

Simultaneous Analysis of Purine and Pyrimidine Compounds Associated with the Freshness and Taste of Marine Foods

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Abstract A high-performance liquid chromatography-based method following the simultaneous analysis of 11 purine and pyrimidine compounds associated with freshness (*K* value for fish or *Xt* ratio for marine invertebrates) and umami taste in marine products was developed. This method, which separated 11 compounds in a 55-min cycle using a gradient of pH 3.8 phosphate buffer, acetonitrile, and water as the mobile phase, showed sufficient reproducibility and quantitative ability. Using this method, increases in the *K* value of an amberjack (*Seriola purpurascens*) and the *Xt* ratio of a squid (*Todarodes pacificus*) because of post-mortem deterioration were evaluated. Furthermore, high- and low-grade lavers (*Porphyra tenera*) known for their substantial guanosine 5'-monophosphate (an umami taste component) content, were compared using this method. Moreover, other nucleotide 5'-monophosphates, which are also umami components, from canned crab (the snow crab *Chionoecetes opilio*) and dried shiitake (*Lentinula edodes*), which is a mushroom containing a lot of guanosine 5'-monophosphate and was used as positive control for analysis by developed method, were successfully analyzed using this method.

Keywords HPLC · Marine products · Invertebrate · ATP decomposition · *K* value · Nucleotide 5'-monophosphate

Introduction

Marine products contain various extractable components, including organic acids, amino acids, and peptides. Particularly, purine and pyrimidine compounds are associated with the quality of marine products. In post-mortem marine products, adenosine 5'-triphosphate (ATP) gradually decomposes into adenosine 5'-diphosphate (ADP), adenosine 5'-monophosphate (AMP), inosine 5'-monophosphate (IMP), inosine (HxR), and hypoxanthine (Hx). Saito et al. (1959) used this ATP decomposition pathway to define the *K* value as an indicator of initial freshness in fish. The *K* value was defined using the equation below.

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

Marine invertebrates such as cephalopods (squids and octopi) and crustaceans (shrimp and crabs) have different pathways of ATP decomposition. Cephalopods have a decomposition pathway that proceeds via adenosine (AdR) rather than IMP (Arai and Saito 1961), and crustaceans have both the decomposition pathways proceeding via IMP and via AdR (Arai and Terasaki 1966). In crustaceans, Hx decomposes into xanthine (*Xt*) (Matsumoto and Yamanaka 1992). Therefore, other indicators are used when the freshness of these marine invertebrates is evaluated, such as the arranged *K* value, which replaces IMP with AdR and *Xt* ratio, as defined by the equation below (Yokoyama et al. 1994).

$$Xt \text{ ratio}(\%) = Xt / (\text{ATP} + \text{ADP} + \text{AMP} + \text{IMP} + \text{AdR} + \text{HxR} + \text{Hx} + Xt) \times 1000$$

Nucleotide 5'-monophosphates are important because they contribute to the umami taste of marine products. IMP and

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guanosine 5'-monophosphate (GMP) have umami tastes (Kuninaka 1960). AMP and cytidine 5'-monophosphate (CMP) have no umami taste by themselves, but when the disodium form of these phosphates coexists with glutamic acid, they synergistically enhance the umami taste of glutamic acid (Terasaki et al. 1965).

As mentioned above, purine and pyrimidine compounds have a great influence on the quality of marine products, and many methods for their analysis have been reported. With regard to the HPLC methods, different methods are reported according to analytical objects: methods for freshness levels of vertebrate (fish) (Lee et al. 1982; Özogul et al. 2000; Ryder 1985), methods for freshness of invertebrate (Yokoyama et al. 1994), and methods for umami taste (Hosokawa et al. 1990; Yamasaki 1994). When purine and pyrimidine compounds in marine products are analyzed to evaluate freshness and umami taste, the analysis requires a great deal of time and equipment because the analysis conditions need to be changed according to the analytical objects and samples.

In this study, we developed an HPLC method for the simultaneous analysis of 11 purine and pyrimidine compounds that are associated with the freshness and umami tastes of marine products. Using this method, we evaluated the

freshness of amberjack (vertebrate) and squid (invertebrate) as well as nucleotide 5'-monophosphate contents, which are umami taste components, of some foods.

Materials and Methods

Chemicals

Phosphoric acid, 70 % perchloric acid, potassium hydroxide, and disodium hydrogenphosphate dodecahydrate, which were JIS special grade, triethylamine, and anhydrous sodium dihydrogen phosphate, which were Wako special grade, and acetonitrile, which was HPLC grade, were obtained from Wako Pure Chemical Industries, Ltd. (Tokyo, Japan). CMP, guanine, Hx, Xt, IMP, HxR, AMP, ADP, AdR, ATP (Sigma Aldrich Japan, Ltd., Tokyo, Japan), and GMP (Wako Pure Chemical Industries, Ltd., Tokyo, Japan) were used as standard compounds.

Standard solutions of purine and pyrimidine compounds were prepared by dissolving the purine and pyrimidine compounds in 20-mM phosphoric acid buffer (pH 7.5). These solutions included $0.4281 \mu\text{mol mL}^{-1}$ CMP, $0.7559 \mu\text{mol mL}^{-1}$ guanine, $0.8601 \mu\text{mol mL}^{-1}$ Hx, $0.3809 \mu\text{mol mL}^{-1}$ Xt,

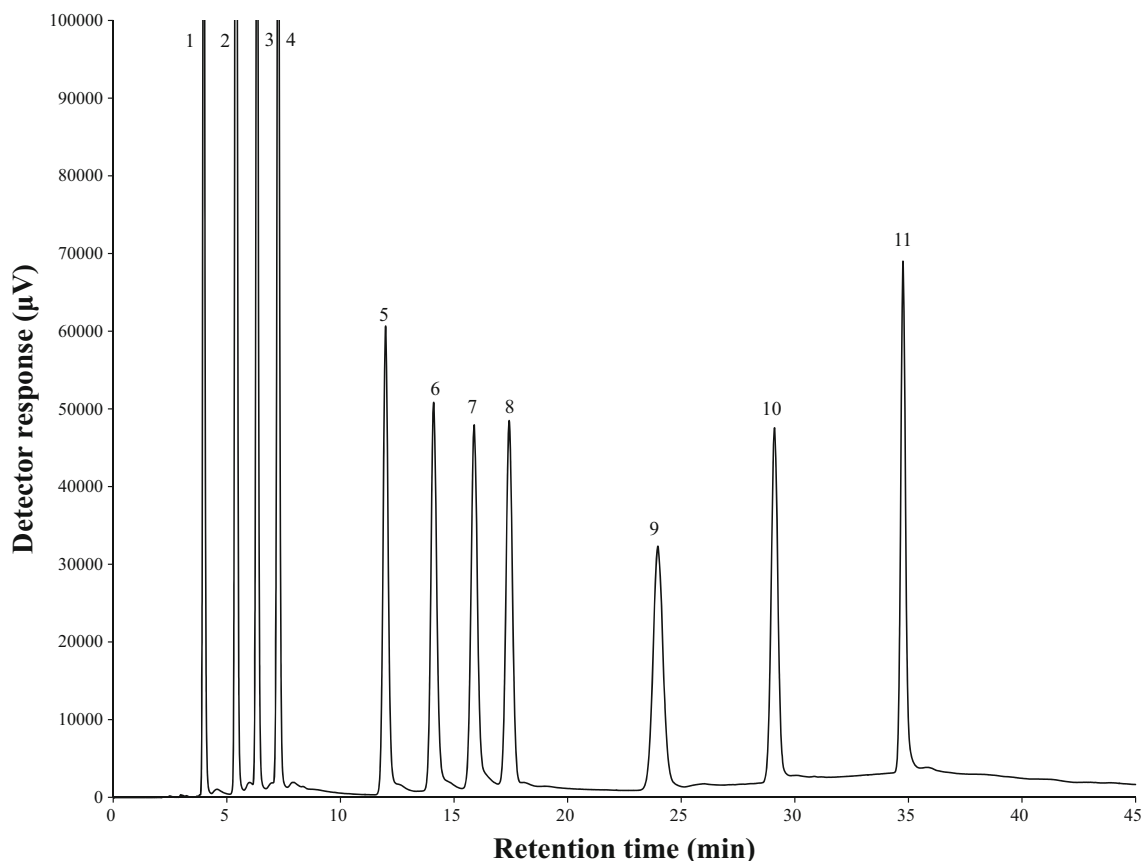


Fig. 1 HPLC chromatogram of purine and pyrimidine compounds. A number of each chromatogram represents each compound: 1, CMP; 2, guanine; 3, Hx; 4, Xt; 5, GMP; 6, IMP; 7, HxR; 8, AMP; 9, ADP; 10, AdR; 11, ATP

Table 1 Results from the evaluation of the precision and specificity of the LC method^a

	tR(min)	tR-RSD (%)	AREA	AREA-RSD(%)	K'	Rs	α
CMP	4.199	0.157	308013	0.897	0.614	4.801	–
Guanine	5.572	0.142	487477	0.877	1.142	7.515	1.860
Hx	6.515	0.147	335354	0.951	1.505	4.846	1.317
Xt	7.451	0.116	335458	1.072	1.865	4.350	1.239
GMP	12.585	0.167	287384	1.051	3.839	15.745	2.058
IMP	14.976	0.148	280130	1.444	4.758	5.213	1.240
HxR	17.293	0.119	277558	0.871	5.649	4.415	1.187
AMP	18.143	0.106	294108	0.848	5.976	1.552	1.058
ADP	26.590	0.094	294201	1.682	9.224	14.323	1.544
AdR	29.627	0.023	288020	2.266	10.391	5.305	1.127
ATP	35.698	0.148	320724	3.538	12.725	14.816	1.225

^aChromatographic parameters of purine and pyrimidine compounds with UV–VIS HPLC (detection, wavelength 270 nm), mixed purine and pyrimidine compound concentration, CMP 0.168 nmol, guanine 0.341 nmol, Hx 0.373 nmol, Xt 0.167 nmol, GMP 0.176 nmol, IMP 0.429 nmol, HxR 0.308 nmol, AMP 0.151 nmol, ADP 0.178 nmol, AdR 0.129 nmol, ATP 0.147 nmol per 20 μ L injection ($n=5$). *tR* retention time, *K'* capacity factor, *Rs* resolution factor, α separation factor

0.4415 μ mol mL⁻¹ GMP, 1.0293 μ mol mL⁻¹ IMP, 0.7728 μ mol mL⁻¹ HxR, 0.3820 μ mol mL⁻¹ AMP, 0.4406 μ mol mL⁻¹ ADP, 0.3320 μ mol mL⁻¹ AdR, and 0.3667 μ mol mL⁻¹ ATP.

HPLC System

Purine and pyrimidine compounds were analyzed using an HPLC system that included a PU-1580 pump, a DG-980-50 degasser, a CO-965 column oven (40 °C), a UV-970 UV–VIS detector (wavelength, 270 nm) (JASCO Corporation, Tokyo, Japan), a DMC675 Mixer (GLscience, Tokyo, Japan), and a 7725i Rheodyne-sampling injector connected to a 20- μ L sample loop (Rheodyne, Rohnert Park, CA, USA). This system was equipped with a CAPCELL PAK UG120 column (250 nm \times

4.6 mm, 5- μ m particle size, Shiseido, Tokyo, Japan) connected to a Guard-Pak guard column containing Nova Pak-C₁₈ (Waters, Milford, MA, USA). The mobile phase consisted of solutions A [water: triethylamine: phosphoric acid, 950:10:5 (v/v); pH 3.8] and B [water: acetonitrile, 90:10 (v/v)] were used. The gradient employed was 0–20 min, isocratic at 0 % B; 20–35 min, 0 % B to 35 % B; 35–45 min, isocratic at 35 % B. The flow rate was 1.0 mL/min. New samples were injected at intervals of 55 min. HPLC was conducted at room temperature.

Method Validation

Validation of the HPLC method included precision and specificity, linearity, range, and limit of detection and quantitation, recovery.

Table 2 Results from the evaluation of the linearity and range of the LC method

	Regression equation	Correlation factor	Linear range (nmol mL ⁻¹)	LOD (nmol mL ⁻¹)	LOQ (nmol mL ⁻¹)
CMP	$Y=2.95 \times 10^{-5} X^b - 1.20 \times 10^{-1}$	0.9999	0.53–8.41	1.9712	6.5707
Guanine	$Y=3.39 \times 10^{-5} X + 1.49 \times 10^{-2}$	1.0000	1.07–68.22	1.9987	6.6622
Hx	$Y=5.81 \times 10^{-5} X - 6.48 \times 10^{-1}$	0.9998	1.17–149.14	4.3692	14.5642
Xt	$Y=2.42 \times 10^{-5} X - 5.82 \times 10^{-2}$	1.0000	0.52–33.45	1.9600	6.5335
GMP	$Y=3.25 \times 10^{-5} X + 1.36 \times 10^{-2}$	1.0000	0.55–25.22	2.0637	6.8789
IMP	$Y=8.40 \times 10^{-5} X + 5.13 \times 10^{-1}$	0.9999	1.34–214.41	5.0253	16.7511
HxR	$Y=5.77 \times 10^{-5} X + 1.05 \times 10^{-1}$	1.0000	0.96–61.66	3.6128	12.0428
AMP	$Y=2.54 \times 10^{-5} X - 2.67 \times 10^{-2}$	0.9999	0.47–60.39	1.7692	5.8972
ADP	$Y=3.10 \times 10^{-5} X - 9.58 \times 10^{-2}$	1.0000	1.12–53.52	2.0908	6.9694
AdR	$Y=2.21 \times 10^{-5} X - 7.10 \times 10^{-2}$	1.0000	0.80–25.74	1.5080	5.0267
ATP	$Y=2.46 \times 10^{-5} X - 4.72 \times 10^{-1}$	0.9999	0.92–29.44	0.8624	2.8748

^aY=mass concentration (nmol mL⁻¹)

^bX=HPLC peak area

Table 3 Results from the evaluation of the recovery of the LC method

Percent	Recovery			
	Original ^a	Half of original	Quarter of original	Average
Add solution				
CMP	94.49	91.02	103.29	96.27
Guanine	92.31	90.16	115.61	99.36
Hx	97.27	92.38	113.77	101.14
Xt	92.17	86.31	86.42	88.30
GMP	89.52	87.94	101.21	92.89
IMP	107.37	104.93	242.97	151.76
HxR	91.80	91.65	101.58	95.01
AMP	92.03	74.09	86.72	84.28
ADP	92.66	89.88	108.40	96.98
AdR	95.67	84.47	91.18	90.44
ATP	93.31	99.66	120.59	104.52

^a Original mixed purine and pyrimidine compound solutions contained CMP (428.07 nmol mL⁻¹), guanine (755.89 nmol mL⁻¹), Hx (860.07 nmol mL⁻¹), Xt (380.86 nmol mL⁻¹), GMP (441.53 nmol mL⁻¹), IMP (1029.29 nmol mL⁻¹), HxR (772.76 nmol mL⁻¹), AMP (382.04 nmol mL⁻¹), ADP (440.57 nmol mL⁻¹), AdR (331.96 nmol mL⁻¹), and ATP (366.71 nmol mL⁻¹). 1-mL aliquot of each mixed standard solutions was added to 1 g of amberjack meat, which was then mixed and treated as described in the Experimental section. Recovery and RSD indicate the average recoveries for the three analyses (%)

For the precision and specificity tests, analytical solutions were prepared by diluting standard solution 50-fold 20 mM phosphate buffer (pH 7.5). Five separate 20- μ L aliquots of each analytical solution were injected and analyzed using HPLC. The retention time (tR), peak area, capacity factor (K'), resolution factor (Rs), and separation factor (α) were estimated using the results of these five replicate analyses. The

parameters were defined using the following standard formulas: $K'=(tR_n - tR_0) / tR_0$, $Rs=2(tR_{n+1} - tR_n) / (W_{n+1} + W_n)$, $\alpha=K'_{n+1}/K'_n$, where tR_n is retention time of each peak, tR₀ is the retention time of the first solvent peak (tR₀=2.60 min), W_n is the width of the peak at baseline. The subscripts n refer to the order of purine and pyrimidine compounds elution.

To test the linearity, range, limit of detection (LOD), and limit of quantification (LOQ), analytical solutions were prepared by diluting standard solutions in gradations from 2-fold to 800-fold with 20 mM phosphate buffer (pH 7.5). A 20- μ L aliquot of each solution was injected and analyzed using HPLC. The linearity, range, LOD, and LOQ were estimated using the results of these analyses. LOD was calculated as the concentration at which the purine and pyrimidine compounds peaks could be detected without any baseline noise disturbances (>3 times the baseline noise). LOQ was calculated as the concentration at which the analytes' responses were >10 times the baseline noise.

For the recovery tests, fortified samples were prepared by adding a 1-mL aliquot of standard solution (original), solution diluted twofold with 20 mM phosphate buffer (pH7.5) (half of original), and solution diluted fourfold with 20 mM phosphate buffer (pH7.5) (quarter of original) to the meat of an amberjack (*Seriola purpurascens*). According to the method of Hu et al. (2013), four pieces of amberjack muscle (1 g each) were weighed into separate 50-mL plastic centrifuge tubes, then a 1-mL aliquot of standard solution (original, half of original, quarter of original, 20-mM phosphate buffer) was added to each. Then, 10 mL of a 1-mol L⁻¹ perchloric acid solution was added to each tube. These samples were homogenized and left for 10 min at room temperature. The samples were centrifuged at 1500 g for 10 min, and the supernatants were transferred to 15-mL plastic test tubes. These sample solutions

Table 4 Purine and pyrimidine compounds contents^a and K value^b of amberjack of varying freshness during cooled storage at 5 °C

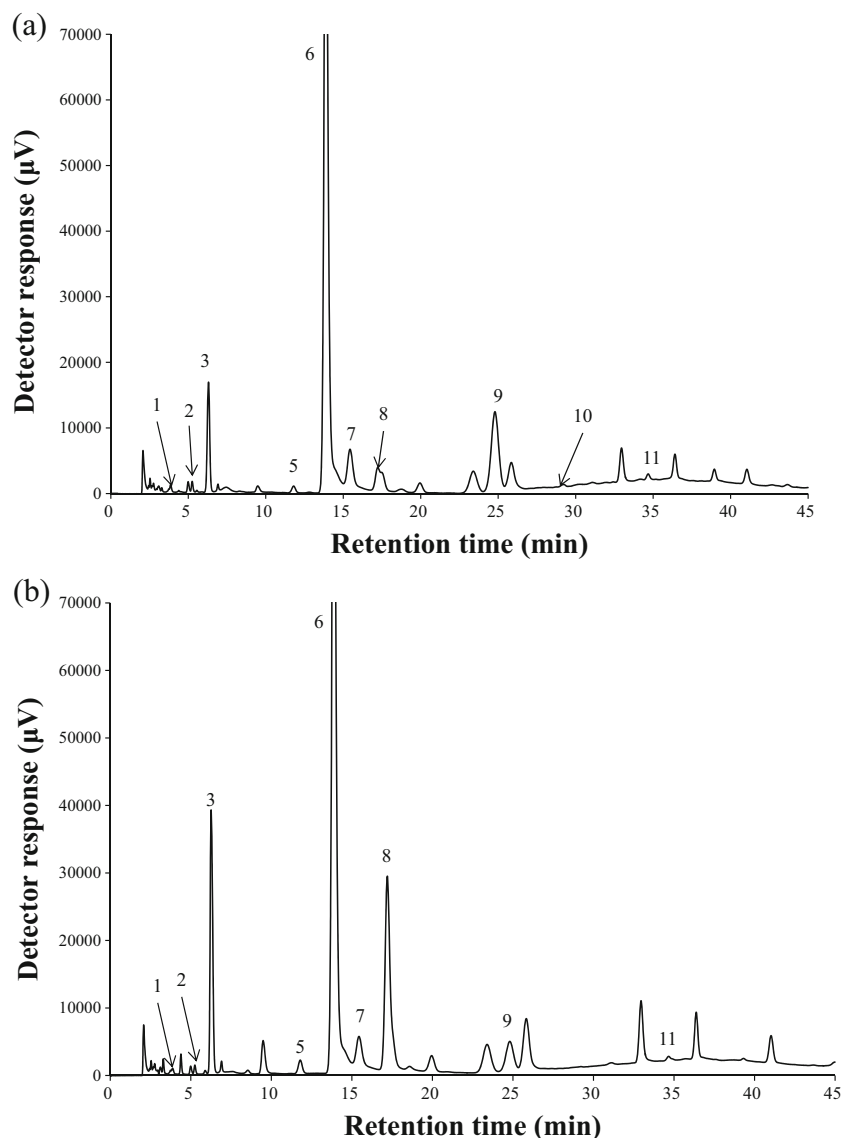
Storage hours	0	24	48	72	96
CMP	0.20±0.07	0.11±0.03	0.14±0.11	nd ^c	0.14±0.07
Guanine	0.68±1.23	0.12±0.05	0.14±0.22	0.40±0.36	0.27±0.11
Hx	0.06±0.03	0.06±0.02	0.26±0.20	0.62±0.47	0.50±0.16
Xt	nd	nd	nd	nd	nd
GMP	0.26±0.15	0.21±0.09	0.29±0.09	0.67±0.53	0.41±0.18
IMP	88.70±28.80	71.93±19.64	72.82±25.40	105.31±63.48	86.77±34.60
HxR	6.21±1.86	5.04±1.19	12.03±4.59	25.54±12.44	27.89±12.41
AMP	0.85±0.18	0.71±0.19	0.56±0.21	0.97±0.50	0.82±0.29
ADP	7.91±3.16	4.64±2.79	4.77±2.77	8.53±6.44	6.20±2.26
AdR	0.15±0.16	0.02±0.01	0.32±0.23	nd	nd
ATP	1.49±1.43	1.08±0.57	6.32±5.72	0.20±0.50	0.18±0.44
K value (%)	8.69±0.88	8.97±0.86	18.54±4.65	26.25±1.89	30.78±2.22

^a Purine and pyrimidine compounds contents represented as mg 100 g⁻¹ muscle

^b The K value was evaluated using the equation below: K value (%)=(HxR+Hx) / (ATP+ADP+AMP+IMP+HxR+Hx) × 100

^c nd not detected

Fig. 2 HPLC chromatograms of purine and pyrimidine compounds in high-freshness- and low-freshness-level amberjacks. **a** High-freshness-level amberjack (before storage). **b** Low-freshness-level amberjack (after storage at 5 °C for 96 h). A number of each chromatogram represents each compound: 1, CMP; 2, guanine; 3, Hx; 4, Xt; 5, GMP; 6, IMP; 7, HxR; 8, AMP; 9, ADP; 10, AdR; 11, ATP



were mixed with 9.8 mL of 1 mol L⁻¹ potassium hydroxide and left in a refrigerator for 30 min. Then, the samples were separated by centrifugation at 1500 g for 10 min. The supernatants were poured into another set of four plastic test tubes and diluted to a total volume of 20 mL with water. Each of these solutions was filtered through a 0.45-µm filter (Millex-LH 0.45 µm, Merck Millipore Ltd., Darmstadt, Germany) and 4 mL of the filtered solution was added to 1 mL of 100 mM phosphate buffer (pH 7.5). These solutions were analyzed with HPLC. The recovery and the relative standard deviation (RSD) were estimated using the results of the analysis.

Evaluation of Amberjack Freshness

The purine and pyrimidine compounds content and *K* value of amberjack samples (*n*=6) of various freshness levels were analyzed. Each amberjack was bled by cutting its gills, and

its internal organs were removed. The dressed fish was cut into loins and then into fan-shaped blocks (weight, ca. 50 g; thickness, 20 mm) from the dorsal muscle. The experimental protocol was approved by the Institutional Animal Care and Use Committee, University of Miyazaki. The fan-shaped blocks were stored in a refrigerator at 5 °C for 96 h and were taken from the refrigerator at each time point (24, 48, 72, and 96 h). Purine and pyrimidine compounds were extracted from the muscle using the method of recovery test. The extracts were analyzed using HPLC and the contents of each component and *K* value were evaluated.

The Purine and Pyrimidine Compound Content of Foods

The purine and pyrimidine compound content of foods was analyzed using an HPLC method. Squid (*Todarodes pacificus*), canned crab (the snow crab *Chionoecetes opilio*),

Table 5 Purine and pyrimidine compound contents of foods^a

	Squid		Laver		Crab (canned)	Shiitake
	High-freshness ^b	Low-freshness ^b	High-grade ^c	Low-grade ^c		
CMP	2.39±0.27	3.17±0.24	17.01±1.67	17.76±2.37	nd ^d	0.75±0.04
Guanine	0.21±0.03	0.43±0.06	2.28±0.03	3.82±0.48	nd	1.50±0.07
Hx	5.88±0.09	29.86±0.80	3.74±0.25	4.25±0.98	nd	0.07±0.03
Xt	0.15±0.03	4.35±0.25	nd	nd	nd	1.11±0.04
GMP	0.96±0.04	0.52±0.00	7.56±0.77	0.86±0.16	7.94±0.46	9.07±0.26
IMP	0.90±0.13	nd	11.03±1.77	2.91±0.74	9.89±0.71	nd
HxR	63.99±0.35	6.48±0.27	47.18±4.22	36.00±5.40	nd	4.53±0.11
AMP	11.01±0.05	0.84±0.08	57.78±5.16	61.25±10.90	0.57±0.01	1.07±0.04
ADP	4.83±0.25	1.51±0.05	117.43±11.02	4.07±1.04	nd	1.94±0.31
AdR	0.08±0.01	nd	11.93±1.15	6.39±0.98	nd	0.42±0.03
ATP	nd	nd	122.12±11.21	nd	nd	nd

^a Purine and pyrimidine compounds contents represented as mg 100 g⁻¹ sample

^b Xt ratio(%) of high and low-freshness squids were 1.77±0.37 and 102.13±2.74, respectively, which calculated from purine and pyrimidine compounds

^c Lavers were organoleptically judged by expert panels according to taste, color, and flavor and classified by high or low grades

^d nd not detected

and dried shiitake (*Lentinula edodes*) were purchased at a local supermarket. The squid's mantle muscle was divided into two pieces, and one piece was treated immediately. The other piece was left for 24 h at room temperature and then treated. Two different quality lavers (*Porphyra tenera*) were gifted from the Ariake fishermen's cooperative, Saga Pref., Japan. These samples were cultured and manufactured for dried laver sheets in the Ariake Sea in Japan in 2007–2009 and were organoleptically judged by expert panels of the fishermen's cooperative according to taste, color, and flavor of each sample. The samples were then classified by high or low grades. Purine and pyrimidine compounds were extracted from squid, laver, canned crab, and shiitake using the method of recovery test. The extracts were analyzed using HPLC.

Results and Discussion

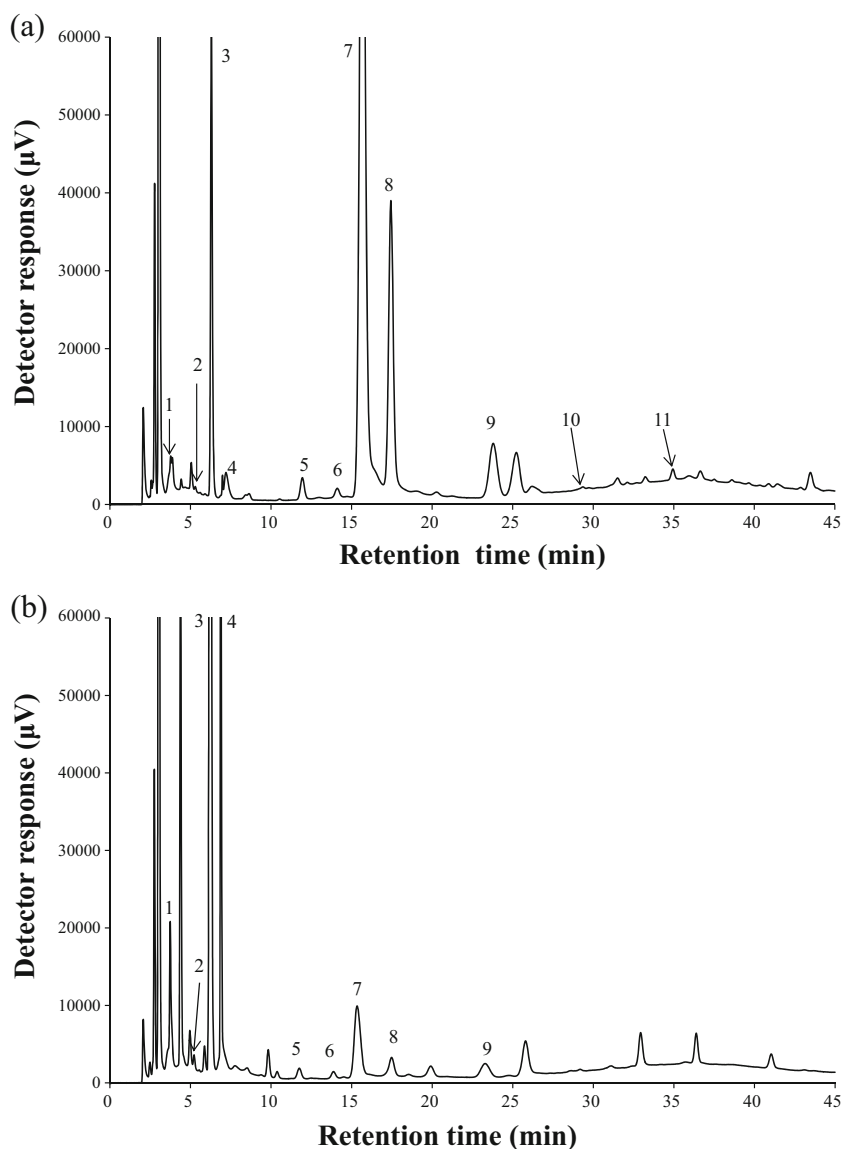
Method Validation

A typical chromatogram of mixed purine and pyrimidine compounds is shown in Fig. 1. Six purine and pyrimidine compounds used to estimate the *K* value were measured on reversed-phase C18 columns using a mobile phase of phosphate buffers (Ryder 1985). However, under these conditions, all peaks without HxR were close to each other so that the purine and pyrimidine compounds related to the freshness of marine invertebrates (Xt and AdR) and those related to umami taste (CMP and GMP) could not be separated simultaneously. To circumvent this issue, Yokoyama et al. (1992) changed the mobile phase to a mixture of citric acid, acetic acid, and

triethylamine and lowered its pH to 4.8. Consequently, the six purine and pyrimidine compounds related to the *K* value and Xt were separated simultaneously. However, CMP and GMP could not be separated using this method. In this study, the six compounds used to estimate the *K* value were separated from other compounds (Xt, CMP, and GMP) by lowering the pH of the mobile phase to pH 3.8, but ADP, AdR, and ATP could not be eluted in 100 min under these conditions. Özogul et al. (2000) could determine the *K* value within a short time using a gradient system with acetonitrile based on Ryder's method (1985). Following this, we attempted to use a gradient system with a mixture of acetonitrile and water as the mobile phase. As a result, 11 purine and pyrimidine compounds were separated with good specificity in a 55-min cycle. This included not only the six compounds used to estimate the *K* value, but guanine, Xt, AdR, CMP, and GMP as well.

To confirm the precision and specificity of the method developed here, tR, peak area, *K'*, Rs, and α were investigated; the results are displayed in Table 1. The RSDs of the tRs for all the purine and pyrimidine compounds were less than 0.20 %. The RSDs of the peak areas for all purine and pyrimidine compounds, except AdR and ATP, were less than 2.00 %, and the RSDs of the peak areas of AdR and ATP were 2.266 and 3.538 %, respectively. The separation of each purine and pyrimidine compound exhibited good specificity. The RSDs of the peak areas obtained using the HPLC method developed here were less than 4 % for each compound, which suggests that the method developed here exhibits high reproducibility. The linearity, range of detection, LOD, and LOQ were investigated; the results are shown in Table 2. The correlation factor for each of the purine and pyrimidine compounds was more

Fig. 3 HPLC chromatograms of purine and pyrimidine compounds in high-freshness- and low-freshness-level squids. **a** High-freshness-level squid. **b** Low-freshness-level squid. A number of each chromatogram represents each compound: 1, CMP; 2, guanine; 3, Hx; 4, Xt; 5, GMP; 6, IMP; 7, HxR; 8, AMP; 9, ADP; 10, AdR; 11, ATP



than 0.999. This suggests the method developed here exhibits good linearity. The linear range varied depending on the purine and pyrimidine compound species. The LODs and LOQs for the various purine and pyrimidine compounds were found to be from 0.8624 to 5.0253 and from 2.8748 to 16.7511, respectively. The mean recovery for each of the purine and pyrimidine compounds was investigated; the results are shown in Table 3. The recoveries of all the purine and pyrimidine compounds, except IMP, from fish were 74.09–120.59 %. The recoveries of IMP ranged from 104.93 to 242.97 %. The average recoveries obtained using the method of Ryder (1985) ranged from 94.2 to 100.1 %, depending on the compound. In our developed method, the average recoveries of all compounds except IMP ranged from 84.28 to 104.52 %, indicating that the method has the sufficient quantitative ability. When a lower concentration of a standard solution was added to the sample, the recovery increased. This

tendency is consistent with the method of Ryder (1985). Concerning IMP, when the concentration of the added standard was original or half of the original, the recovery was 107.37 or 104.93 %, respectively. When the added standard solution was a quarter of the original, the recovery was 242.97 %. The reason for this sharp increase might be that the peak area of the added IMP was too small compared with that of the natural IMP. This may occur because the absorption of 270-nm lights on IMP is lower than that on the other compounds, but dead fish have very large amounts of IMP.

Evaluation of Amberjack Freshness

The purine and pyrimidine compounds contents and K values of amberjack of varying freshness during cooled storage at 5 °C are shown in Table 4. Typical chromatograms from high freshness-level fish and low-freshness-level fish are shown in

Fig. 4 HPLC chromatograms of purine and pyrimidine compounds in high-grade and low-grade lavers. **a** High-grade laver. **b** Low-grade laver. A number of each chromatogram represents each compound: 1, CMP; 2, guanine; 3, Hx; 4, Xt; 5, GMP; 6, IMP; 7, HxR; 8, AMP; 9, ADP; 10, AdR; 11, ATP

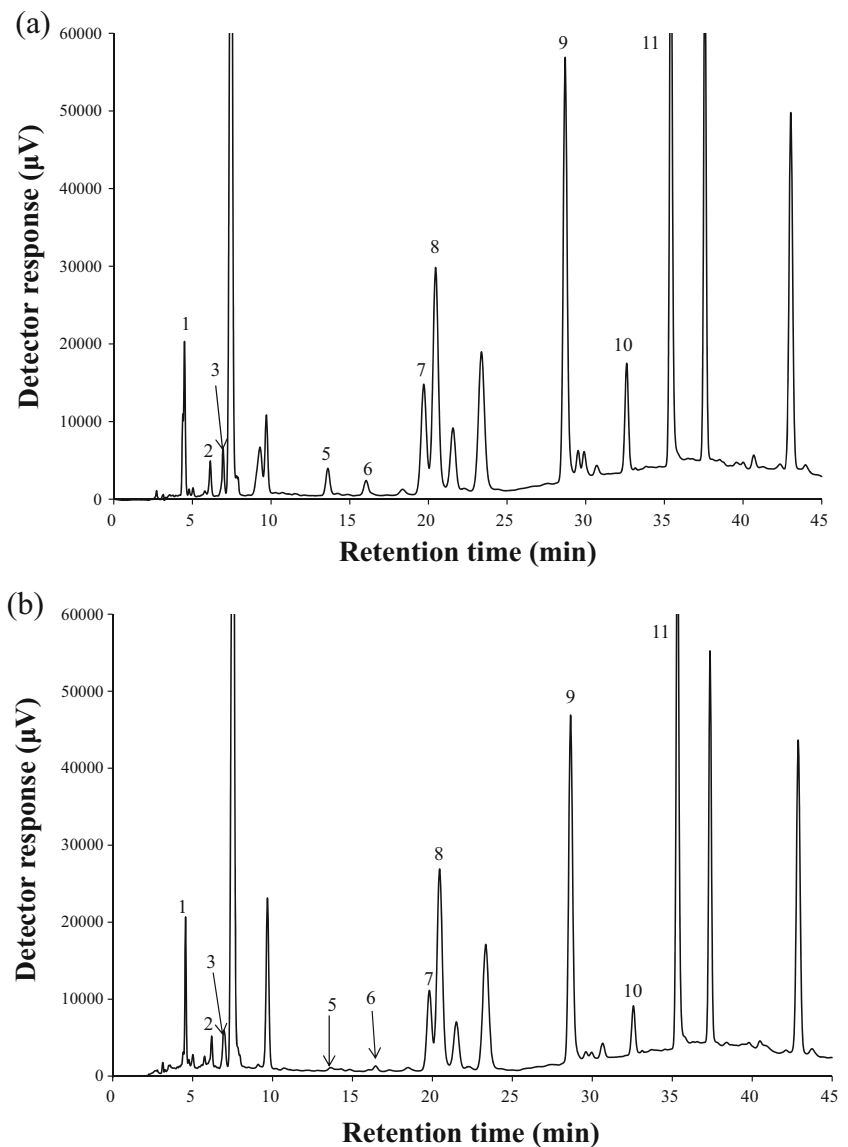


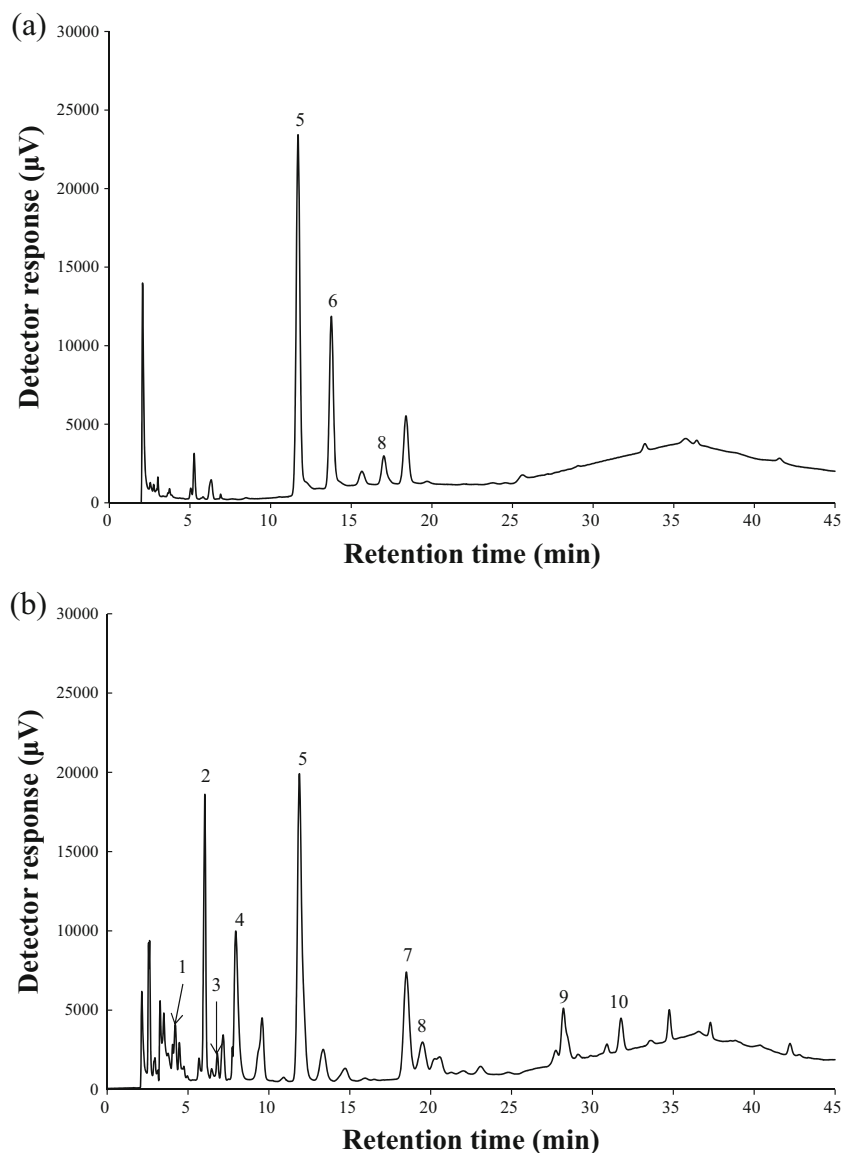
Fig. 2a, b, respectively. The content of CMP was low, and Xt was not detected from 0 to 96 h. The small content of AdR were detected at 0, 24, and 48 h but then vanished at 72 h. The contents of AMP, ADP, and ATP varied with each time point. The content of IMP changed, but there was no significant difference. The contents of Hx did not change between 0 and 24 h. The content of HxR at 0 h was higher than that of 24 h, but there were no significant differences between them. Then, the contents of Hx and HxR increased gradationally from 24 to 96 h. Consistent with this, the *K* value of amberjack did not also change between 0 and 24 h and then increased gradationally to 96 h. The GMP content was analyzed simultaneously with the other purine and pyrimidine compounds associated with the *K* value. The GMP content changed very little from 0 to 48 h (approximately $0.25 \text{ mg } 100 \text{ g}^{-1}$ muscle), increased at 72 h ($0.67 \pm 0.53 \text{ mg } 100 \text{ g}^{-1}$ muscle), and then decreased at 96 h ($0.41 \pm 0.18 \text{ mg } 100 \text{ g}^{-1}$ muscle).

The *K* value is commonly used as an indicator of the initial freshness of fish. Many methods of *K* value determination have been studied. In particular, many HPLC methods have been reported (Lee et al. 1982; Özogul et al. 2000; Ryder 1985). In this study, the *K* values of amberjack stored at 5°C for 96 h were measured using the method developed here. The *K* value before storage was $8.69 \pm 0.88 \%$. It increased to $26.25 \pm 1.89 \%$ after 72 h and reached $30.78 \pm 2.22 \%$ after 96 h. This suggests that the method developed here can analyze the *K* value of fish sufficiently.

Contents of Purine and Pyrimidine Compounds in Foods

Purine and pyrimidine compound contents of foods were investigated using the method developed here (Table 5). Typical chromatograms of high-freshness- and low-freshness-level squids are shown in Fig. 3a, b, respectively. High-freshness-

Fig. 5 HPLC chromatograms of purine and pyrimidine compounds in foods. **a** Canned crab. **b** Shiitake. A number of each chromatogram represents each compound: 1, CMP; 2, guanine; 3, Hx; 4, Xt; 5, GMP; 6, IMP; 7, HxR; 8, AMP; 9, ADP; 10, AdR; 11, ATP



level squid was found to have lower amounts of CMP, Hx, and Xt and higher amounts of HxR, AdR, and ATP than low-freshness-level squid. The Xt ratios of high-freshness- and low-freshness-level squids were 1.77 ± 0.37 and $102.13 \pm 2.74\%$, respectively. Yokoyama et al. (1994) defined the Xt ratio as an indicator of freshness for marine invertebrates and evaluated the freshness of squid, which could not be measured using the *K* value. In this study, the Xt ratio of squid was measured using the method developed here. Using this method, the Xt ratios of high-freshness- and low-freshness-level squids could be easily distinguished. Therefore, the freshness of marine invertebrates such as squid, shrimp, and crab can be evaluated using the method developed here.

Typical chromatograms of high- and low-grade lavers are shown in Fig. 4a, b, respectively. High-grade laver was found to have greater amounts of GMP and IMP than low-grade laver.

Noda et al. (1975) measured the GMP content of laver (*Porphyra spp.*) and found that natural laver contained 5.7–12.5 mg GMP per 100 g of sample. In this study, high-grade laver was found to contain 7.56 ± 0.77 mg GMP per 100 g of sample, which is consistent with the levels measured by Noda et al. (1975).

Typical chromatograms of canned crab are shown in Fig. 5a. Canned crab was found to contain large amounts of GMP and IMP and some amount of AMP. Canned crab did not contain the other purine and pyrimidine compounds. Canned crab was found to have little purine and pyrimidine compounds but large amounts of GMP, IMP, and AMP, which are related to the umami taste. Mouri et al. (1965) measured the purine and pyrimidine compounds content of crab at each stage (raw, during treatment for canning, and after treatment for canning). As a result, the content of most compounds decreased, but the AMP content increased during the canning

process. Results obtained using the method developed here agreed with those of Mouri et al. (1965). The high GMP and IMP contents are thought to be from the umami seasoning. This suggests that the method developed here can analyze a food additive.

Finally, to confirm the applicability of the method for terrestrial foods, shiitake, which contains large amounts of nucleotide 5'-monophosphates, was analyzed using the method developed here. Typical chromatograms of shiitake are shown in Fig. 5b. Shiitake was found to contain some amounts of CMP, GMP, and AMP. GMP, in particular, was found in large quantities. Sasaki et al. (2000) measured the GMP content of dried shiitake and found it to be 10.72 g 100 g⁻¹. In this study, the GMP content of dried shiitake was found to be 9.07 ± 0.26 g 100 g⁻¹, which agrees with the result of Sasaki et al. (2000). Kasuga et al. (1999) also measured the nucleotide content of dried shiitake and reported that dried shiitake contained some amount of CMP. In our study, a small amount of CMP was detected in dried shiitake, which agrees with the result of Kasuga et al. (1999).

These results demonstrate that the HPLC method developed here is useful for the analysis of purine and pyrimidine compounds in foods associated with the umami taste.

Conclusion

The results of this study show that the method developed here separates more kinds of purine and pyrimidine compounds than the usual methods, but require less equipment and time. Moreover, the method developed here shows sufficient reproducibility and quantitative ability.

Using this method, compounds associated with the freshness of marine vertebrates and invertebrates as well as compounds associated with the umami taste can be analyzed simultaneously. Therefore, the method developed here would be applicable to the evaluation of correlations between freshness and umami taste for a wide range of marine products as well as the evaluation of umami taste for terrestrial foods.

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Compliance with Ethical Standards

Conflict of Interest Mami Ishimaru declares that she has no conflict of interest. Masato Haraoka declares that he has no conflict of interest. Hideo Hatate declares that he has no conflict of interest. Ryusuke Tanaka declares that he has no conflict of interest.

Ethical Approval This article contains some studies with fish subjects. All procedures performed in studies involving animals (fish) were in accordance with the ethical standards of the institution or practice at which the studies were conducted.

Informed Consent Not applicable.

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