<b>0.024</b>	2.72 (1, 39)	0.107	0.11 (1, 39)	0.738
Prolyl-leucine	7.48	88.6	21.2 (7.8)	23.5 (6.1)	13.6 (4.7)	11.7 (8.4)	18.55 (1, 39)	<b>0.001</b>	0.03 (1, 39)	0.875	0.93 (1, 39)	0.340
Tyrosine	7.28	86.9	127.4 (39.6)	135.6 (30.8)	102.4 (50.0)	80.7 (37.9)	9.85 (1, 39)	<b>0.003</b>	0.36 (1, 39)	0.553	1.40 (1, 39)	0.244
Prolyl-glycine	7.31	88.1	9.6 (3.4)	15.6 (9.1)	21.5 (20.6)	15.4 (6.3)	4.23 (1, 39)	<b>0.046</b>	0.03 (1, 39)	0.855	2.54 (1, 39)	0.120
Urocanic acid	2.61	79.0	21.2 (10.9)	19.9 (9.3)	44.1 (24.6)	30.5 (14.8)	11.56 (1, 39)	<b>0.001</b>	2.47 (1, 39)	0.124	1.53 (1, 39)	0.222
Lysine	13.27	78.6	579.9 (234.1)	693.1 (180.2)	438.8 (255.9)	368.5 (250.9)	9.92 (1, 39)	<b>0.003</b>	0.08 (1, 39)	0.779	1.66 (1, 39)	0.207

<sup>a</sup>Retention time.

Significant effects are bolded. ATAS, ambient temperature and salinity; ATES, ambient temperature and elevated salinity; ETAS, elevated temperature and ambient salinity; and ETES, elevated temperature and salinity.

previous research showing minimal effects of altered salinity on nutritional properties (e.g. protein and fatty acid composition) contained in *P. armatus* meat (Champion et al., 2020). Exposure to different salinities is known to affect crab tissue by altering the concentrations of free amino acids and other metabolites to maintain cells in an iso-osmotic condition and the null result here is contrary to other studies (Holt and Kinsey, 2002). For example, Chinese razor clams (*Sinonovacula*

*constricta*)—a burrowing bivalve occurring in soft sediment environments throughout eastern Asia—cultured under elevated salinity conditions (PSU = 23) had more abundant volatiles than those cultured under lower salinities (PSU = 13; Ran et al., 2019). Furthermore, lowered salinity evoked softer meat texture and weaker flavour characteristics, including umami taste, in green mud crabs (*Scylla paramamosain*; Luo et al., 2021).

It is possible the trend toward an effect of temperature, but not salinity, on compounds pertaining to the flavour of *P. armatus* was due to the selection of the treatment levels. Presently, temperatures of 27°C (i.e. the elevated temperature treatment) approximate, or exceed, the upper thermal limit experienced by *P. armatus* throughout its temperate eastern Australian distribution, particularly at the site of collection for our experiment (Fiebig, 2010). By comparison, salinity levels within estuarine habitats throughout this region vary considerably in response to drought and rainfall events, and are known to exceed the elevated salinity treatment level applied here (PSU = 40) Wallis Lake (Fiebig, 2010). Therefore, the physiology of *P. armatus* may be better adapted to cope with changes in salinity, relative to the changes in temperature, at the levels applied. Nevertheless, our experimental treatment levels were selected to align with projections of climate-driven environment change in eastern Australia (Gillanders et al., 2011; Hobday and Lough, 2011), and so, for *P. armatus*, future changes in temperature are likely to make greater contributions to altered sensory quality than changes in salinity.

We also acknowledge salinity levels in estuarine environments are projected to become increasingly variable under climate change, with elevated levels predicted during intense drought events, and reduced levels in response to increasing flooding events. While the magnitude of variation in salinities experienced in estuarine environments is expected to increase under climate change, here we have only considered the elevated salinity scenario likely to occur during periods of drought. Testing the effects of lower salinities is also important for fully addressing the effects of future salinity levels on aroma- and taste-active compounds pertaining to the flavour of *P. armatus* under climate change.

In addition to coherent choices of the upper bounds of environmental change, careful consideration of experimental durations is essential when testing the effects of climate change on seafood sensory properties. Predicted increases in the magnitude of short-term environmental extremes and the frequency of conditions that are outside species' tolerance ranges under climate change (Kerr, 2011) highlight the importance of short-term (i.e. weeks to months) experiments for quantifying species-specific responses (as applied here; Jentsch et al., 2007; Thompson et al., 2013). However, the physical effects of climate change on marine systems also include underlying shifts in mean environmental conditions. Selecting experimental treatment levels and durations that align with the projected effects of climate change for specific study systems and species are necessary to ensure results are indicative of likely future changes (Shalders et al., 2022). Moreover, incorporating the effects of climate change across trophic levels is an additional consideration for future research. Here *P. armatus* were *M. macleaya* that had not been exposed to experimental treatments. Thus, it remains unknown how the potential effects of environmental change on prey contribute to

changes in the composition of volatile and non-volatile compounds in *P. armatus* meat.

## Implications of climate-driven changes to seafood flavour

Seafood flavour is inextricably linked with consumer choices and market demand (Birch and Lawley, 2012; Murray et al., 2017). In Australia and Canada, seafood taste is a leading driver of appeal and a key determinant of the amount of seafood that humans consume (Christenson et al., 2017; Murray et al., 2017). By influencing the amounts and types of seafood that humans consume, flavour characteristics also contribute to the associated health benefits derived through seafood consumption (Christenson et al., 2017). In this way, climate-driven changes to the sensory quality of seafood have the potential to impact consumer preferences for harvested species, and also affect the nutritional benefits humans derive from seafood consumption.

Climate-change impacts may be positive or negative depending on the magnitude and direction of the effects of environmental change on seafood sensory properties. If the magnitude of climate-driven changes to seafood sensory properties are large enough, there is potential for the economic viability of fisheries to become impacted through reductions in the value and/or market demand for their products. However, such impacts will depend on the relative vulnerability of harvested species to climate change, which will be determined by relative physiological sensitivities and exposures (Foden et al., 2019). While experimental assessments are required to quantify the sensitivity of sensory properties in harvested species to environmental change, regions of the global ocean where species are most exposed to the effects of climate-driven ocean warming are already apparent (Hobday and Pecl, 2014; Pecl et al., 2014). It is within these regions, which include the Sea of Japan, south-eastern Australia and the Bering Sea among others, that seafood industries should be most aware of the potential for climate-driven changes to seafood aroma and taste.

## Data availability statement

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and accession number(s) can be found below: <https://doi.org/10.5281/zenodo.6775167>.

## Author contributions

CC, MB and MC conceptualised the research question and designed the experimental approach. DF, MT, KK and UP

undertook laboratory analysis of volatile and non-volatile metabolites. CC, DF and MT statistically analysed the data. CC and TS wrote the manuscript with input from all authors. All authors contributed to the article and approved the submitted version.

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## Conflict of interest

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