18. ORGANIC PROPERTIES OF SEDIMENTS AND AMINO ACIDS IN INTERSTITIAL WATERS FROM THE FLANK OF THE COSTA RICA RIFT, GALAPAGOS SPREADING CENTER (ODP SITES 677 AND 678)¹

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INTRODUCTION

The basement at Ocean Drilling Program (ODP) Sites 677 and 678 originated from the Galapagos spreading center of the Costa Rica Rift and has moved about 200 km over the last 6 m.y. (Fig. 1) (Shipboard Scientific Party, 1987, 1988; Scientific Drilling Party, 1987). Sediments about 300 m thick cover basement so young that basal sediments at Sites 677 and 678 have been reheated up to 60°-70°C at Site 677 and altered to limestone and/or chert (Shipboard Scientific Party, 1988). Sediments from both sites indicate (1) a high sedimentation rate (about 48 m/m.y.) and (2) biogenic silica and carbonate as the main constituents of sediments (Table 1) (Shipboard Scientific Party, 1988). Heatflow observations and measurements of interstitialwater chemistry around the sites show that Site 677 is in a lower heatflow zone (166 mW/m²; 1°12.14'N, 83°44.22'W) whereas Site 678 is located in a zone of higher heat flow (250 mW/m²; 1°13.01'N, 83°43.39'W) (Langseth et al., 1988; Shipboard Scientific Party, 1988). In the flank hydrothermal systems, circulating solution is moving upward through the sedimentary column in zones of higher heat flow while it is moving downward in zones of lower heat flow (Anderson and Skilbeck, 1981).

The chemistry of the interstitial waters is modified by several processes such as (1) diagenetic reactions and (2) advective and (3) diffusive transports of dissolved constituents. Analyses of Ca^{2+} and Mg^{2+} in interstitial waters from Sites 677 and 678 show that their profiles are mainly controlled by advective transport (Shipboard Scientific Party, 1988). In contrast, the interstitial-water profiles for NH⁴⁺, Si, and PO₄⁻³ are highly affected by reactions in the sediments. Site 677 offers a good opportunity to investigate amino acids in the interstitial waters because sediments of similar compositions have been deposited at constant rates of sedimentation. There are few previous works on amino acid distributions in interstitial waters (Henrichs and Farrington, 1979; Michaelis et al., 1982; Henrichs et al., 1988).

In this chapter, we report (1) Rock-Eval analysis and (2) the composition of total hydrolyzable and dissolved free amino acids (THAA and DFAA, respectively) in the interstitial waters. Our objectives are to discuss (1) the possible origin of organic materials, (2) the characteristics of THAA and DFAA, and (3) their relationships in interstitial waters.

METHODS

Sampling and Storage

Interstitial-water samples from Sites 677 and 678 were collected aboard JOIDES Resolution by hydraulic squeezing at the same tempera-

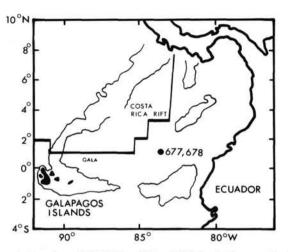


Figure 1. Location of ODP Sites 677 and 678 in the Panama Basin of the Costa Rica Rift. Water depths at Sites 677 and 678 are 3461 and 3435 m, respectively.

ture at which the sediment was recovered. All interstitial-water samples were filtered through 0.22- μ m Millipore cellulose acetate filters in the shipboard laboratory. For amino acid analysis of interstitial water, a 5-mL aliquot of the water sample was placed into a precombusted glass ampule. The ampule was sealed after flushing with helium, stored in a refrigerator aboard ship, and returned to the shorebased laboratory for analysis.

Analytical Procedures

Amino Acids

Interstitial-water DFAA was analyzed after the addition of 7 μ L of ultrapure (amino acid-free) 17% HCl to 0.7 mL of an interstitial-water sample. Each amino acid concentration was determined by direct injection into an automated liquid chromatograph (Hitachi Model 835). The reagent blank was suitably corrected.

For THAA, a 1-mL interstitial-water sample, together with 1 mL of ultrapure 17% HCl, was placed in a precombusted glass ampule, and hydrolysis was conducted at 110°C for 22 hr under an argon atmosphere. Then, the solution was gently evaporated using a rotary evaporator at a temperature lower than 42°C, and the residue was redissolved in 1 mL of 0.01 M HCl. The resulting aliquot was injected into the amino acid analyzer. The standard deviation for amino acid analysis based on replicated measurements of a standard solution (about μ mol/L for amino acids) was less than 10%, and the detection limit of our method was about 0.05 μ mol/L. Low values in free and hydrolyzable amino acids near the detection limit may not be dependable. Nevertheless, we retain these values because they confirm that the amino acid concentration is low (Ishizuka et al., 1988).

Rock-Eval Analysis

After squeezing, remnants of the interstitial-water sediment samples were oven-dried at 60°C and ground into powder. These samples were analyzed by pyrolysis in the shipboard Rock-Eval unit.

¹ Becker, K., Sakai, H., et al., 1989. Proc. ODP, Sci. Results, 111: College Station, TX (Ocean Drilling Program).

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Unit	Depth (mbsf)	Lithology	Age	Porosity ^a (%)	Density ^a (g/cm ³)	Sedimentation rate (m/m.y.)
Hole 677A						
I	0-153.8	Alternating clayey biogenic calcareous siliceous oozes and clayey biogenic siliceous calcareous oozes	early Pliocene-late Pleistocene	90	1.2	48
11	153.8-308.5	Siliceous nannofossil and nannofossil chalk	late Miocene	70	1.8	48
ш	303.3-308.5	Cherty limestone and nannofossil chalk	late Miocene	50	1.8	48
IV	308.5-309.5	Iron oxide- and smectite- rich sediments inter- mixed with glassy basement basalts	late Miocene			
Hole 678B						
I	0-27.7	Clayey calcareous siliceous ooze	late Pliocene-late Pleistocene	83	1.3	
П	95.5-111.8	Clayey diatom nannofossil chalk	late Miocene	77	1.4	
III	111.8-169.8	Limestone	late Miocene	75	1.5	
IV	169.8-170.1	Basalt fragments and conglomerates	late Miocene			

Table 1. Characteristics of lithologic units, Holes 677A and 678B.

^a Mean values.

RESULTS

Pyrolysis Assay (Rock-Eval)

The total organic carbon (TOC) content of the sediment residues from the interstitial-water squeezing process is highly variable, but appears to have a generally decreasing trend with depth. TOC values range from 0.10 to 0.81 wt%, except for a single sample with 2.92 wt% (Table 2 and Fig. 2). The maximum TOC value occurs at about 80 m below seafloor (mbsf). The values are similar to those from Hole 678B.

The total hydrocarbon yields, S1 (free, distillable hydrocarbons) + S2 (generatable hydrocarbons), from Holes 677A and 678B are low (0–9 mg hydrocarbons/g rock) and show the same profile as the TOC. The oxygen index (OI = S3/TOC) is highly scattered, with a mean value of 644. Most of the organic matter is highly oxidized, as indicated by the high OI values. The hydrogen index (HI = S2/TOC) ranges from 0 to 292, and its mean value is 126 (Table 2 and Fig. 2).

Pyrolysis offers an indication of organic maturity through T_{max} , the temperature at which there is maximum hydrocarbon generation. T_{max} values from Sites 677 and 678 are up to 419°C (Table 2 and Fig. 2), indicating that the organic matter is thermally immature and has not entered into the stage of peak hydrocarbon generation.

THAA in Interstitial Waters

The concentrations of individual THAA are given in Table 3. The total concentration of THAA ranges from 1.80 to 31.50 μ mol/L, averaging 8.34 μ mol/L. Relatively high values are found at near the surface and at depths of 25.8, 196.6, and 274.2 mbsf (Fig. 3). The THAA profile is different from those of dissolved Ca²⁺ and Mg²⁺ measured in interstitial waters, which are mainly controlled by downward advective flow (Shipboard Scientific Party, 1987).

In spite of high fluctuations of THAA, the relative abundance of individual amino acids is similar throughout the hole. The mean values of acidic, basic, neutral, aromatic, and sulfurcontaining amino acids are 2.28, 1.25, 4.62, 0.24, and 0.01 μ mol/L, respectively.

DFAA in Interstitial Waters

The concentrations of individual DFAA are given in Table 4. The total concentration of amino acids ranges from 0.20 to 11.8 μ mol/L, averaging 3.19 μ mol/L. Relatively high values are found near the surface and at depths of 25.8, 179.2, and 285.0 mbsf (Fig. 3). The total DFAA profile vs. sub-bottom depth correlates with that of total THAA.

The data in Table 4 reveal the following general order of amino acid abundance: neutrals > basics > acids > aromatics > sulfur-containing. The mean values of acidic, basic, neutral, aromatic, and sulfur-containing amino acids are 0.18, 0.67, 2.24, 0.08, and 0.00 μ mol/L, respectively.

DISCUSSION

Organic Material from Sites 677 and 678

The TOC values from Holes 677A and 678B are, in general, high compared with those recorded in open-marine deep-sea sediments, which are usually less than 0.3 wt% (McIver, 1975). The high values are probably caused (1) by high oceanic productivity, such as in equatorial upwelling areas, and (2) by rapid burial of organic carbon (C_{org}) as a result of high sedimentation rates. These reasons are supported by two observations: (1) that biogenic siliceous and calcareous fragments, which have supplied much of the organic carbon, constitute the sediments at Sites 677 and 678 and (2) that much framboidal pyrite formed as a result of the reduction of sulfate ions in the interstitial water by organic materials occurring throughout the hole.

The HI and OI values determined by Rock-Eval analyses are plotted in Figure 4 on a van Krevelen diagram (Tissot and Welte, 1984). This diagram implies that most of the organic matter in the sediments at Sites 677 and 678 is of a type II/III mixture, that is, a mixture of marine and terrigenous organic matter. A large contribution of terrestrial organic matter seems unlikely, however, because the terrigenous inorganic constituents (e.g., feldspar and quartz) make up only about 1%-4% of the total sediments, whereas marine biogenic fragments comprise about 70%-90%, based upon our microscopic examination. In addi-

Core, section, interval (cm)	Depth (mbsf)	TOC (wt%)	carb	S2 nydro- ons/ ock)	S3 (mg CO ₂ / g rock)	S1 + S2	HI (mg S2 hydrocarbons/g organic carbon)	OI (mg S3 CO ₂ /g organic carbon)	T _{max} (°C)
111-677A-									
1H-3, 145-150	4.5	0.41	0.18	0.85	2.93	1.03	207	714	389
2H-1, 80-89	7.0	0.70	0.21	1.23	4.13	1.44	175	590	396
3H-4, 120-125	21.4	0.90	0.28	2.00	4.04	2.28	222	448	413
4H-1, 55-74	25.8	0.58	0.19	0.94	3.72	1.13	162	641	411
5H-1, 51-69	35.3	0.49	0.24	0.75	4.45	0.99	153	908	385
6H-3, 120-125	48.4	0.74	0.19	0.93	3.81	1.12	125	514	406
9H-3, 120-125	76.9	2.91	0.55	8.51	4.80	9.06	292	164	419
12H-4, 120-125	106.9	0.32	0.13	0.44	3.46	0.57	137	1081	391
13H-4, 145-150	116.7	0.27	0.09	0.38	2.20	0.47	140	814	393
14H-4, 145-150	126.2	0.70	0.16	1.20	3.11	1.36	171	444	413
15H-4, 120-125	135.4	0.28	0.09	0.31	2.82	0.40	110	1007	389
16X-4, 145-150	145.2	0.38	0.11	0.33	2.27	0.44	86	597	392
17X-3, 145-150	148.7	0.30	0.45	0.27	2.01	0.72	90	670	392
18X-3, 120-125	158.0	0.39	0.08	0.35	2.36	0.43	89	605	393
19X-4, 145-150	169.5	0.17	0.08	0.17	2.06	0.25	100	1211	384
20X-4, 145-150	179.2	0.32	0.09	0.43	1.32	0.52	134	412	406
22X-3, 120-125	196.6	0.41	0.10	0.21	1.33	0.31	51	324	395
23X-4, 145-150	208.1	0.13	0.10	0.11	1.08	0.21	84	830	388
25X-4, 120-125	227.1	0.12	0.03	0.16	1.14	0.19	133	950	400
27X-4, 145-150	246.8	0.29	0.04	0.40	1.10	0.44	137	379	406
29X-3, 145-150	264.5	0.14	0.03	0.16	1.10	0.19	114	785	396
31X-4, 120-125	285.0	0.10	0.04	0.17	1.08	0.21	170	1080	400
33X-2, 145-150	301.6	0.11	0.02	0.08	0.89	0.10	72	809	392
33X-3, 133-150	303.0	0.18	0.02	0.14	0.92	0.16	77	511	416
111-678B-									
1H-1, 45-50	0.5	0.86	0.29	1.61	4.70	1.90	187	546	417
1H-2, 145-150	3.0	0.67	0.37	1.19	4.57	1.56	177	682	398
2H-1, 145-150	19.7	0.61	0.14	0.62	3.41	0.76	101	559	397
2H-4, 145-150	24.2	0.51	0.14	0.58	3.62	0.72	113	709	388
3H-1, 145-150	97.0	0.45	0.04	0.30	1.80	0.34	66	400	399
3H-6, 145-150	104.5	0.51	0.07	0.53	1.85	0.60	103	362	402
4W-3, 95-107	138.9	0.33	0.03	0.13	1.82	0.16	39	551	391
4W-CC, 3-15	168.9	0.26	0.00	0.00	0.77	0.00	0	296	337

Table 2. Rock-Eval analysis.

tion, marine sediments off coasts generally have a predominance of marine organic matter. The characteristics of OI and HI of original organic material may be overprinted by the mechanisms of (1) the effect by microbiological degradation (Whelan and Tarafa, 1986) and (2) degradation by montmorillonite catalvsis (Whelan and Hunt, 1983).

Amino Acids in Interstitial Waters

Characteristics of THAA

The characteristics of THAA from Hole 677A are as follows:

1. THAA concentrations are similar to the values found for biogenic carbonate-rich sediments from the Atlantic Ocean (Ishizuka et al., 1988).

2. Glutamic and aspartic acids, glycine, and serine are the major constituents.

3. THAA composition of interstitial waters is fairly uniform throughout the hole (Table 3).

These characteristics are explained by (1) the THAA compositions and concentrations of the original seawater trapped with the sediment particles when they were deposited, (2) the THAA compositions and concentrations of proteinaceous materials in sediment particles, (3) the production rate of THAA from sediment by chemical and biological processes, (4) the decomposition rate of THAA in interstitial waters, and (5) the water content or porosity. The porosity of the sediments from Hole 677A averages about 90% in lithologic Unit I and decreases to 50% in Unit III (Table 1). There is no correlation between porosity and composition of THAA in the interstitial waters of Hole 677A.

THAA of deep-sea water from the Pacific Ocean shows (1) that alanine accounts for up to 60%-70% of the total amino acids in seawater and (2) that the total concentration of THAA is less than 100 nmol/kg (Bada et al., 1982). Because the concentrations of total THAA and DFAA of interstitial water from Hole 677A are about 8.34 and 3.19 μ mol/L, respectively, the concentration of THAA in the original seawater must have been one or two orders lower in magnitude. Reaction between interstitial water and sediments during early diagenesis plays an important role in the large discrepancy between the amino acid composition of THAA of seawater and interstitial waters.

Sources of THAA

Planktonic and terrestrial organic matters are considered to be the most important amino acid contributors in view of the Rock-Eval analyses, smear slide observations, and the inorganic chemistry of sediments from Sites 677 and 678. Fragments of the calcareous and siliceous microfossils make up about 70%– 90% of the total sediments. Aluminous smectites, quartz, feldspars, and rutile were transported from land and deposited at Sites 677 and 678 (Beiersdorf and Rosch, 1983; Beiersdorf and Natland, 1983). Relative abundances of acidic, basic, and neutral amino acids of DFAA and THAA in the interstitial waters are plotted as a function of sub-bottom depth in Figure 5. The relative abundances are uniform, especially in the upper 220 m

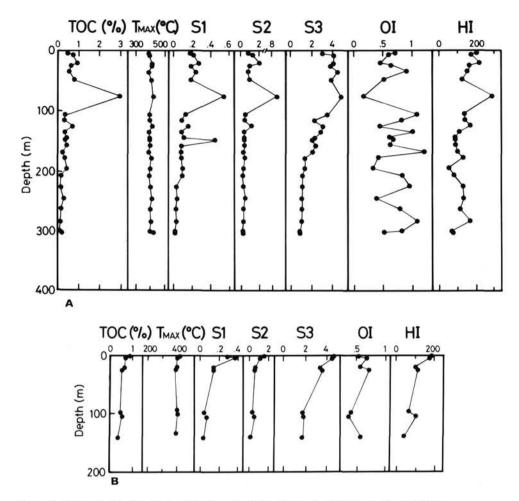


Figure 2. TOC, HI, OI, S1, S2, and S3 vs. sub-bottom depth. A. Hole 677A. B. Hole 678B.

of sediments, in spite of large fluctuations in the concentrations. The relation between neutral, acidic, and basic amino acids is shown in Figure 6. The correlation factors of neutral to acidic and basic of THAA are 0.90 and 0.72, respectively. These results indicate that simple decomposition of proteinaceous materials in the sediments plays a more important role than discrimination by bacteria in determining the compositions of THAA.

Examination of the qualitative data in Table 3 reveals the order of abundance as neutral > acidic > basic > aromatics > sulfur-containing. In comparison, examination of planktonic amino acid composition in Table 5 (Morris, 1975; Raymont et al., 1975) gives a predominance of neutral over acidic and acidic over basic.

As far as the neutral fraction is concerned, the relative abundance of straight-branch fractions dominates. The ratio between glycine and alanine in Hole 677A is similar to that found in plankton and marine suspensions (Raymont et al., 1975; Ittekkot et al., 1984). The second dominant fraction is the hydroxy amino acids. Within the hydroxy fraction there is a general predominance of serine, as is seen in the plankton, in contrast to the predominance of threonine over serine in sediments enriched in algal cellular debris (Gonzalez et al., 1983). Threonine, serine, and glycine are the main components of a proteinsilica complex in diatom cell walls that should be particularly resistant to environmental degradation (Siezen and Magne, 1978). The diatomaceous silica fragments account for 19% of the total sediments in lithologic Unit I and 14% in Unit II. These lines of evidence suggest that proteins from diatoms contribute as one of the important sources of THAA of interstitial waters. Branched-chain amino acids, including valine, leucine, and isoleucine, in Hole 677A interstitial-water THAA are as abundant as those in plankton.

Within the acidic fraction the relative abundance of glutamic acid is twice that of aspartic acid in plankton. In addition, significant amounts of glutamic acid are contained by marine bacteria (Henrichs and Farrington, 1979) and by oceanic suspended particulate matter (Siezen and Magne, 1978). This contrasts with the predominance of aspartic acids over glutamic acid found in terrestrial humic hydrolates, river suspensions, and sediments rich in terrestrial organic material (Peake et al., 1972; Khan and Sowden, 1972; Pelet and Debyser, 1977). Glutamic acid has three times the concentration of aspartic acid, accounting for 18% of the total THAA, which is consistent with the composition of plankton.

Within the basic fraction of Hole 677A ornithine is the major amino acid and arginine is the minor one in spite of the predominance of arginine over ornithine in marine plankton. The ratio from Hole 677A probably resulted from the decomposition from arginine and transformation to ornithine during early diagenesis.

Aromatic and sulfur-containing amino acids are depleted in THAA in the interstitial waters from Hole 677A. Nissenbaum et al. (1972) reported the predominance of phenylalanine with the oxidizing conditions of deposition. The low content of phenylalanine of THAA in Hole 677A is probably due to reducing conditions resulting from the large supply of organic carbon, in

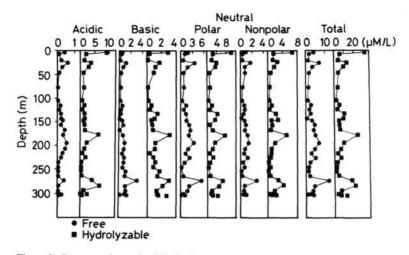


Figure 3. Concentrations of acidic, basic, neutral, and total amino acids of THAA and DFAA in interstitial waters vs. sub-bottom depth in Hole 677A.

consideration of the abundance of framboidal sulfides (about 1 wt%) occurring throughout the hole.

Comparison of DFAA to Other Fields

The DFAA concentrations are of the same order of magnitude as those from the coastal region of Buzzards Bay (Massachusetts) and the upwelling regions off Peru, although the values from the surface to 20 cm in the sediment of coastal and upwelling regions are one or two orders higher (Henrichs et al., 1984; Henrichs and Farrington, 1987). The DFAA of the North Atlantic Ocean abyssal plain is one order of magnitude lower (Ishizuka et al., 1988). These concentrations are probably highly affected by TOC. Relatively organic-rich sediments (TOC > 0.5 wt%) show DFAA values on the order of 10 μ mol/L whereas the DFAA concentrations are very low (<1 μ mol/L) in organicpoor sediments (TOC < 0.2 wt%).

Relative abundances of acidic, basic, and neutral amino acids in the interstitial waters are uniform, especially in the upper 220 m of sediments, as evidenced by THAA (Fig. 5). The correlation factors of neutral to acidic and neutral to basic are 0.67 and 0.74, respectively (Fig. 6). The DFAA compositions of Hole 677A are different than those from estuary and hemipelagic sediments. Glycine, alanine series, and orthine are the major constituents of the DFAA in Hole 677A. Acidic amino acids—including aspartic (3 mol%) and glutamic (5 mol%) acids—are minor constituents. In hemipelagic and coastal sediments, glutamic acid is commonly the major constituent (Henrichs et al., 1984; Henrichs and Farrington, 1987).

Relation between THAA and DFAA

Interesting characteristics of the relation between THAA and DFAA in Hole 677A are (1) the good correlation between total THAA and total DFAA, (2) a much lower concentration of acidic amino acid fractions in DFAA than in THAA, (3) higher concentrations of hydroxyl and straight-branched amino acid fractions in DFAA than in THAA, and (4) very low concentrations of aromatic and sulfur-containing fractions in DFAA and THAA.

The relative abundance of acidic amino acids in DFAA is only one-third that of THAA in Hole 677A (Fig. 5), although the acidic amino acids are more stable than the basic amino acids in the free state (Abelson, 1959). Based on the results of adsorption experiments of amino acids in distilled water, basic (positively charged) amino acids were strongly adsorbed (40%-80% removal) and neutral (uncharged) amino acids were taken up appreciably (10%-15%) by montmorillonite. Acidic (negatively charged) acids were adsorbed very little (Hedges and Hare, 1987). Stevenson and Cheng (1969) found that basic amino acids are more apt to be incorporated into the lattice of clay minerals than acidic amino acids are in the Argentine Basin sediments. Montmorillonites are the major clay minerals in Hole 677A, accounting for 2%-40% of the total sediment composition, but they are not capable of controlling the composition of amino acids by adsorption. Furthermore, quartz does not have the ability to selectively adsorb aspartic acid-rich organic matter (Carter, 1978).

Acidic amino acids inhibit the reaction with calcite, which can be best explained by a specific epitaxial effect in which aspartic or glutamic acid forms a protective overgrowth on the carbonate surface (Jackson and Bischoff, 1971). Carter (1978) and Carter and Mittere (1978) indicated that the carbonate surface appears to selectively adsorb aspartic acid-enriched organic matter whereas noncarbonates do not have this property. Biogenic carbonate fractions in the sediments from Hole 677A make up 40% of the total sediments, and the low concentration of acidic amino acids of DFAA is probably due to reaction or adsorption with carbonates.

SUMMARY AND CONCLUSIONS

Biogenic siliceous and calcareous sediments were drilled at Sites 677 and 678 on the flank of Costa Rica Rift during ODP Leg 111.

1. The TOC values of the sediments range from 0.10 to 2.91 wt%. Mean values of OI and HI are 644 and 126, respectively. T_{max} values from Sites 677 and 678 are up to 419°C, indicating that the organic matter is thermally immature.

2. The total concentration of THAA ranges from 1.80 to 31.50 μ mol/L, averaging 8.34 μ mol/L. The mean values of acidic, basic, neutral, aromatic, and sulfur-containing amino acids are 2.28, 1.25, 4.62, 0.24, and 0.01 μ mol/L, respectively.

3. The total concentration of DFAA ranges from 0.20 to 11.80 μ mol/L, averaging 3.19 μ mol/L. The profile of total DFAA is relatively consistent with that of total THAA from Hole 677A. The mean values of acidic, basic, neutral, aromatic, and sulfur-containing amino acids are 0.18, 0.67, 2.24, 0.08, and 0.0 μ mol/L, respectively.

4. The most plausible source of THAA is marine plankton, based upon the comparison of amino acid compositions between THAA, plankton, and river suspensions.

Table 3. Concentrations of THAA in interstitial-water samples from Hole 677A.

Depth (mbsf)	1H-3, 145-150 4.5	2H-1, 80-89 7.0	3H-4, 120-125 21.4	4H-1, 55-74 25.8	5H-1, 51-69 35.3	6H-3, 120-125 48.4	9H-3, 120-125 76.9	12H-4, 120-125 106.9	13H-4, 145-150 116.7	14H-4, 145-150 126.2	15H-4, 120-125 135.4	16X-4, 145-150 145.2	17X-3, 145-150 148.7	18X-3, 120-125 158.0
Acidic (µmol/L)														
Aspartic acid	3.29	0.36	0.26	0.78	0.73	0.18	0.21	0.18	0.45	0.19	0.56	0.53	0.26	0.31
Glutamic acid	5.86	1.48	1.03	2.75	2.04	0.85	0.85	0.34	0.91	0.52	0.84	0.78	0.74	1.09
Basic														
Ornithine	0.23	0.15	0.07	0.99	0.67	0.72	0.14	0.13	0.41	0.20	0.83	0.50	0.42	0.52
Lysine	0.83	0.14	0.08	0.25	0.16	0.05	0.05	0.04	0.11	0.04	0.16	0.21	0.14	0.14
Histidine	0.52	0.06	-	0.36	0.31	0.13	0.04	0.01	0.09	0.14	0.36	0.08	0.07	0.18
Arginine	1.33	0.03		0.09			0.02	-	0.02	-	0.14	0.22	0.01	0.02
Neutral														
Hydroxy														
Threonine	2.34	0.38	0.25	0.52	0.48	0.10	0.13	0.07	0.23	0.08	0.47	0.42	0.16	0.21
Serine	2.75	0.34	0.13	1.45	1.21	0.19	0.28	0.19	0.78	0.33	1.74	1.14	0.59	0.88
Straight														
Glycine	4.87	1.09	0.71	2.19	1.58	0.56	0.55	0.40	0.82	0.52	2.09	1.60	0.89	1.25
Alanine	2.69	0.83	0.46	1.12	0.77	0.34	0.33	0.22	0.48	0.36	1.01	1.15	0.46	0.69
Branch														
Valine	1.61	0.02	0.02	0.60	0.44	0.18	0.34	0.14	0.35	0.35	0.46	0.58	0.20	0.48
Iso-leucine	1.29	0.12	0.10	0.75	0.16	0.08	0.10	0.09	0.13	0.10	0.65	0.76	0.13	0.07
Leucine	2.27	0.55	0.56	0.66	0.50	0.39	0.09	0.05	0.51	0.31	0.46	0.68	0.43	0.55
Others														
Aspargine Aromatic	—	-	-	-	-	-	-	-	0.09		-	-	-	-
Tyrosine	0.76		-	0.16	0.14		-			-	-	0.14	0.10	-
Phenylalanine	0.47	_	_		-				-	-	-	_	_	-
Sulfur	N													
Cystein		-	—	277		777			-		-	-	-	-
Methionine	0.14	-			_			100			-	0.06	0.01	0.01
Taurine	0.02	0.01	0.01	0.04	0.02	0.01	-		0.01		0.06	0.06	0.01 0.15	0.01 0.06
β-alanine	0.10	0.29	0.34	0.29	0.22	0.58	0.23	0.09	0.08	0.14	0.12	0.17 0.22	0.15	0.06
β -aminobutyric acid	0.14	0.12	0.14	0.17	0.19		0.16	0.08	0.09	0.12				
Total	31.51	5.97	4.16	13.17	9.62	4.36	3.52	2.03	5.56	3.40	10.07	9.24	4.89	6.60

5. The important compositional characteristic of much lower concentrations of acidic amino acid fractions in DFAA than in THAA in the interstitial waters from Hole 677A is probably due to the reaction between acidic amino acids and carbonate.

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Table 3 (continued).

19X-4,	20X-4,	22X-3,	23X-4,	24X-4,	25X-4,	26X-4,	27X-4,	28X-3,	29X-3,	30X-3,	31X-4,	32X-4,	33X-2,	33X-4,	33X-3
145-150 169.5	145-150 179.2	120-125 196.6	145-150 208.1	145-150 217.7	120-125 227.1	145-150 237.1	145-150 246.8	120-125 254.6	145-150 264.5	145-150 274.2	120-125 285.0	145-150 294.9	145-150 301.6	145-150 304.6	133-150 303.0
0.32	2.03	0.41	0.42	0.13	0.33	-	_	_	0.03	0.93	0.43	0.11	_	0.46	0.08
1.18	4.32	1.75	1.09	0.45	1.36	0.35	0.09	-	0.51	3.20	16.25	1.03	0.17	1.94	0.68
0.52	0.54	0.73	0.63	0.35	0.63	0.68	0.52	1.28	0.96	2.37	1.11	0.95	1.06	1.88	1.25
0.19	1.01	0.18	0.18	0.10	0.18	0.17	0.16	0.32	0.18	0.38	0.53	0.21	0.16	0.35	0.14
0.04	0.43 1.12	0.17 0.05	0.14 0.05	0.07 0.01	0.18 0.02	0.19 0.03	0.07 0.02	0.03 0.03	0.23 0.02	0.41 0.14	0.32 0.17	0.17 0.02	0.21 0.03	0.46 0.03	0.08 0.03
0.25	1.61	0.39	0.22	0.14	0.22	_	_	_	_	0.56	0.83	0.75	_	0.23	_
0.94	2.02	1.38	0.83	0.43	0.91	0.44	0.09	-	0.41	2.88	1.72	0.78	0.41	1.61	0.63
1.24	3.31	1.67	1.09	0.66	1.29	0.56	0.29	0.20	0.68	2.80	2.12	1.07	0.57	1.97	0.90
0.74	1.91	1.05	0.60	0.46	0.69	0.18	0.56	-	0.20	1.30	2.81	0.49	0.22	0.90	0.44
0.32	1.56	0.48	0.36	0.16	0.46	—		—	0.08	0.53	0.64	0.16	0.13	—	0.12
0.06 0.19	1.06 1.88	0.59 0.55	0.04 0.48	0.01 0.52	0.07 0.47	0.06 0.11	0.01 0.03	-	0.10 0.13	0.75 0.70	0.69 0.63	0.74 0.17	0.60 0.10	0.60 0.53	0.54 0.13
-	-	_	5 		_	_	-		_	—	1.25	-		-	_
_	0.40	0.13	0.10	-	-		_	_	_	0.64	0.46	0.24	0.51	0.47	0.49
	0.23	_	—	-	-	0.07	0.10	0.10	0.16	0.27	0.21	0.44	0.16	0.36	-
-	-		-		-	-			_	÷	-	-			
0.02	0.03	0.02	-		0.02	-			_	0.02	0.82	—		-	-
0.03 0.05	0.03	0.02	0.04	0.08	0.03 0.05	0.09	_	0.01	0.03	0.04	0.82	0.20	0.01	_	0.02
0.14	0.08	0.21	0.20	0.10	0.11					0.04	0.10			_	
6.21	23.61	9.89	6.47	3.67	7.00	2.93	1.94	1.97	3.79	17.95	31.10	7.53		11.79	5.53
0.41	25.01	2.02	0.47	5.07	7.00	4.95	1.94	1.97	5.13	17.95	51.10	1.00	4.54		5.55

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Depth (mbsf)	1H-3, 145-150 4.5	2H-1, 80-89 7.0	3H-4, 120-125 21.4	4H-1, 55-74 25.8	5H-1, 51-69 35.3	6H-3, 120-125 48.4	9H-3, 120-125 76.9	12H-4, 120-125 106.9	13H-4, 145-150 116.7	14H-4, 145-150 126.2	15H-4, 120-125 135.4	16X-4, 145-150 145.2	17X-3, 145-150 148.7	18X-3, 120-125 158.0
Acidic														
Aspartic acid Glutamic acid	0.24 0.16	0.10	0.08 0.13	0.43 0.16	0.18 0.08	0.03 0.04	0.02	_	0.09 0.06	0.01 0.06	0.11 0.08	0.20	0.13 0.14	0.18
Basic														
Ornithine Lysine Histidine Arginine	0.21 0.08 0.11 0.01	0.13 0.06 0.06 0.03	0.07	0.68 0.23 0.16 0.00	0.41 0.12 0.09	0.09 0.05 0.05	0.14 0.05 0.04	0.12 0.04 0.01	0.29 0.09 0.09 0.02	0.20 0.04 0.04	0.02 0.09 0.08		0.35 0.07 0.07 0.01	0.48 0.14 0.18 0.02
Neutral														
Hydroxy Threonine Serine	0.21 0.67	0.11 0.29	0.05 0.11	0.33 1.44	0.12 0.65	0.19	0.03 0.28	0.08	0.08 0.57	0.02 0.26	0.11 0.52	=	0.14 0.43	0.21 0.88
Straight														
Glycine Alanine	0.91 0.67	0.51 0.32	0.37 0.27	1.76 1.11	0.90 0.58	0.40 0.34	0.47 0.33	0.22 0.22	0.73 0.48	0.48 0.36	0.75 0.47	=	0.66 0.46	1.25 0.69
Branch														
Valine Iso-leucine Leucine	0.35 0.04 0.18	0.01	0.01	0.11 0.47	0.03 0.10	0.01 0.36	0.01	0.01	0.04 0.10	0.02 0.31	0.03	111	0.01 0.07	0.28 0.07 0.47
Others														
Aspargine	\rightarrow	-	-	0.04	-	-	\sim -	-	-		0.07	-	0.09	0.13
Aromatic														
Tyrosine Phenylalanine Sulfur	=	1 1	-	Ξ	Ξ	-		1.1		-	-	Ξ		-
Cystein Methionine Taurine β-alanine	0.02 0.10	 0.18	 0.03	 0.10		Ē	1111	1111		 0.02	 0.02		 	F F F F
β -aminobutyric acid Total	$\frac{0.07}{4.03}$		$\frac{-}{1.12}$				1.37		2.64	1.82	2.35	0.20	2.65	4.98

Table 4. Concentrations of DFAA in interstitial-water samples from Hole 677A.

Table 4 (continued).

19X-4, 145-150 169.5	20X-4, 145-150 179.2	22X-3, 120-125 196.6	23X-4, 145-150 208.1	24X-4, 145-150 217.7	25X-4, 120-125 227.1	26X-4, 145-150 237.1	27X-4, 145-150 246.8	28X-3, 120–125 254.6	29X-3, 145-150 264.5	30X-3, 145-150 274.2	31X-4, 120-125 285.0	32X-4, 145-150 294.9	33X-2, 145-150 301.6	33X-4, 145-150 304.6	33X-3, 133-150 303.0
0.21 0.20	0.21 0.08	0.32 0.19	0.24 0.18	0.13 0.14	0.18			Ξ		0.35		Ξ		0.21	_
	0.40	0.63	0.53	0.35	0.48	0.47	0.38	0.89	0.65	2.37	0.70	0.70	0.55	1.17	0.13
_	0.12	0.15	0.13	0.07	0.14	0.13	0.10	0.23	0.17	0.33	0.17	0.21	0.14	0.03	-
	0.06	0.17	0.11	0.07	0.18	0.15	0.05	_	0.15	0.37	0.17	0.17	0.21	0.18	
0.02	0.01	0.02	0.03	0.01	0.02	_	0.02	-	-	0.04	<u></u>	0.02	0.03	-	-
0.25	0.24	0.39	0.22	0.14	0.21		_	_	_	0.36	—	0.75	_	0.17	
0.94	0.83	1.38	0.75	0.43	0.88	0.44	0.09		0.38	2.28	0.88	—	0.29	0.48	0.13
1.24	1.03	1.67	1.01	0.66	1.25	0.55	0.15	_	0.35	2.23	0.92	0.78	0.33	0.41	0.14
0.73	0.63	1.05	0.60	0.46	0.69	0.18	0.56	-	0.11	1.12	0.43	0.29	0.08	0.25	0.10
_	0.36	0.15	0.32	<u></u>	0.28	-	-	_	0.07	—	0.17	0.16	0.13	—	-
-	0.05	0.12	0.04	0.01	0.07	0.03			0.03	0.75	0.07	0.07	0.60	0.36	0.18
-	0.45	0.20	0.13	0.07	0.47	0.07	—		-	0.68	0.16	—	-	0.21	0.07
0.19	0.13	0.19	0.13	0.09	0.13	-		-	-	-	-	-	-	-	-
	_		-		_	7	<u>_</u>	-	_	0.64	0.46	0.24	_	0.15	-
-	_	-	1	-	-	0.07	0.10	0.06	0.11	0.25	-	0.15	0.02	0.03	-
-	_	-			-	\rightarrow			-	—	-	_	-	-	-
_	1		—		—	\rightarrow		$\sim - 1$		0.02	-	-	-		-
	0.01	0.02	-		-	0000		-		$\sim \rightarrow \sim$		_		-	100
0.02	0.02	0.02		0.02	-	-				—		-	-		100
0.03	- 4.63	- 6.67			1.00	—		—	-	0.03	0.10	_		-	-
3.83	4.63	6.67	4.42	2.65	4.98	2.09	1.45	1.18	2.02	11.82	4.23	3.54	2.38	3.65	0.75
					4.70		1.42		4.04					1.1.1.1.1	0.000

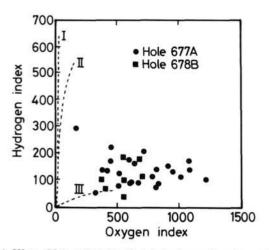


Figure 4. HI vs. OI (van Krevelen diagram) of organic matter at Holes 677A and 678B.

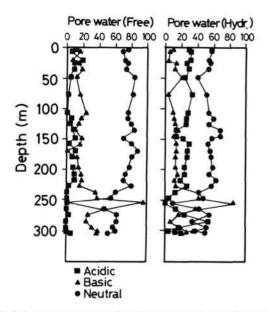


Figure 5. Relative abundances of acidic, basic, and neutral fractions of DFAA and THAA amino acids vs. sub-bottom depth in Hole 677A.

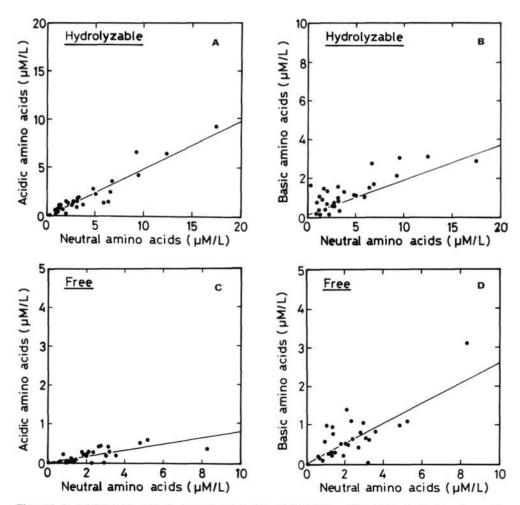


Figure 6. A. Acidic amino acids vs. neutral amino acids of THAA from Hole 677A. B. Basic amino acids vs. neutral amino acids of THAA from Hole 677A. C. Acidic amino acids vs. neutral amino acids of DFAA from Hole 677A. D. Basic amino acids vs. neutral amino acids of DFAA from Hole 677A.

	DFAA (mol%)	THAA (mol%)	Marine plankton ^a (mol%)	Marine suspension ^b (mol%)	River suspension ⁶ (mol%)
Acidic	8.2	22.6	20.4	25.3	24.9
Aspartic acid	2.9	4.8	9.0	16.7	13.1
Glumatic acid	5.2	17.7	11.5	8.6	11.8
Basic	22.4	19.0	17.4	14.5	3.0
Ornithine	15.3	12.8	1.0	0.1	0.9
Lysine	3.8	3.3	8.2	8.6	2.1
Histidine	3.0	2.0	2.2	1.9	
Arginine	0.3	0.9	6.0	3.9	
Neutral	66.7	51.5	48.8	51.6	61.2
Hydroxy					
Threonine	3.5	3.1	4.9	5.0	4.0
Serine	15.4	10.8	6.2	7.2	6.7
Straight			1.00	a 1 mar 1 mar 1 m	
Glycine	22.6	16.3	12.1	17.9	18.3
Alanine	15.2	7.5	8.5	9.6	14.8
Branch	1.0			5.0	6.2
Valine	1.8	4.3	5.5	5.0	
Iso-leucine	3.0	3.4	4.6	2.5	3.0
Leucine	4.2	5.8	7.1	4.4	8.3
Others Aspargine	1.0	0.2			
	1.0	2.4	6.0		2.5
Aromatic	1.9	2.6	6.8	5.2	3.5
Tyrosine	0.9	1.5	3.2	2.5	0.8
Phenylalanine	1.0	1.2	3.5	2.7	2.7
Sulfur	0.0	0.0	3.7	•	0.3
Cystein	0.0	0.0	1.1	*	•
Methionine	0.0	0.0	2.6	*	0.3

Table 5. The relative abundance of amino acids in interstitial-water DFAA and THAA, marine plankton, and suspended matter.

Note: * = Data not reported. ^a Raymont et al. (1975). ^b Below 3000 m water depth, equatorial Pacific Ocean (Ittekkot et al., 1984). ^c Average of river suspensions (Peake et al., 1972).