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THE EFFECT OF TEMPERATURE ON SALINITY-INDUCED CHANGES IN THE FREE AMINO ACID POOL OF MYA ARENARIA

A Thesis

Presented to

The Faculty of the School of Marine Science The College of William and Mary in Virginia

In Partial Fulfillment

Of the Requirements for the Degree of

Master of Arts

By William D. DuPaul 1968

APPROVAL SHEET

This thesis is submitted in partial fulfillment of the requirements for the degree of

Master of Arts

Author

Approved, August 1968

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Dester S. Haven, M.S.

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ABSTRACT

Warm and cold acclimated M. arenaria were transferred from sea water of 20 °/oo salinity to sea water of 30 °/oo salinity at 8, 18 and 25 C. The observed accumulation of NPS was not linear in relation with time and a three component process in the accumulation of NPS is proposed. The time lag between two of the components produced a change in the rate of NPS accumulation around the 36th hour. The increase in alanine concentration accounted for 80 to 90 % of the observed NPS increase. The high correlation between the decrease in aspartic acid and the increase in alanine indicates a direct relationship in the formation of alanine from aspartic acid. The rate-temperature functions of NPS accumulation were not the type expected for warm and cold acclimated poikilotherms. The reverse translation pattern of the rate-temperature curves indicates that warm acclimated animals can accumulate NPS at a greater rate than the cold acclimated ones. The conclusion made from data obtained through the use of temperature as an environmental variable is consistent with the hypothesis that an enzyme system mediates the supply of amino acids for isosmotic intracellular regulation.

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THE EFFECT OF TEMPERATURE ON SALINITY-INDUCED CHANGES IN THE FREE AMINO ACID POOL OF <u>MYA ARENARIA</u>

INTRODUCTION

The participation of small organic molecules in isosmotic intracellular regulation of many marine invertebrates was reviewed by Florkin and Schoffeniels (1965). They pointed out that, in marine invertebrates, the intracellular inorganic ion concentration is lower than the extracellular concentration and that the balance of osmotically active materials leading to an isosmotic concentration mainly consists of organic nitrogen compounds. A correlation between free amino acids (FAA) and ambient salinity consistent with the role of FAA in osmoregulation has been shown under laboratory conditions (Florkin and Schoffeniels, 1965). This correlation does not appear to be a laboratory artifact since high concentrations of total FAA in <u>Crassostrea virginica</u> correspond to high natural environmental salinities (Lynch and Wood, 1966).

The source of FAA and the mechanism for their accumulation has been widely discussed. On the basis of work done on <u>Eriocheir sinensis</u>, Schoffeniels (1960) concluded that the osmotically active amino acids are of intracellular origin and that the environmental osmotic pressure <u>per se</u> is not responsible for the amino acid concentration; Na or K is necessary for the accumulation process. Possible sources of FAA are: (1) FAA are actively taken up from the external medium, (2) a greater percentage of FAA and nitrogenous material normally excreted or released is retained, (3) the FAA are obtained from the digestive tract and, (4) other FAA pools contribute to the FAA pool involved in isosmotic intracellular regulation.

The effect of temperature on the FAA pool of marine animals is relatively unknown. Duchateau and Florkin (1955) reported for <u>Eriocheir</u> <u>sinensis</u> a decrease in proline at 2-3 Celsius (C) when compared to the same species kept at 10 C. More recently, Anders et al. (1962) found that a temperature change from 28 C to 22 C caused a rise in the level of tissue amino acids for the fish <u>Platypoecilus</u> and <u>Xiphophorus</u>. Rao (1963) found that the blood of the fresh water mussel, <u>Lamellidens</u> <u>marginalis</u>, showed an increase in total amino acids after warm acclimation (35 C) compared to cold acclimated (20 C) ones. Saroja and Rao (1965) reported that the body fluid FAA in the earthworm <u>Lampito mauritii</u> showed a decrease of 142.3 % in warm acclimated animals (35 C) when compared to animals kept at normal conditions (28 C).

The purpose of this research was to determine the effect of temperature on the time course of the increase of FAA caused by an increase in salinity. The experiments are designed to permit characterization of the rate-temperature functions according to Precht (1958) and Prosser and Brown (1961). The determination of temperature effects on 18 identified amino acids found in <u>Mya arenaria</u> Linne may also help elucidate the mechanism of accumulation of FAA in a more definitive manner.

METHODS AND MATERIALS

Animals

Specimens of <u>Mya arenaria</u> were collected from a single intertidal mud-flat on the York River (Virginia). At the time of collection (September 1967), water temperatures ranged from 20 to 23 C. One hundred and ninety animals were selected for size uniformity (62 ± 6 mm) and kept in a laboratory table of running sea water with a salinity of 20.4 \pm 0.2 ^o/oo and a temperature of 21.0 \pm 1.0 C for 8 days prior to temperature acclimation.

Two groups of animals were kept at 25 C and 8 C for 8 days and, in regard to their general metabolism, are referred to as warm and cold acclimated. Acclimation is virtually complete for most aquatic animals in a few days (Prosser and Brown, 1961, p. 242). Experiments consisted of placing warm and cold acclimated animals in glass battery jars containing 30 $^{\rm O}$ /oo natural sea water at temperatures of 25, 18 and 8 C. Samples were taken every 12 hours of the experiment for a duration of 60 to 84 hours. Other groups of <u>M</u>. <u>arenaria</u> were kept at temperatures of 30, 20 and 8 C for various lengths of time. These groups supplied additional information and served as experimental controls. Table I summarizes the acclimation, experimental and control conditions.

Every 12 hours during the experiment, the animals were given a mixture of algae consisting of <u>Monochrysis</u>, <u>Isochrysis</u> and <u>Phaeodactylum</u>. Continuous aeration and daily changes of sea water prevented an excessive buildup of waste products.

Tissue Preparation

Two portions of the anterior adductor muscle were dissected from <u>M. arenaria</u>. One portion, ca. 120 mg, was blotted, weighed, dried at 90 C for 60 hours (to constant weight), and reweighed to determine the water content. The other portion (<100 mg) was blotted, weighed, and placed in 2 ml of 80 % ethanol to extract the ninhydrin positive substances (Awapara, 1948). The per cent water content obtained for one portion of the adductor muscle was used to calculate water content and dry weight for the extracted tissue.

Ninhydrin Positive Substances (NPS)

NPS were determined from an aliquot (0.1 ml) of the ethanol extract by the method described by Clark (1964). Each determination was done in duplicate. A standard of glycine was serially diluted (0.01 to 0.15 μ M/ml) and was analyzed with each group of tissue extracts. Colorimetric determinations were made at 570mµ using a Bauch and Lomb Spectronic 20 colorimeter.

Values of NPS as glycine equivalents are expressed in µM/g tissue water and statistical significance at the 95 % level was determined by analysis of variance "t" test and multiple range analysis (Snedecor, 1960). Values for NPS were computed using the mean per cent tissue water (75.2 %) of 36 animals from an environmental salinity of 20.4 ^O/oo and a temperature of 21.0 C. These were the conditions in the laboratory sea water tables prior to acclimation and establishment of experimental conditions. By using values corrected for changes in tissue water, changes in the concentration of NPS due to fluctuations in tissue water are eliminated. Thus values for NPS, unless otherwise stated, are net values corrected for water loss. Any change in value represents a real change in the concentration of NPS and is not affected by tissue water changes. This correction for water loss does not change the character of the NPS accumulation as a process, but only decreases the change in relative amounts and the variability between individual animals.

Individual Amino Acids

A qualitative and quantitative analysis of amino acids was made using an automatic ion exchange analyzer (Technicon Auto-Analyzer ^R). An aliquot (0.1 ml) of the ethanol extract was mixed with 0.2 ml 0.1 N HCl and 0.25 uM L-amino guanido propionic acid was added as an internal standard. This mixture was dried, then redissolved with 2.0 ml 0.1 N HCl and 0.25 uM Norleucine added to serve as a second internal standard; samples were then lyophylized. The samples were redissolved in 0.1 N HCl and injected under N₂ pressure into a 0.6 cm x 129 cm heated (60 C) glass column filled with Technicon Chromobeads B ^R. A gradient of sodium citrate buffers (pH 2.875 to 5.000) with a flow rate of 0.5 ml/min served as an eluent.

The color developed with ninhydrin was determined at 440 and 570 mu through 15-mm light path continuous flow cuvettes. Amino acids were identified by comparing unknown peaks with those of a known 21 amino acid standard solution (General Biochemicals, Chagrin Falls, Ohio).

As with the NPS, all amino acid values are calculated on the basis of 75.2 % tissue water and are corrected for loss of tissue water.

RESULTS -

After temperature acclimation, the warm acclimated animals showed a significant change in tissue water with a mean value of 73.3 % as compared to 75.2 % in unacclimated animals. Cold acclimated animals showed no significant change in tissue water content. The loss of tissue water under experimental conditions did not account for all the increase in NPS concentration but was responsible for 4-11 % of the change.

There were no significant differences in NPS among warm, cold and unacclimated animals with or without correction for tissue water change. Total NPS values for 19 M. <u>arenaria</u> before temperature acclimation ranged from 232.6 to 353.3 μ M/g tissue water. NPS for cold acclimated animals ranged from 286.0 to 344.6 μ M/g tissue water and for warm acclimated animals, 294.5 to 320.2 μ M/g tissue water. Values for warm and cold acclimated animals were pooled to obtain a common baseline for a comparison to conditions of increased salinity.

NPS values obtained at both acclimation conditions and conditions of increased salinity are summarized in Table II. There were definite increases in NPS concentration over the time course of the experiment. Figures 1 and 2 show an initial increase of NPS concentration which reached a peak at 12 or 24 hours depending upon experimental conditions; the greater the temperature change, the sooner this peak was reached. With the exception of cold acclimated animals at 8 C, a significant drop in NPS concentration occurred at 36 hours only to be followed by

a second increase. This second increase was also influenced by temperature but, unlike the initial increase, higher levels and a greater rate of NPS accumulation occurred with warmer temperatures regardless of acclimation temperature. This latter pattern of response remains essentially the same for three groups of animals each acclimated to a different experimental temperature (Fig. 3).

The results of two- and three-way analysis of variance for the NPS values obtained at various experimental conditions are presented in Table III. The warm and cold acclimated groups and the group of animals acclimated to each experimental temperature showed significant differences in NPS concentration for time, temperature and for interaction between time and temperature. In the analysis in which only the warm and cold acclimated groups were combined, significant differences in NPS concentration remained for time, temperature and for interaction between time and temperature. Significant differences at the 5 % level between adjacent times and temperatures were tested by use of multiple range analysis. The results of these tests are included in Table II.

The 18 identified amino acids in <u>M</u>. <u>arenaria</u> accounted for 69 to 95 % of the measured NPS. In Table IV, the quantities of the 18 FAA are presented for acclimated and unacclimated conditions. Glycine and alanine made up the greatest part of the FAA pool and constituted 70 to 76 % of the total. Other FAA in order of decreasing concentrations were taurine, glutamic acid, arginine, aspartic acid and serine, each contributed 1 to 9 % of the total.

There were no statistical differences in the total of the 18 identified amino acids when the warm and cold acclimated groups were compared with the unacclimated group. However, there were statistical differences in the total of the 18 amino acids between the warm and cold acclimated animals. In addition to the 18 identified FAA, the chromatogram showed other peaks that are, at this time, unidentified. Values ranged from 7.99 to 24.86 µM/g tissue water (norleucine equivalents). These unidentified materials constituted about 5 % of the measured NPS. There were no statistical differences between the three groups of animals when the combined totals of the unknown and known amino acids were considered.

Because of the large variability in the concentration of individual FAA, no significant differences were detected when the unacclimated animals were compared to the warm and cold acclimated animals. However, there were significant differences in two important FAA between warm and cold acclimated animals. In the warm acclimated group the mean values of taurine and glycine were lower than in the cold acclimated group by 12.05 and 27.30 μ M/g tissue water respectively. Although the majority of the values presented in Table V are from only one chromatogram, some idea as to the variability of each FAA is obtained at time zero for warm and cold acclimated animals (Table IV). Also, the values for warm acclimated animals at 18 C (36 hours) represent the mean and standard deviation for three individuals (Table V).

FAA totals showed definite changes over the time course of the experiment (Table V) and the pattern of accumulation was similar to that of the NPS accumulation.

Of the individual FAA, glycine and alanine were the most concentrated at experimental conditions. Alanine made up 33.7 to 59.8 % of the FAA pool while glycine made up 22.2 to 48.9 % of the pool. The response of glycine to an increase in salinity was variable. There was a slight increase in the concentration of glycine but, its mole percentage of

the FAA pool decreased over the time course of the experiment. Data for glycine is presented in Table V.

Over the time course of the experiments, the mole % of alanine in the FAA pool increased for warm and cold acclimated animals. The mole % of alanine in the FAA pool was significantly different for both groups of animals at each of the three experimental temperatures; by the end of the experiment, higher experimental temperatures resulted in higher percentages of alanine in the FAA pool. For the warm acclimated animals between 12 and 36 hours of the experiment, there was an inversion in the quantities of alanine in relation to temperature and the curves demonstrated a change in slope (Fig. 5a). This is similar to the behavior pattern of the total FAA and NPS in relation to time and temperature (Fig. 1, 4a). The alanine in the cold acclimated animals presented a somewhat similar response to experimental conditions. The alanine content of the animals at 18 C demonstrated a pronounced two-step increase in concentration whereas at 8 and 25 C there appeared to be a simple increase and no noticeable change in the slope of the curve (Fig. 5b).

Other amino acids such as taurine, glutamic acid and arginine each made up 2 % or more of the FAA pool. The data for these amino acids are presented in Table V. There was a slight but noticeable increase in the quantities of serine and phenylalanine during experimental conditions. The concentration of serine approximately doubled from its baseline value with values as high as $6.51 \,\mu$ M/g tissue water but, it seldom was responsible for more than 2 % of the FAA pool. The increase in phenylalanine was less pronounced and more erratic than that of serine and was consistently less than 2 % of the FAA pool. The remaining FAA, cysteic acid, methionine sulfoxide, proline, valine, isoleucine, leucine, tryptophane, ornithine, lysine, histidine and threonine showed little if any change at experimental conditions and each contributed less than 2 % of the FAA pool. The only amino acid that consistently decreased in concentration with time during the experiments was aspartic acid. Initial values ranged from $17.38 \pm 3.65(3)$ to $11.87 \pm 4.39(3)$ µM/g tissue water for warm and cold acclimated animals respectively. These values dropped to between 1.10 and 5.52 µM/g tissue water depending upon experimental conditions. Data for aspartic acid is presented in Table V and Fig. 6a, b. Table I. Summary of acclimation and experimental conditions.

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- Table II. NPS (μ M/g tissue water) in the anterior adductor muscle of <u>M</u>. arenaria after transfer from 20 to 30 ^O/oo sea water. Values are the mean and SD for three animals except for time zero where N=19.
 - (I) No significant differences at the 95 % level from previous time values.
 - (II) No significant differences at the 95 % level between experimental temperatures.

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$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	322.7 <u>+</u> 24.0 391.9 <u>+</u> 34.3 333. (I)	391.9 <u>+</u> 34.3 333.	333.	6 <u>+</u> 18.9	352.7 <u>+</u> 15.8	382.1 <u>+</u> 8.9 (II)	371.8 <u>+</u> 8.4 (II)	386.8 <u>+</u> 29.5
$17.3 \qquad 359.9\pm12.3 \qquad 417.6\pm6.3 \qquad 423.5\pm12.7 \qquad 408.46\pm6.9 \\ (I) \qquad (I) \qquad (II) \qquad (II) \qquad (II) \qquad \\ 381.6\pm29.8 \qquad 410.4\pm4.0 \qquad 425.8\pm17.9 \\ (II) \qquad (II) \qquad (II) \qquad (II) \qquad \\ (II) \qquad (II) \qquad (II) \qquad (II) \qquad \\ 384.3\pm16.4 \qquad 448.9\pm4.6 \qquad 451.7\pm17.3 \\ (I) \qquad (II) \qquad (II) \qquad (II) \qquad \\ \end{array}$	325.8 <u>+</u> 22.9 368.8 <u>+</u> 22.2 318.5 (I) (I)	368.8 <u>+</u> 22.2 318.5 (I)	318.5	5±17.6	350.7 <u>+</u> 3.8 (I) (II)	349.3 <u>+</u> 12.1 (II)	357.9 <u>+</u> 26.6 (I) (II)	364.3 <u>+</u> 3.5
21.6 381.6 <u>+</u> 29.8 410.4 <u>+</u> 4.0 425.8 <u>+</u> 17.9 (II) (II) 384.3 <u>+</u> 16.4 448.9 <u>+</u> 4.6 451.7 <u>+</u> 17.3 (I) (II) (II)	341.6 <u>+</u> 2.0 372.5 <u>+</u> 10.4 412.6 (I)	372.5 <u>+</u> 10.4 412.6 (I)	412.6	5±17.3	359.9 <u>+</u> 12.3 (I)	417.6 <u>+</u> 6.3 (II)	423.5 <u>+</u> 12.7 (II)	408.46 <u>+</u> 6.9
384.3 <u>+</u> 16.4 448.9 <u>+</u> 4.6 451.7 <u>+</u> 17.3 (I) (II) (II)	358.5 <u>+</u> 24.4 361.9 <u>+</u> 4.4 392.9 (II) (I) (I) (I) (I)	361.9 <u>+</u> 4.4 392.9 (I) (I) (I)	392.9 (I))	381.6 <u>+</u> 29.8	410.4 <u>+</u> 4.0 (II)	425.8 <u>+</u> 17.9 (II)	
					384.3 <u>+</u> 16.4 (I)	(II)	451.7 <u>+</u> 17.3 (II)	

Table III. Two- and three-way analysis of variance on NPS data.

	Source	Mean	Degrees of	Level of Significance
		Square	Freedom	(%)
Warm acclimated	Time	12215.78	ک	0.5
	Temperature	1742.15	2	10.0
	Time x Temperature	1433.45	10	2.5
Cold acclimated	Time	5350.41	'n	0.5
	Temperature	3463.18	2	0.5
	Time x Temperature	1532.71	10	0.5
Warm and Cold				
Acclimated	Time	16438.81	S	0.5
	Temperature	5022.01	2	0.5
	Acclimation	15950.52	1	0.5
	Time x Temperature	2787.26	10	0.5
	Time x Acclimation	183.55	Ŷ	NS
	Temperature x Acclimation	1127.38	2	NS
	Time x Temperature x Acclimation	179.90	10	NS
Acclimated at Each experimental				
Temperature	Time	7461.81	S	0.5
	Temperature	8533.60	2	0.5
	Time x Temperature	855.05	10	5.0

Table IV. FAA (μ M/g tissue water) in the anterior adductor muscle of <u>M</u>. <u>arenaria</u> before and after temperature acclimation. Each value represents mean and SD.

Free amino	Before accl	imation	Warm accli 25 C	mation	Cold accli 8 C	mation
	N=8	Mole ⁰ /o	N=3	Mole ⁰ /o	N=3	Mole ⁰ /a
Cyste ic acid	1.62 <u>+</u> 0.57	0.66	2.04 <u>+</u> 0.74	0.90	1.40+0.20	0.56
Taurine	16.05 <u>+</u> 4.76	6.59	11.05 <u>+</u> 5.62	4.86	23.10 <u>+</u> 3.59	9.31
Methionine sulfoxide	0.38 <u>+</u> 0.23	0.15	0.74 <u>+</u> 0.20	0.33	0.26*	0.10
Aspartic acid	11.77 <u>+</u> 4.90	4.88	17.38 <u>+</u> 3.65	7.66	11.88 <u>+</u> 4.38	4.78
Threonine	2.41 <u>+</u> 0.88	0.98	3.93 <u>+</u> 0.90	1.73	1.91 <u>+</u> 0.06	0.77
Serine	2.95<u>+</u>1.19	1.21	1.86 <u>+</u> 0.66	0.82	2.51 <u>+</u> 0.66	1.01
Glutamic acid	14.59 <u>+</u> 2.46	6.13	13.91 <u>+</u> 1.44	6.12	13.90 <u>+</u> 1.92	5.61
Glycine	86.00 <u>+</u> 22.61	35.31	7 3.68<u>+</u>9.7 0	32.43	10 0.98<u>+</u>1. 46	40.70
Alanine	90.58 <u>+</u> 14.04	37.19	85.03 <u>+</u> 11.90	37.43	85.81 <u>+</u> 19.01	34.59
Valine	0.56 <u>+</u> 0.31	0.23	0.85 <u>+</u> 0.32	0.37	0.49*	0.19
Isoleucine	0.42 <u>+</u> 0.25	0.17	0.54 <u>+</u> 0.25	0.24	0.28*	0.11
Leucine	0.59 <u>+</u> 0.34	0.24	0.68 <u>+</u> 0.13	0.30	0.45 <u>+</u> 0.21	0.18
Tyrosine	0.43 <u>+</u> 0.36	0.17	0.72 <u>+</u> 0.13	0.32	,0.30*	0.12
Phenylalanine	2.08 <u>+</u> 0.73	0.85	2.21 <u>+</u> 0.56	0.97	1.55 <u>+</u> 0.37	0.62
Ornithine	1.44 <u>+</u> 1.30	0.59	0.97 <u>+</u> 0.16	0.43	0.40*	0.16
Lysine	1.51 <u>+</u> 0.77	0.62	1.97 <u>+</u> 0.35	0.87	1.31 <u>+</u> 0.81	0.52
Tryptophan	1.34 <u>+</u> 0.85	0.55	1.25 <u>+</u> 0.46	0.55	0.50*	0.20
Histidine	0.45 <u>+</u> 0.19	0.18	0.25 <u>+</u> 0.12	0.11	0.20*	0.08
Arginine	9.01 <u>+</u> 2.19	3.28	8.47 <u>+</u> 0.71	3.73	7.52 <u>+</u> 0.63	3.03
Total % of NPS	247.99 <u>+</u> 39.35 85.45 <u>+</u> 7.61		227.16 <u>+</u> 3.41 73.63 <u>+</u> 1.51		248.0 <u>6+</u> 17.20 86.07 <u>+</u> 5.67	
Total plus unidentified					651 10110 OF	
p eaks ≠ Z of NPS	261.51 <u>+</u> 35.85 90.49 <u>+</u> 8.12		242.49 <u>+</u> 3.65 79.70 <u>+</u> 3.75		90.03 <u>+</u> 8.06	

* estimated from peaks too small to measure accurately # ammonia excluded

Table V. FAA (µM/g tissue water) in the anterior adductor muscle of <u>M</u>. <u>arenaria</u> over the time course of the experiment. Values where N=3 are the mean and SD; where N=2, values are the mean of two animals.

Time (bours	· _ ·	Total amino acids (plus	Glycine	Mole Z	Alanine	Mole %	Taurine	Glut ani c Acid	Arginine	Aspartic Acid
Vera		242.49	78.68	32.44	85.03	35.06	11.05	13.91	8.47	17.38
accl. 25 C	0	<u>+</u> 3.65	1 9.79		<u>+</u> 11.90		<u>+</u> 5.26	4.4	- <u>+</u> 0.71	±3.65
8 C				•						
Ĩ	12	395.35	126.08	31.85	136.53	34.53	16.50	20.98	9.63	12.08
	. 9E	302.92	87.69	28.51	131.50	43.41	23.40	15.53	10.11	6.36
	60	334.37	100.75	30.13	146.24	43.74	18.56	15.93	10.28	6.80
	7 8	350.93	105.64	26.02	160.69	45.78	14.88	18.37	10.34	5.52
18 C	13	295.10	91.10	29.55	130.51	42.35	10.70	15.73	8.53	3.47
1	26	100.90	87.64	28.28	142.35	45.93	16.61	12.00	8.82	2,66
N=3	36	296.05	73.09	24.69	130.98	44.24	27.21	15.32	7.46	3.14
•		+7.30	+6.97		+11.39		+7.34	+2.55	+1.85	+1.34
	48	361.61	99.45	27.49	$\overline{1}73.22$	47.89	12.76	22.04	10.07	2.64
	60	339.30	93.80	27.64	154.46	45.52	23.79	19.36	11.21	2.00
-	78	351.75	91.82	26.10	180.83	51.41	10.68	20.15	12.44	2.72
25 C	5	01 626	50 111	F0 07	0/. 95	30.95	19 27	10 31	5 07	50.6
	12	21.212	00 57	10°01	717 43	00.40 00 00	70.7T	11 16	a/ a	50°7
j	24 26	202 212	10.00	20.25	118 70	00.00 00 7 c	40 CV	10 75	0.40 15	در.ر ۱ ۱
		0/.010	00.00	10.00	41.011	59 10	31 61	10.01	10.15	+1·1
	9 3	400.30	88.66	22.15	239.42	59.81	10.10	14.01	5.82	1.63
Cold		254.40	100.98	39 • 69 ́	85.81	33.73	23.10	13.90	7.52	11.87
8 C	0	±10.03	0 1 .11		119.01		60.61	±1.92	Fa•07	66.97
່ ບ ອ	24	266.66	89.38	33.52	91.09	34.13	27.54	15.11	9.95	9.68
N=2	36	298.09	95.06	31.88	119.28	40.01	27.60	16.28	8.02	5.39
	48	266.84	86.83	32,54	115.74	43.37	10.47	12.53	7.98	5.64
	9.	336.85	123.34	36.62	138.21	41.03	21.22	17.32	10.01	3.68
18 C		161 01	14 041	36 10	16 21	20 C7	16.06	16 26		
	37	10.114	14.021	26 00	10.4C1	00.04	10.90 23 07	70°71		1.03
	1 ig 7 f	00.000	04-1C1	20.99 27 08	CD.CC1	. 00 07	10.20	17 01	12.90	7 C
	87	359.98	131.95	37.20	140.11	39.50	19.04	14.86	10.82	1.59
	09	387.21	106.19	28.01	195.27	51.17	21.74	19.85	11.26	1.52
25 C										
ì	12	292.96	114.49	39.08	120.58	41.16	18.82	10.37	10.34	3.67
Ĺ	0 d	1000 100 c3 c	110.02	54.35 00 cc	157.03	41.11	20.02	05.21	00	4.43
	09	330.66	78.80	23.83	171 58	40.J/	11.11	13.40	9.14	

Figure 1. Accumulation of NPS (μ M/g tissue water) in the anterior adductor muscle of warm acclimated <u>M</u>. <u>arenaria</u> after a transfer from 20 to 30 ^O/oo sea water.



Figure 2. Accumulation of NPS (μ M/g tissue water) in the anterior adductor muscle of cold acclimated <u>M</u>. <u>arenaria</u> after a transfer from 20 to 30 ^o/oo sea water.



Figure 3. Accumulation of NPS (uM/g tissue water in the anterior adductor muscle of M. arenaría after a transfer from 20 to 30 ^o/oo sea water. Each group of animals was acclimated to each experimental temperature.



Figure 4. Accumulation of total FAA including unknown chromatogram peaks (μ M/g tissue water) in the anterior adductor muscle of <u>M</u>. arenaria after a transfer from 20 to 30 °/oo sea water.

- A. Cold acclimated.
- B. Warm acclimated.



Figure 5. Accumulation of alanine (μ M/g tissue water) in the anterior adductor muscle of <u>M</u>. <u>arenaria</u> after a transfer from 20 to 30 ^o/oo sea water.

- A. Warm acclimated.
- B. Cold acclimated.



Figure 6. Decrease of aspartic acid (µM/g tissue water) in the anterior adductor muscle of <u>M</u>. <u>arenaria</u> after a transfer from 20 to 30 ^o/oo sea water. A. Warm acclimated.

B. Cold acclimated.



DISCUSSION

Several authors have demonstrated that there is a linear relationship between salinity and NPS or total FAA in muscle tissue of the marine pelecypods <u>Mytilus edulis</u> (Lange, 1963), <u>Crassostrea virginica</u> (Lynch and Wood, 1966) and <u>Mya arenaria</u> (Virkar and Webb, unpublished data). <u>M. arenaria</u> accumulates NPS and total FAA in response to a salinity increase in a fashion that is not linear with time. Except for cold acclimated animals at 25 C, the accumulation of NPS and total FAA changed in rate or significantly decreased in absolute concentration around the 36th hour (Figs. 1, 2, 3 and 4). The exception for cold acclimated animals at 25 C may be an artifact of the 12 hour sampling interval.

The patterns of NPS and total FAA accumulation indicate that there may be multiple steps involved in this process. The slight transitory increase in the glycine concentration during the first 24 hours of experimental conditions along with a more pronounced increase in the concentration of alanine accounted for more than 80 % of the NPS increase. Also, the slight transitory increase of the FAA pool as a whole during the first 12 to 24 hours contributed to the observed initial increase in NPS concentration. The change in the rate of NPS increase that occurred around the 36th hour was caused either by the decrease or lack of increase in the concentration of alanine. After 36 hours, the concentration of alanine increased sharply and accounted for approximately 90 % of the NPS increase. The interpretation of multiple steps in NPS accumulation

is valid as shown by the significant interaction between time and temperature in the analysis of variance (Table III) and multiple range analysis (Table II).

From the above analysis, it is apparent that there may be at least two major processes involved in the accumulation of NPS; a "fast component" that operates during the first 24 hours after a salinity increase and a "slow component" that obtains its effective rate after 36 hours. Since <u>M. arenaria</u> is an osmoconformer (Hegemann, 1964), the fast component may be useful in preventing excessive tissue water loss and cell volume changes. Data on adductor muscle tissue water indicate that no further significant water loss occurs after 24 to 36 hours of exposure to an increase of salinity. This is consistent with the idea that the active adjustment of the intracellular osmotic pressure to new osmotic pressure could prevent a change in hydration of the cells (Jeuniaux, Bricteux-Gregoire and Florkin, 1961) and effect cell volume regulation (Lange, 1968).

There was a transitory increase in the concentration of taurine around the 36th hour of experimental conditions which coincided with the general decrease of the other osmotically active substances. Since the initial concentration of taurine was lower in warm acclimated animals than in cold acclimated ones, the increase was more noticeable in the warm acclimated group. This is interpreted as another indication of the presence of multiple components in the accumulation process of osmotically active substances. The increase was not large (7-20 μ M/g tissue water) and appeared to be acting in response to the decrease or lack of increase in the concentration of alanine at the time when the proposed fast and slow components are in a transition stage.

Three proposed components to account for the observed accumulation of FAA in M. arenaria are presented in Fig. 7. The proposed fast component (A) may be partly from the release of osmotically active FAA from an osmotically inactive form. During the first 36 hours of the experiment, this supply of osmotically inactive amino acids is exhausted. This, combined with the continual loss or leakage of FAA from the osmotically active pool, produces a net loss or decrease in the concentration of FAA in the cell around the 36th hour. At approximately the same time, the long ranged slow component (B) begins to supply FAA faster than they are lost as noted by the sharp net increase in the concentration of alanine. The time lag between these two components produces the change in slope in the accumulation of osmotically active FAA (D). The initial peak and subsequent decrease in the concentration of FAA is not interpreted as an overshoot phenomenon because the 12 to 24 hour concentration is considerably lower than the final concentration. A third component (C) is the transitory increase in the concentration of taurine at about 36 hours. Without this increase, the drop in total osmotically active FAA would be even greater. It is possible that, in M. arenaria, taurine may be used as an auxiliary and become an important osmotically active substance when other FAA are not available in sufficient quantities to accomplish isosmotic intracellular regulation. In contrast, taurine has a major role in the isosmotic intracellular regulation at increased salinities in Mytilus edulis (Lange, 1963) and Crassostrea virginica (Lynch and Wood, 1966). Lange (1963) reported that <u>Mytilus edulis</u> establishes its high concentration of NPS primarily by an increase in taurine and in this regard, exerts a sparing action on the animals use of "essential" amino acids. This may also be true for <u>Crassostrea</u> virginica, as Lynch and Wood (1966) showed that taurine is the most concentrated amino acid

above a salinity of 19 o /oo and there the concentration of glycine, alanine and glutamic acid remains relatively constant.

On the basis of experiments performed on isolated nerves of Eriocheir sinensis it was concluded that the amino acids contributing to the total osmotic pressure are of intracellular origin (Schoffeniels, 1960). Gilles and Schoffeniels (1966) identified an aspartate decarboxylase in the ventral nerve chain of Homarus vulgaris. They concluded that L-alanine synthesis depends on at least two major pathways: (a) a transamination of pyruvate and (b) a decarboxylation of aspartate. The decarboxylation of aspartic acid to alanine has also been reported for microorganisms (Mahler and Cordes, 1966). The decarboxylation of aspartic acid could be an important pathway in the supply of alanine as an osmotically active substance in the muscle tissue of M. arenaria. At all experimental temperatures there was a close relationship between the decrease of aspartic acid and the increase in the concentration of alanine. The correlation coefficients for the relationship between aspartic acid and alanine concentrations in five of six experiments range from r = -0.74to r = -0.98. In the sixth experiment with warm acclimated animals at 25 C, the correlation disappears as the concentration of alanine exceeds 200 μ M/g tissue water and the aspartic acid appears to have a lower limit of 1.10 µM/g tissue water. Nevertheless, the correlation indicates a direct pathway in the formation of alanine from aspartic acid. The fact that the decrease of aspartic acid is less at 8 C than at 18 and 25 C (Fig. 6a, b) is an indication that the conversion of aspartic acid to alanine is temperature sensitive. These data are in agreement with the general hypothesis presented by Schoffeniels (1967) that the increased intracellular content of osmotically active amino acids

after a transfer to a hypertonic medium is ascribed partly to a decrease in the breakdown or loss and partly to the increase in the synthesis of amino acids.

The rate-temperature function for NPS accumulation during the first 24 hours of experimental conditions (fast component) does not follow the standard patterns for warm and cold acclimated poikilotherms as described by Prosser and Brown (1962) and Precht (1958). When the warm and cold acclimated \underline{M} . <u>arenaria</u> experienced a temperature change, the rate of NPS accumulation increased regardless of the direction of the temperature change (Fig. 8a). The data for alanine indicate that the amino acid follows the pattern of accumulation similar to that of the NPS. The reason for the rate-temperature inversion is unknown at this time, but it may be the result of the additional stress caused by the sudden temperature change.

Intertidal populations of \underline{M} . <u>arenaria</u> in an area subject to fresh water runoff could be exposed to simultaneous salinity and temperature fluctuations. The fast component of NPS accumulation may serve as a response to both a temperature and salinity stress but it must be kept in mind that, in the range tested, temperature alone does not elicit this response. Unpublished data on experimental control animals show a slight but significant increase in NPS only when the temperature reaches 30 C; the changes occur around the 48th hour of control conditions. The concentration and composition of the NPS pool remains relatively stable within the temperature and time limits of these experiments (8 to 25 C for approximately 10 days).

The rate-temperature functions of NPS accumulation in Figure 8b were constructed from data between time zero and 60 hours. The resulting

pattern of reverse translation (e.g. the curve of the cold acclimated animals lies to the right and below that of warm acclimated ones) may be the result of other factors complicating the response (Prosser and Brown, 1962). This is the case for the rate-temperature functions of NPS accumulation as it was affected by two changed parameters, temperature and salinity. The combined effect of each environmental condition presents rate-temperature functions which do not correspond to the normal patterns for poikilotherms (Prosser and Brown, 1962) and subsequent interpretations may not be applicable. An arithmetic plot of NPS accumulation is presented in Figure 9. The optimum temperature range for NPS accumulation is different for warm and cold acclimated animals. Warm acclimated animals had a greater accumulation rate at 18 and 25 C than at 8 C. The accumulation rates at 18 and 25 C were not appreciably different. The rate of NPS accumulation for cold acclimated animals was greatest at 25 C and was different from the rate at 18 C. There were no appreciable differences in the accumulation rates between 8 and 18 C. It would appear that temperatures above 25 C are optimum for NPS accumu**lation** in cold acclimated animals but preliminary studies indicate that under these experimental conditions higher temperatures have lethal effects on the animals.

The effect of temperature on the rate of NPS accumulation, which is the type expected for an enzyme mediated pathway within certain temperature limits, is another indication that the supply of amino acids for isosmotic intracellular regulation in <u>M. arenaria</u> is controlled by an enzyme system. This is in agreement with a statement by Lange (1968) that the size of the FAA pool in the cell is at least partly controlled by the activity of amino acid synthesizing enzymes. The reason why the warm acclimated animals have a faster rate of NPS accumulation than cold acclimated ones at the same temperature is unknown at this time. One reason may be that the amino acid synthesizing enzymes have to be manufactured to meet the demand of isosmotic intracellular regulation. If this is so, the warm acclimated animals having a more active metabolism would probably be able to manufacture the enzymes needed faster than the cold acclimated animals.

In conclusion, data obtained through the use of temperature as an environmental variable is consistent with the hypothesis that an enzyme system mediates the synthesis of amino acids for isosmotic intracellular regulation. The indications that alanine is synthesized from aspartic acid in the muscle tissue of \underline{M} . <u>arenaria</u> imply that the major osmotically active amino acid, alanine, is neither obtained by active uptake from the environment nor supplied completely from sources of osmotically in-active amino acids. However, these possibilities as well as others must be considered as a source of osmotically active amino acids especially for other marine invertebrates.

Figure 7. Proposed multiple components in the accumulation of NPS and FAA in M. arenaria.

- A. Fast component.
- B. Slow component.
- C. Taurine component.D. Composite of multiple component process.



- Figure 8. Rate-temperature function of NPS accumulation for warm

 - and cold acclimated M. arenaria.
 A. Initial increase (fast component).
 B. Accumulation over entire time course of the experiment.



Figure 9. Rate-temperature of NPS accumulation for warm and cold acclimated <u>M</u>. <u>arenaria</u> over the entire time course of the experiment.



SUMMARY

- The accumulation of FAA and NPS in the adductor muscle of <u>M</u>. <u>arenaria</u> in response to an increased salinity was not linear with time.
- Three components in the process of NPS and FAA accumulation were proposed: (1) a fast component active during the first 24 hours,
 (2) a slow component effective from 36 hours and (3) a taurine component effective during the time lag of the fast and slow components.
- The increase in the alanine concentration accounted for 80 to 90 % of the observed increase in NPS concentration.
- 4. The high correlation between the decrease in the concentration of aspartic acid and the increase in the concentration of alanine indicates a direct relationship in the formation of alanine from aspartic acid. It is suggested that aspartic acid undergoes a decarboxylation forming alanine.
- 5. The rate-temperature functions of NPS accumulation for warm and cold acclimated <u>M</u>. <u>arenaria</u> did not conform to standard patterns described for warm and cold acclimated poikilotherms.

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