

An Estuarine Benthic Index of Biotic Integrity for the Mid-Atlantic Region of the United States. I. Classification of Assemblages and Habitat Definition

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ABSTRACT: An objective of the Mid-Atlantic Integrated Assessment Program (MAIA) of the U.S. Environmental Protection Agency is to develop an index for assessing benthic community condition in estuaries of the mid-Atlantic region of the United States (Delaware Bay through Pamlico Sound). To develop such an index, natural unimpaired communities must first be identified and variability related to natural factors accounted for. This study focused on these two objectives; Llansó et al. (2002) describe the index. Using existing data sets from multiple years, classification analyses of species abundance and discriminant analysis were employed to identify major habitat types in the MAIA region and evaluate the physical characteristics that structure benthic infaunal assemblages. Sampling was restricted to soft bottoms and to the index development period, July through early October. The analyses revealed salinity and sediment composition as major factors structuring infaunal assemblages in mid-Atlantic estuaries. Geographical location was a secondary factor. Nine habitat classes were distinguished as a combination of 6 salinity classes, 2 sediment types, and the separation of North Carolina and Delaware-Chesapeake Bay polyhaline sites. The effect of sediment types on faunal assemblages was restricted to polyhaline sites, which were separated into two sediment groups above and below 90% sand content. Assemblages corresponding to each of these 9 habitats were identified in the context of widely recognized patterns of dominant taxa. Differences between North Carolina and Delaware-Chesapeake Bay polyhaline assemblages were attributed to the relative contributions of species and not to differences in species composition. No zoogeographic discontinuities could be identified. Our results reinforce the findings of recent studies which suggest that, with respect to estuarine benthic assemblages, the boundary between the Virginian and the Carolinian Provinces be moved to a new location south of Pamlico Sound.

Introduction

Over the past decade there has been an effort to characterize environmental conditions and gradients of coastal regions. Many assessments of environmental condition have focused on benthic assemblages because the benthos respond predictably to natural and anthropogenic stress (Pearson and Rosenberg 1978; Dauer 1993; Wilson and Jeffrey 1994). Several efforts have been aimed at developing indices of benthic condition for estuaries and coastal regions (e.g., Weisberg et al. 1997; Engle and Summers 1999a; Van Dolah et al. 1999; Paul et al. 2001; Smith et al. 2001). Although some of these indices are applicable to large coastal regions, the emphasis has been on the characteriza-

tion of single estuaries and local conditions, especially in the mid-Atlantic region of the United States (Weisberg et al. 1997; Adams et al. 1998). Regional assessments, however, are critical for effective management of coastal ecosystems, and this requires a comprehensive strategy that integrates key, region-wide, environmental problems. One such regional assessment is the Mid-Atlantic Integrated Assessment (MAIA) Program of the U.S. Environmental Protection Agency.

MAIA is a cooperative effort of federal and state agencies for the evaluation of ecosystem health. MAIA was established in 1995 to assess the condition of the natural resources of the mid-Atlantic region of the U.S. (Bradley and Landy 2000). The MAIA region encompasses the watersheds of the Delaware Bay, Chesapeake Bay, the coastal bays of the Delmarva peninsula, and the Albemarle-Pamlico estuarine system (Fig. 1). One of the objectives

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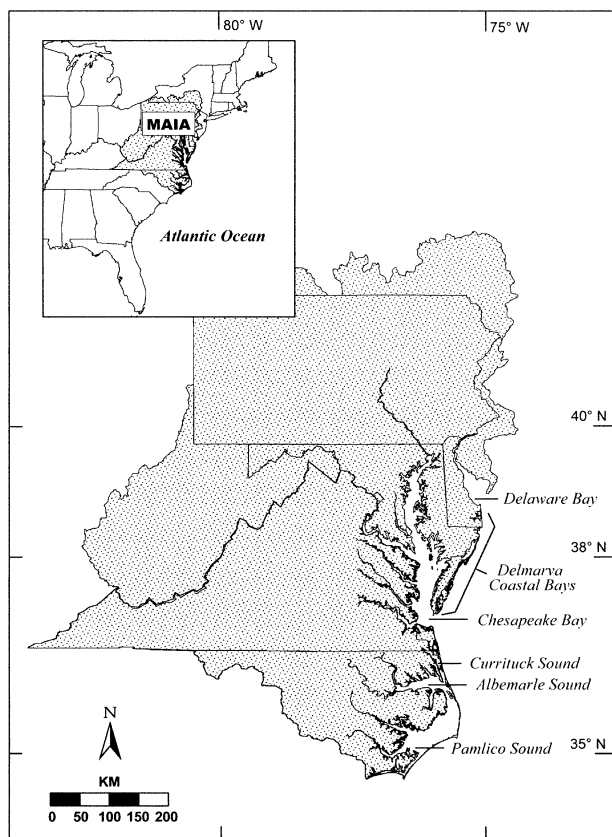


Fig. 1. Mid-Atlantic Integrated Assessment (MAIA) region.

of MAIA is to characterize the condition of mid-Atlantic estuaries using a common set of measurements applied over the entire region. To achieve this objective, we developed an index of benthic community condition for application throughout the MAIA region. The index is described in a separate manuscript (Llansó et al. 2002). We chose an index based on a scoring system approach similar to that of the benthic index of biotic integrity (B-IBI) developed for Chesapeake Bay (Weisberg et al. 1997) because of its simplicity, ease of interpretation, and previous successful application for assessing benthic community condition in Chesapeake Bay (Dauer and Llansó in press; Llansó et al. in press) and throughout estuaries of the southeastern U.S. (Van Dolah et al. 1999).

The development of an index of benthic community condition for the MAIA region requires the identification of reference conditions in key community attributes to which attributes from putatively degraded sites can be compared. In addition, habitat factors that control unimpaired assemblage structure need to be identified to ensure that natural differences in benthic assemblages related to habitat type do not confound index interpretation.

Other index development efforts applicable to portions of the MAIA region have identified sediment type, salinity, and latitude as factors structuring estuarine benthic assemblages, and have partitioned the data into habitat classes defined by these factors (e.g., Weisberg et al. 1997; Van Dolah et al. 1999). Although numerous studies have focused on the identification of factors that control benthic infaunal assemblages in estuaries of the mid-Atlantic region (e.g., Mountford et al. 1977; Maurer et al. 1978; Chester et al. 1983; Tourtellotte and Dauer 1983; Holland et al. 1987), most are system-specific and have not described habitat types and associated assemblages on a region-wide basis.

The objective of the present study is to identify major habitat types in the MAIA region based on benthic infaunal classification analysis and the evaluation of the physical characteristics that structure these assemblages. The resulting habitat classification will be the basis for the partition of data and the formulation of an index of benthic community condition that accounts for variability due to habitat type. We also describe the benthic assemblages and discuss these results in terms of differences between zoogeographic provinces.

Methods

Data from seven sampling programs were assembled for this study (Table 1). One sampling program was specifically established in 1997 for the MAIA project, while the remaining were part of long-term monitoring programs that have sampled estuaries in the region from as early as 1984. Only sites that included benthic information and were sampled during the summer, defined as July 1 through October 7, were used in this study. Summer is the time of the year when benthic assemblages and associated ecological processes show the greatest response to both natural and anthropogenic stress. For this reason, our analysis used only data collected during the summer season. The resulting database contained 1,999 sites, 19 of which were sampled in multiple years yielding a total of 2,083 sampling events. Data for sites sampled more than once during the summer were averaged. All sampling programs used a 440-cm² Young grab (one to three grabs per site), sieved samples through a 0.5-mm mesh screen, and identified organisms to the lowest possible taxonomic level. Details on specific sampling protocols and measurements taken in each of the programs can be found in the references provided in Table 1.

Data sets used to describe community structure were first standardized to ensure common species nomenclature. Incidental and epifaunal organisms were eliminated from the data, and taxa such as

TABLE 1. Data sources used in this study.

Program	Time Period	Sampling Location	References
Mid-Atlantic Integrated Assessment	1997–1998	Chesapeake and Delaware estuaries	Strobel 1998
Environmental Monitoring and Assessment, Carolinian Province	1997–1998	Chowan and Neuse Rivers	Hyland et al. 1998, 2000
Environmental Monitoring and Assessment, Carolinian Province	1994–1997	Pamlico-Albemarle Sounds and tributaries	Hyland et al. 1998, 2000
National Oceanic and Atmospheric Administration, National Status and Trends	1997–1998	Delaware estuary, Chesapeake Bay	Hartwell et al. 2001
Maryland Long-Term Benthic Monitoring	1984–1998	Maryland portion of Chesapeake Bay	Ranasinghe et al. 1999
Virginia Benthic Monitoring	1985–1998	Virginia portion of Chesapeake Bay	Dauer et al. 1998
Environmental Monitoring and Assessment, Virginian Province	1990–1993	Virginian Province	Paul et al. 1999
Coastal Bays Joint Assessment	1993	Delaware and Maryland coastal bays	Chaillou et al. 1996

the Anthozoa and Nemertina for which there was variable taxonomic effort between collection programs, were combined at the phylum or class levels (a complete list is available from the author). We excluded epifauna from all the analyses of the index development effort, including those leading to the characterization of assemblages and habitat factors in this study, because these organisms are not fully exposed to chemical contaminants in sediments and were not expected to be sensitive indicators of sediment quality.

Reference (non-degraded) sites were then selected according to dissolved oxygen, sediment contaminant, and sediment toxicity criteria. These criteria were similar to those used to define reference sites in previous studies (Weisberg et al. 1997; Van Dolah et al. 1999). We excluded degraded sites from the present study so that natural gradients in community composition could be identified without the confounding influence of anthropogenic effects. Only sites for which all sampling events had associated dissolved oxygen and chemistry data were considered. Sites were defined as reference if, for all sampling events, dissolved oxygen concentrations were greater than 3.0 ppm, no chemical contaminant concentration exceeded Long et al.'s (1995) effects range-median concentrations, no more than two chemical contaminants exceeded Long et al.'s (1995) effects range-low concentrations, and sediments were not toxic in *Ampelisca abdita* or Microtox bioassays. Amphipod bioassays followed guidelines provided in American Society for Testing Materials (1993) protocols, and were considered to indicate toxicity when mean test survival was significantly different from and less than 80% of control survival. Microtox bioassays were considered to indicate toxicity when the EC_{50} of test sediments (sediment concentration that reduces bacterial light production by 50% relative to water controls) was ≤ 0.2 for sediments with silt-clay content $\geq 20\%$, the EC_{50} was ≤ 0.5 for sediments with silt-clay content $< 20\%$, or the EC_{50} of test

sediments was significantly different from controls (Ringwood et al. 1997). A dissolved oxygen criterion of 3.0 ppm for point-in-time measurements is consistent with other benthic index approaches and with the general literature which suggests that concentrations below 3.0 ppm adversely affect benthos (Diaz and Rosenberg 1995).

Benthic infaunal assemblages and habitat groups were identified using classification (cluster) analysis (Boesch 1977a). Four hundred sixty reference sites were used in the analysis to identify affinities between sites based on mean species abundance. Also, inverse (by species) classifications were produced to identify the species responsible for site segregation. All analyses were conducted using the computer program COMPAH (Eugene Gallagher, Environmental Science Program, University of Massachusetts, Boston, Massachusetts). The Bray-Curtis similarity coefficient and the flexible sorting method (with β coefficient = -0.25 , see Boesch 1977a) were used. Abundance data were standardized to values per square meter and transformed to $\log(x+1)$ to lessen the sensitivity of the similarity measure to large abundances. In addition, nodal analysis (Williams and Lambert 1961; Boesch 1977a) was used to relate the groups derived from normal and inverse classifications. Dominant species were identified using the Biological Index Value (McCloskey 1970). The index is a reflection of the frequency with which a species is present in the assemblage, and it was computed by assigning 10 points to the first numerically dominant species at each site, nine points to the second, etc. A score of 0 was assigned to species ranking below the tenth position. Tied species received the average of their individual scores. Ranking scores for the 10 most abundant species were then summed over all sites for each of the groups derived from the classification of sites.

Prior to cluster analysis, taxonomic designations with more than one level of identification were treated by removing the higher level (e.g., genus)

and keeping the lower level (e.g., species) of identification. Taxa occurring in no more than 10 sampling events across data sets were also eliminated from the analysis. This last procedure eliminated 59% of the taxa while keeping 96% of the cumulative total abundance in the analysis. The elimination of species occurring in low frequencies usually helps clarify the distribution patterns of the more common species (Boesch 1977a). Habitat classes were delineated by examining salinity, substrate, depth, and station location information of sites grouped in cluster analysis, and by multiple discriminant analysis to correlate site group separation with environmental variables (Green and Vascotto 1978; Hyland et al. 1991). Variables used in discriminant analysis were salinity, percent silt-clay (transformed to arcsine \sqrt{p} to approximate normality, where p is a proportion), percent total organic carbon (TOC), depth, and latitude. Total structure coefficients, which are the correlations between the original variables and the discriminant functions, served as a measure of the relative contribution of each variable to group separation.

Results

HABITAT CHARACTERIZATION

Site classifications showed patterns of spatial variation that could be explained primarily on the basis of differences in salinity and secondarily on the basis of differences in sediment composition (Fig. 2). Some sites were separated by geographical location. Water depth and sampling year did not appear to be factors in the grouping of sites.

Thirteen major groups of sites defined by the benthic assemblages could be distinguished in the dendrogram. Ten of these groups corresponded closely to one of six salinity classes defined according to a modified Venice System: tidal freshwater (0.0–0.5‰), oligohaline (0.5–5‰), low mesohaline (5–12‰), high mesohaline (12–18‰), polyhaline (18–27‰), and euhaline (> 27‰). Two additional groups indicated conditions that were transitional between adjacent salinity classes (Fig. 3), and one group consisted of outliers of mixed salinity. Although the ten salinity groups were distinct in the dendrogram, some sites with salinity measurements corresponding to different salinity classes were grouped together. Adding the rare species (see methods) or decreasing the number of taxa available for classification did not result in improved site segregation.

A relationship between sediment characteristics and site segregation was discernible within the polyhaline and euhaline salinity groups (Fig. 4). Two groups of sites within each of these salinity classes were distinct in the dendrogram at broad levels of

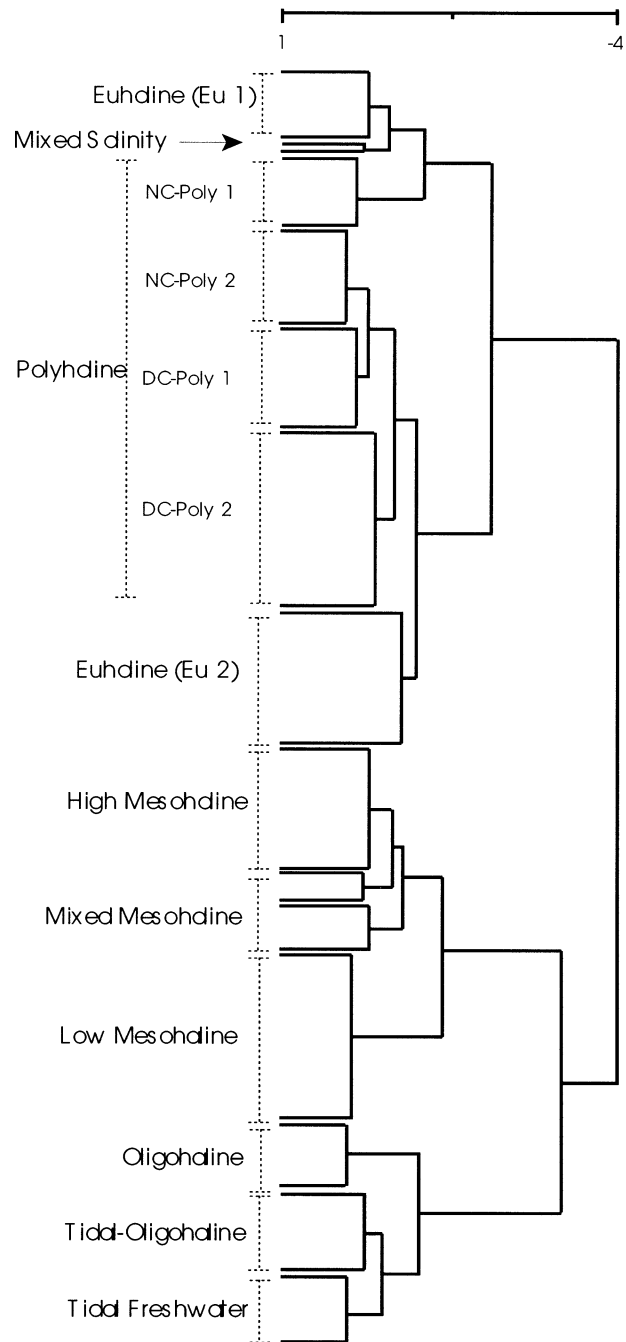


Fig. 2. Cluster analysis of 460 reference (non-degraded) sites in the Mid-Atlantic Integrated Assessment (MAIA) region. Sites were separated according to salinity. Polyhaline and euhaline sites were further separated according to sediment type and geographical location. Individual sites and small groupings within major cluster groups have been omitted for clarity. Eu 1 = high-energy euhaline; Eu 2 = Delaware-Chesapeake Bay euhaline; NC-Poly 1 = North Carolina polyhaline silt and clay; NC-Poly 2 = North Carolina polyhaline sand; DC-Poly 1 = Delaware-Chesapeake Bay polyhaline sand; DC-Poly 2 = Delaware-Chesapeake Bay polyhaline silt and clay.

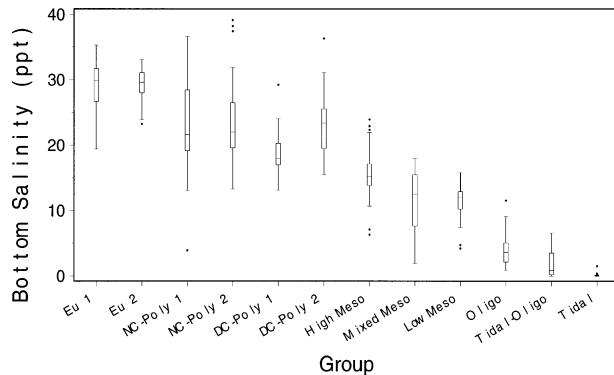


Fig. 3. Salinity ranges for 12 groups of sites identified by cluster analysis. Boxes are interquartile ranges (25 and 75 percentiles); range bars (1.5 interquartile range) show maximum and minimum of non-outliers; small horizontal lines within boxes are medians; dots are outliers. Eu 1 = high-energy euhaline; Eu 2 = Delaware-Chesapeake Bay euhaline; NC-Poly 1 = North Carolina polyhaline silt and clay; NC-Poly 2 = North Carolina polyhaline sand; DC-Poly 1 = Delaware-Chesapeake Bay polyhaline sand; DC-Poly 2 = Delaware-Chesapeake Bay polyhaline silt and clay; High Meso = high mesohaline; Mixed Meso = transitional group between high and low mesohaline; Low Meso = low mesohaline; Oligo = oligohaline; Tidal-Oligo = transitional group between tidal freshwater and oligohaline; Tidal = tidal freshwater.

mud and sand. Polyhaline sites with high sand content were generally separated from polyhaline sites with varying amounts of silt and clay above 10%. A group of euhaline sites with high sand content was distinct in the dendrogram. The effect of sed-

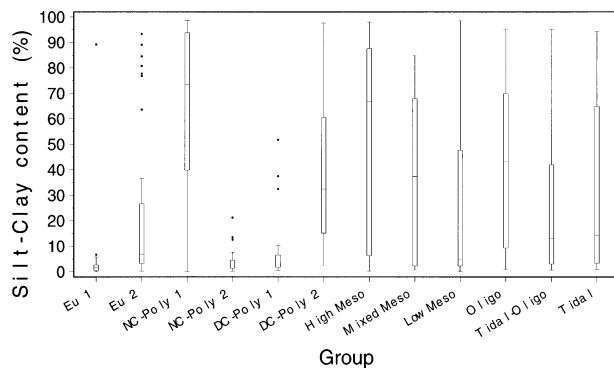


Fig. 4. Silt-clay ranges for 12 groups of sites identified by cluster analysis. Boxes are interquartile ranges (25 and 75 percentiles); range bars (1.5 interquartile range) show maximum and minimum of non-outliers; small horizontal lines within boxes are medians; dots are outliers. Eu 1 = high-energy euhaline; Eu 2 = Delaware-Chesapeake Bay euhaline; NC-Poly 1 = North Carolina polyhaline silt and clay; NC-Poly 2 = North Carolina polyhaline sand; DC-Poly 1 = Delaware-Chesapeake Bay polyhaline sand; DC-Poly 2 = Delaware-Chesapeake Bay polyhaline silt and clay; High Meso = high mesohaline; Mixed Meso = transitional group between high and low mesohaline; Low Meso = low mesohaline; Oligo = oligohaline; Tidal-Oligo = transitional group between tidal freshwater and oligohaline; Tidal = tidal freshwater.

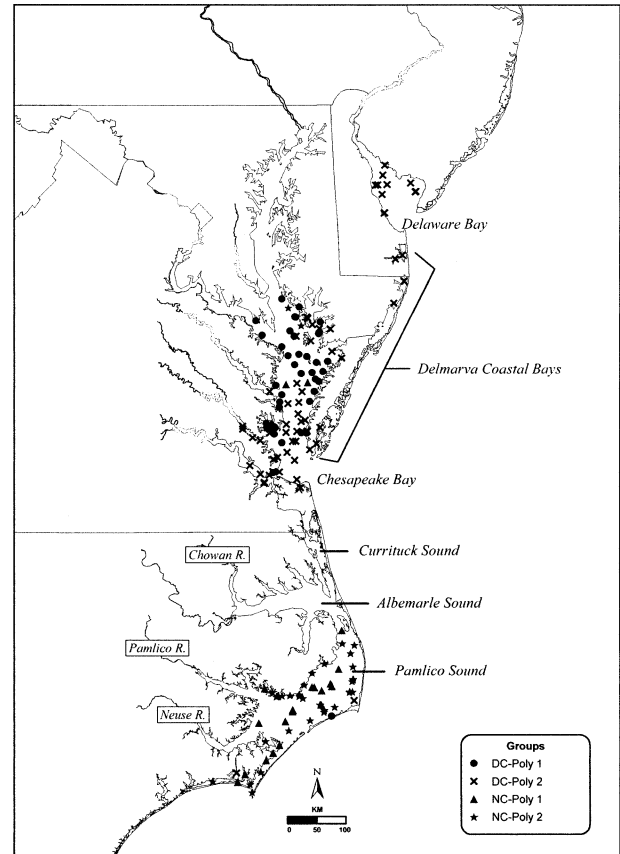


Fig. 5. Geographical distribution of 4 of 12 habitat groups identified by cluster analysis. DC-Poly 1 = Delaware-Chesapeake Bay polyhaline sand; DC-Poly 2 = Delaware-Chesapeake Bay polyhaline silt and clay; NC-Poly 1 = North Carolina polyhaline silt and clay; NC-Poly 2 = North Carolina polyhaline sand.

iment type was not entirely clear as there was overlap between sand and silt-clay groups. In particular, the separation between euhaline site groups appeared to be related to the energy of the habitat rather than to percent silt and clay. The euhaline site group with high sand content consisted of shallow coastal sites or sites near barrier-island inlets exposed to wave action. The overriding factor structuring assemblages at these locations is likely to be sediment mobility. Notwithstanding the overlap in sediment types, a distinction between sand and silt-clay groups based on silt-clay interquartile (25–75%) range distributions could be made for the polyhaline sites (Fig. 4).

In addition to salinity and sediment groups, polyhaline and euhaline sites in the Pamlico-Albemarle Sound region were generally separated from all others (Fig. 5). No separation based on geographical location was discernible for the lower salinity groups. High salinity groups in the Pamlico-Albemarle Sound region exhibited a substantial

TABLE 2. Total structure coefficients of abiotic environmental variables on the first 3 discriminant functions (DF), which accounted for 98% of the cumulative variance. These are the correlations between the discriminating variables and the discriminant functions. Correlations considered important on each function are underlined. TOC = total organic carbon.

Variable	DF I	DF II	DF III
Salinity (‰)	<u>0.994</u>	0.070	-0.007
Silt-Clay (arcsine \sqrt{p})	-0.156	<u>0.936</u>	0.259
Depth (m)	0.194	0.327	<u>0.072</u>
Latitude (decimal degrees)	-0.069	-0.276	<u>0.882</u>
TOC (%)	-0.245	<u>0.776</u>	-0.043

amount of overlap, with many euhaline and polyhaline sites clustering together.

Multiple discriminant analysis of physical parameters indicated highly significant separation of groups ($p < 0.0001$) as defined by the numerical classification. Three discriminant functions accounted for 98% of the cumulative variance. The first function accounted for 83% of the among-group variance and was related to salinity (Table 2). The second discriminant function accounted for 12% of the among-group variance and revealed the influence of silt and clay in the high salinity habitats. The second function was also related to TOC content of the sediment, though the influence of TOC is probably a reflection of the strong correlation between TOC and percent silt-clay. The third discriminant function accounted for 3% of the variance and was related to latitude. The variance accounted for by additional functions was considered negligible and added little to the interpretation.

Based on the above results, sites were classified into six habitat classes according to salinity: tidal freshwater, oligohaline, low mesohaline, high mesohaline, polyhaline, and euhaline. The polyhaline habitat can further be divided into sand and silt-clay strata segregated according to latitude into two groups: North Carolina estuaries and Delaware-Chesapeake Bay (inclusive of coastal bays). A distinction could not be made between polyhaline and euhaline habitats in the North Carolina estuaries. Nine habitat classes based on macroinfaunal species composition and abundance were defined for the MAIA region (Table 3). The high-energy euhaline habitat was not included in this classification because sediment properties (e.g., median grain size, sorting coefficient) that may aid in the classification of sites into this habitat were not measured by the sampling programs used in this study. We include this habitat in the characterization of species assemblages below.

SPECIES ASSEMBLAGES

Using 143 taxa, twenty-three species groups were distinguished in the inverse classification analysis.

TABLE 3. Habitats defined in the Mid-Atlantic Integrated Assessment (MAIA) region. na = not applicable.

Habitat	Bottom Salinity (‰)	Silt-Clay (< 62 μ) Content by Weight (%)
Tidal freshwater	0–0.5	na
Oligohaline	≥ 0.5 –5	na
Low mesohaline	≥ 5 –12	na
High mesohaline	≥ 12 –18	na
Delaware-Chesapeake Bay polyhaline sand	≥ 18 –27	0–10
Delaware-Chesapeake Bay polyhaline silt and clay	≥ 18 –27	>10
North Carolina polyhaline sand	≥ 18	0–10
North Carolina polyhaline silt and clay	≥ 18	>10
Delaware-Chesapeake Bay euhaline	≥ 27	na

This distinction was based on the separation of species groups from other groups in the analysis, rather than on a predetermined level of similarity. The dendrogram is not shown here but a list of species by group is presented in Table 4. Nodal analysis revealed most species groups as having clear affinities with some of the site groups, providing an explanation for the observed site classification. Some species, particularly those in group 17 (Table 4), had no particular affinity with any one site group but presented a moderate to high frequency of occurrence among sites in the mesohaline to euhaline salinity range. Species exhibiting a moderate to high frequency of occurrence across sites were euryhaline species of broad distribution in the higher salinity habitats and in most sediment types. These species were often numerically dominant.

Species groups 1, 4, and 6 (Table 4) characterized high-energy euhaline sites in terms of frequency of occurrence. These sites exhibited a lower number of species than euhaline sites elsewhere. Numerically dominant species at these sites were the bivalves *Tellina agilis* and *Spisula solidissima*, nemertean, the polychaete *Leitoscoloplos* sp., and the amphipods *Parahastorius longimerus*, *Rhepoxynius hudsoni*, and *Acanthohastorius millsi* (Table 5). Species groups 9 and 17 characterized low-energy euhaline sites from Delaware to Chesapeake Bay. Species in groups 5, 7, 15, and 16 had high affinity for these sites, occurring in low frequency but seldom occurring elsewhere. Numerically dominant species of the Delaware-Chesapeake Bay euhaline site group were the polychaete *Mediomastus ambiseta* and the bivalve *T. agilis*.

North Carolina polyhaline sites (≥ 18 ‰) were characterized by species groups 3 and 17. Species in group 3 occurred in low frequency, but were mostly restricted to these sites. Silt and clays differed from sands in having lower number of species and in dominance patterns. Numerically-dom-

TABLE 4. Species groups identified by inverse classification analysis of 460 sites used in this study.

Group 1	Group 11	Phoronis spp.
<i>Acanthohaustorius millsii</i>	<i>Drilonereis longa</i>	<i>Heteromastus filiformis</i>
<i>Paraonis fulgens</i>	<i>Nassarius vibex</i>	<i>Streblospio benedicti</i>
<i>Parahaustorius longimerus</i>	<i>Glycera dibranchiata</i>	<i>Neanthes succinea</i>
<i>Spio pettiboneae</i>	<i>Lyonsia hyalina</i>	Nemertina
Group 2	<i>Scoloplos rubra</i>	Oligochaeta
<i>Ogyrides alphaerostris</i>	<i>Gemma gemma</i>	<i>Mulinia lateralis</i>
<i>Parahesione luteola</i>	<i>Haminoea solitaria</i>	Group 18
Group 3	<i>Leptosynapta tenuis</i>	<i>Ampelisca abdita-vadourum</i> complex
<i>Aglaophamus verrilli</i>	<i>Scolecopsis texana</i>	<i>Leucon americanus</i>
<i>Apopriospio pygmaea</i>	Group 12	<i>Pectinaria gouldii</i>
Sipuncula	<i>Aligena elevata</i>	<i>Rictaxis punctostriatus</i>
<i>Magelona</i> spp.	<i>Chaetopterus variopedatus</i>	<i>Leitoscoloplos robustus</i>
<i>Pseudeurythoe ambigua</i>	<i>Bhawania heteroseta</i>	<i>Leitoscoloplos</i> sp.
Group 4	<i>Prionospio perkinsi</i>	Anthozoa
<i>Americhelidium americanum</i>	<i>Nephtys cryptomma</i>	<i>Listriella barnardi</i>
<i>Nephtys picta</i>	<i>Scolecopsis bousfieldi</i>	<i>Spiochaetopterus costarum</i>
<i>Rhepoxynius hudsoni</i>	Group 13	<i>Polydora cornuta</i>
Group 5	<i>Clymenella torquata</i>	Group 19
<i>Brania wellfleetensis</i>	<i>Listriella clymenellae</i>	<i>Ameroculodes</i> species complex
<i>Caulieriella</i> sp. B (Blake)	<i>Macoma tenta</i>	<i>Lepidactylus dytiscus</i>
<i>Neanthes arenaceodentata</i>	<i>Saccoglossus kowalevskii</i>	<i>Laeonereis culveri</i>
<i>Cirrophorus</i> spp.	<i>Sigambra tentaculata</i>	<i>Cyclaspis varians</i>
Group 6	Group 14	<i>Eteone foliosa</i>
<i>Ancinus depressus</i>	<i>Cabira incerta</i>	Group 20
<i>Spisula solidissima</i>	<i>Monticellina baptistae-dorsobranchialis</i>	<i>Chiridotea almyra</i>
Group 7	complex	<i>Rangia cuneata</i>
<i>Amastigos caperatus</i>	<i>Microphiopholis atra</i>	<i>Tubificoides</i> spp.
<i>Spio setosa</i>	Group 15	Group 21
<i>Aricidea catherinae</i>	<i>Anadara transversa</i>	<i>Cyathura polita</i>
<i>Euceramus praelongus</i>	<i>Podarke obscura</i>	<i>Marenzelleria viridis</i>
<i>Polygordius</i> spp.	<i>Capitella capitata</i> complex	<i>Macoma mitchelli</i>
Group 8	<i>Mysella planulata</i>	<i>Leptocheirus plumulosus</i>
<i>Asabellides oculata</i>	<i>Parapionosyllis longicirrata</i>	<i>Eteone heteropoda</i>
<i>Nephtys incisa</i>	<i>Prionospio heterobranchia</i>	<i>Macoma balthica</i>
<i>Nucula proxima</i>	<i>Dipolydora socialis</i>	Group 22
<i>Upogebia affinis</i>	<i>Polycirrus</i> spp.	<i>Hargeria rapax</i>
Group 9	<i>Unicicola serrata</i>	<i>Hobsonia florida</i>
<i>Ampelisca verrilli</i>	Group 16	Group 23
<i>Tellina agilis</i>	<i>Arabella iricolor-multidentata</i> complex	<i>Aulodrilus pigueti</i>
<i>Tharyx</i> sp. A Morris	<i>Demonax microphthalmus</i>	<i>Tanytarsus</i> spp.
<i>Oxyurostylis smithi</i>	<i>Tagelus divisus</i>	<i>Dicrotendipes</i> spp.
<i>Ensis directus</i>	<i>Ceratonereis irritabilis</i>	Chironomidae
<i>Owenia fusiformis</i>	<i>Carazziella hobsonae</i>	<i>Corbicula fluminea</i>
<i>Phyllodoce arenae</i>	<i>Cyathura burbancki</i>	<i>Polypedium</i> spp.
<i>Mercenaria mercenaria</i>	<i>Diopatra cuprea</i>	<i>Chironomus</i> spp.
Group 10	<i>Scoletoma tenuis</i>	<i>Coelotanytus</i> spp.
<i>Ancistrosyllis hartmanae</i>	<i>Exogone dispar</i>	<i>Limnodrilus hoffmeisteri</i>
<i>Branchiostoma caribaeum</i>	<i>Glycera americana</i>	Tubificidae immature without
<i>Loimia medusa</i>	<i>Macroclymene zonalis</i>	capilliform chaetae
<i>Spiophanes bombyx</i>	<i>Melinna maculata</i>	<i>Cryptochironomus</i> spp.
<i>Parvilucina multilineata</i>	<i>Notomastus</i> sp. A Ewing	<i>Procladius</i> spp.
<i>Podarkeopsis levifuscina</i>	<i>Sabaco elongatus</i>	Tubificidae immature with capilliform
<i>Pseudeurythoe paucibranchiata</i>	Group 17	chaetae
	<i>Acteocina canaliculata</i>	Ceratopogonidae
	<i>Mediomastus ambiseta</i>	<i>Isochaetides freyi</i>
	<i>Glycinde solitaria</i>	<i>Cladotanytarsus</i> spp.
	<i>Paraprionospio pinnata</i>	

TABLE 5. Mean abundance m^{-2} (\pm SD) and Biological Index Value (BIV) of numerically-dominant macroinfauna in each of nine habitat classes, plus the high-energy euhaline, defined by classification of sites in the Mid-Atlantic Integrated Assessment (MAIA) region. Only sites where the species was present (n) were used in the calculation of means. N = total number of sites in a habitat class. BIVs were obtained by assigning rank scores to numerically dominant species for each site, and summing scores for the 10 most abundant species over all sites in each habitat class. Species are listed by BIV, from highest to lowest. Eu 1 = high-energy euhaline; Eu 2 = Delaware-Chesapeake Bay euhaline; NC-Poly 1 = North Carolina polyhaline silt and clay; NC-Poly 2 = North Carolina polyhaline sand; DC-Poly 1 = Delaware-Chesapeake Bay polyhaline sand; DC-Poly 2 = Delaware-Chesapeake Bay polyhaline silt and clay; High Meso = high mesohaline; Low Meso = low mesohaline; Oligo = oligohaline; Tidal = tidal freshwater.

Habitat	Species	Mean Abundance (SD)	BIV	n
Eu 1 (N = 26)	<i>Tellina agilis</i>	129.7 (216.1)	118.5	17
	Nemertina	175.0 (513.3)	103.5	16
	<i>Leitoscoloplos</i> sp.	117.2 (140.2)	97.5	14
	<i>Parahaustorius longimerus</i>	365.3 (457.8)	82.0	9
	<i>Rhepoxynius hudsoni</i>	167.0 (213.2)	66.0	10
	<i>Spisula solidissima</i>	193.7 (280.4)	56.0	7
	<i>Acanthohaustorius millsi</i>	1904.7 (1580.0)	50.8	6
	<i>Ampelisca abdita-vadourum</i> complex	26.5 (42.8)	41.0	6
	<i>Nephtys picta</i>	36.7 (38.6)	40.8	10
	<i>Paraonis fulgens</i>	151.5 (139.6)	39.0	5
Eu 2 (N = 50)	<i>Mediomastus ambiseta</i>	1011.1 (1352.6)	347.5	48
	<i>Tellina agilis</i>	1059.1 (2623.7)	221.9	41
	Oligochaeta	273.8 (365.2)	199.3	47
	<i>Tharyx</i> sp. A Morris	124.1 (138.3)	107.0	36
	<i>Ampelisca verrilli</i>	170.6 (272.2)	101.0	38
	<i>Mulinia lateralis</i>	180.6 (257.4)	99.0	37
	<i>Exogone dispar</i>	378.2 (615.8)	97.5	24
	<i>Acteocina canaliculata</i>	127.3 (129.4)	91.0	39
	<i>Ampelisca abdita-vadourum</i> complex	325.2 (713.9)	87.5	33
	<i>Streblospio benedicti</i>	132.0 (175.4)	87.0	31
NC-Poly 1 (N = 25)	Nemertina	94.5 (101.9)	180.6	22
	<i>Paraprionospio pinnata</i>	83.8 (85.2)	148.0	20
	<i>Acteocina canaliculata</i>	159.8 (146.8)	132.0	16
	<i>Sigambra tentaculata</i>	52.8 (47.2)	101.2	17
	<i>Phoronis</i> spp.	44.7 (49.5)	71.8	15
	<i>Mediomastus ambiseta</i>	108.6 (223.4)	67.1	12
	<i>Magelona</i> spp.	39.0 (33.1)	49.5	10
	<i>Pseudeurythoe ambigua</i>	62.9 (52.7)	44.0	9
	<i>Streblospio benedicti</i>	65.9 (67.4)	35.0	5
	<i>Macoma tenta</i>	33.5 (37.5)	34.1	7
NC-Poly 2 (N = 36)	<i>Acteocina canaliculata</i>	306.4 (316.1)	241.5	36
	<i>Parvilucina multilineata</i>	332.2 (301.2)	229.4	33
	<i>Mediomastus ambiseta</i>	378.8 (565.1)	192.5	27
	<i>Phoronis</i> spp.	364.5 (1018.7)	180.5	30
	Nemertina	76.0 (59.8)	132.4	35
	<i>Paraprionospio pinnata</i>	69.9 (54.6)	102.7	27
	<i>Tellina agilis</i>	80.6 (75.8)	80.3	22
	<i>Mulinia lateralis</i>	93.3 (143.0)	79.0	24
	<i>Glycinde solitaria</i>	53.3 (51.4)	64.4	26
	<i>Ampelisca verrilli</i>	81.6 (85.5)	46.5	17
DC-Poly 1 (N = 38)	<i>Acteocina canaliculata</i>	468.1 (543.8)	249.0	37
	<i>Phoronis</i> spp.	405.2 (544.2)	230.6	36
	<i>Glycinde solitaria</i>	190.1 (134.2)	188.1	38
	<i>Mediomastus ambiseta</i>	622.6 (1461.5)	153.5	27
	Oligochaeta	323.5 (443.5)	135.0	30
	<i>Rictaxis punctostriatus</i>	124.0 (116.3)	97.3	29
	<i>Heteromastus filiformis</i>	111.9 (198.5)	80.0	31
	<i>Paraprionospio pinnata</i>	112.6 (150.9)	60.1	26
	<i>Branchiostoma caribaeum</i>	302.0 (605.6)	55.5	15
	<i>Gemma gemma</i>	4402.6 (16050.5)	53.0	14
DC-Poly 2 (N = 66)	<i>Mediomastus ambiseta</i>	1553.3 (2613.2)	532.7	64
	<i>Paraprionospio pinnata</i>	303.0 (306.2)	339.5	63
	<i>Acteocina canaliculata</i>	353.1 (443.2)	301.5	58
	<i>Glycinde solitaria</i>	165.9 (134.3)	272.9	65
	Oligochaeta	250.1 (490.1)	146.4	42
	<i>Leucon americanus</i>	132.3 (254.6)	130.9	47

TABLE 5. Continued.

Habitat	Species	Mean Abundance (SD)	BIV	n
High Meso (N = 46)	<i>Streblospio benedicti</i>	340.8 (677.8)	118.0	32
	<i>Neanthes succinea</i>	141.6 (347.9)	99.8	44
	<i>Ampelisca abdita-vadorum</i> complex	113.9 (141.6)	99.3	41
	Nemertina	89.0 (92.1)	96.1	58
	<i>Streblospio benedicti</i>	330.2 (412.6)	297.0	42
	<i>Mulinia lateralis</i>	220.3 (407.5)	214.5	31
	<i>Heteromastus filiformis</i>	134.8 (384.5)	166.5	33
	<i>Mediomastus ambiseta</i>	565.1 (1110.2)	156.8	25
	<i>Glycinde solitaria</i>	71.7 (83.8)	143.3	33
	<i>Macoma mitchelli</i>	166.0 (220.4)	137.0	22
	<i>Paraprionospio pinnata</i>	168.6 (222.1)	128.3	22
	Oligochaeta	275.6 (703.2)	120.8	21
	Nemertina	47.6 (54.5)	113.5	32
	<i>Neanthes succinea</i>	77.1 (81.6)	97.0	20
Low Meso (N = 60)	<i>Macoma michelli</i>	656.8 (568.0)	443.0	58
	<i>Heteromastus filiformis</i>	544.4 (587.4)	428.2	60
	<i>Streblospio benedicti</i>	230.7 (284.0)	282.2	56
	<i>Marenzelleria viridis</i>	281.3 (320.8)	275.0	56
	<i>Cyathura polita</i>	138.6 (116.9)	232.5	55
	<i>Neanthes succinea</i>	215.0 (273.7)	224.8	54
	Oligochaeta	832.0 (1623.7)	212.0	45
	<i>Leptocheirus plumulosus</i>	418.0 (493.7)	205.5	34
	Nemertina	83.4 (83.9)	150.5	58
	<i>Laeonereis culveri</i>	165.2 (170.5)	123.5	28
	<i>Tubificoides</i> spp.	444.7 (600.4)	176.5	23
Oligo (N = 23)	<i>Marenzelleria viridis</i>	301.6 (476.8)	145.0	19
	<i>Cyathura polita</i>	194.7 (291.8)	129.5	20
	<i>Rangia cuneata</i>	108.5 (137.0)	109.0	17
	<i>Leptocheirus plumulosus</i>	698.1 (837.3)	92.0	13
	<i>Coelotanypus</i> spp.	294.7 (327.9)	66.5	10
	<i>Limnodrilus hoffmeisteri</i>	556.2 (585.3)	47.5	6
	Tubificidae imm. without capilliform chaetae	62.7 (99.4)	43.5	9
	<i>Hobsonia florida</i>	176.4 (215.3)	42.0	7
	Nemertina	38.7 (49.6)	42.0	9
	Tubificidae imm. without capilliform chaetae	1710.1 (2047.3)	211.0	23
	<i>Limnodrilus hoffmeisteri</i>	312.0 (349.6)	165.0	24
Tidal (N = 25)	<i>Corbicula fluminea</i>	415.8 (412.3)	136.8	18
	<i>Coelotanypus</i> spp.	127.5 (122.3)	118.5	21
	<i>Polypedium</i> spp.	268.7 (399.5)	99.5	15
	Tubificidae imm. with capilliform chaetae	195.5 (260.2)	93.0	15
	<i>Cyathura polita</i>	110.1 (102.2)	88.0	15
	<i>Procladius</i> spp.	64.8 (86.4)	74.0	20
	<i>Chironomus</i> spp.	456.4 (1332.9)	66.8	14
	<i>Cryptochironomus</i> spp.	38.7 (34.5)	62.8	18

inant species of North Carolina polyhaline silt and clay sites were nemerteans, the polychaete *Paraprionospio pinnata*, and the gastropod *Acteocina canaliculata* (Table 5). Dominant species of North Carolina polyhaline sand sites were the gastropod *A. canaliculata* and the bivalve *Parvilucina multilineata*.

Delaware-Chesapeake Bay polyhaline sites (18–27‰) were characterized by species groups 17 and 18, and by other species with a lower frequency of occurrence among these sites. Low frequency species groups 8 and 10–14 differentiated many of these sites from other site groups. Species groups 8 and 12–14 had a higher affinity for silt and clays, while species groups 10 and 11 had a higher affinity

for sands. Numerically dominant species of Delaware-Chesapeake Bay silt and clay sites were the polychaetes *M. ambiseta* and *P. pinnata*, and the gastropod *A. canaliculata* (Table 5). Sand sites were dominated by *A. canaliculata*, the phoronid *Phoronis* spp., and the polychaete *Glycinde solitaria*.

Species group 17 (Table 4) characterized both high and low mesohaline sites. In addition, low mesohaline sites included many species in groups 19 and 21. Numerically dominant species of high mesohaline sites were the polychaete *Streblospio benedicti* and the bivalve *Mulinia lateralis* (Table 5). Low mesohaline sites were dominated by the bivalve *Macoma mitchelli* and the polychaete *Heteromastus fil-*

iformis. Species groups 20 and 22 characterized oligohaline sites with dominance of *Tubificoides* spp. and the polychaete *Marenzelleria viridis* (Table 5). Species group 23 characterized the tidal freshwater with dominance of mature and immature (most tubificids without capilliform chaetae) *Limnodrilus hoffmeisteri*.

Discussion

The classification analysis of estuarine sites in the mid-Atlantic region of the U.S. revealed assemblages of organisms that were mainly associated with salinity gradients and sediment type, reinforcing results from previous studies. Weisberg et al. (1997), for example, distinguished five salinity habitats in the Chesapeake Bay that correspond closely to the salinity habitats described in this study and to the Venice System for the classification of estuarine waters. The salinity habitats defined in this study were the tidal freshwater, oligohaline, low mesohaline, high mesohaline, polyhaline, and euhaline. Weisberg et al.'s study (1997) did not distinguish euhaline sites within the Chesapeake Bay. In the present study we found that the separation of euhaline sites was closer to the 27‰ breakpoint than to the generally accepted 30‰. Weisberg et al. (1997) also classified assemblages according to sediment type at the cut-off level of 60% sand. Our classification analysis could only separate sites with a percentage of sand content above 90% in the polyhaline region. In our study dominant species in the mesohaline to euhaline salinity range were not restricted to one substrate, but were broadly distributed in different types of sediment with peaks of abundance in either sand or mud. There was also considerable overlap in the distribution of species across adjacent salinity types. These results suggest that infaunal assemblages in the estuaries of the mid-Atlantic region represent a continuum along the environmental gradient. This pattern has been widely reported for the Hudson River estuary (Ristich et al. 1977), Delaware Bay (Maurer et al. 1978), Chesapeake Bay (Boesch 1977b; Holland et al. 1987; Diaz and Schaffner 1990), South Carolina estuaries (Calder et al. 1977), and Georgia's Ogeechee River estuary (Dörjes and Howard 1975).

Our analysis of benthic community composition was restricted to macroinfaunal assemblages, to the summer index period, and to non-degraded sites in the mid-Atlantic region of the U.S. Using these types of data, our characterization of dominant species across salinity and sediment types in the MAIA region agrees well with characterizations from previous investigations. A consistent pattern of dominant taxa along the estuarine gradient of U.S. east coast temperate estuaries emerges from this and previous studies (Tenore 1972; Calder et

al. 1977; Ristich et al. 1977; Maurer et al. 1978; Holland et al. 1987). Tidal freshwater habitats are dominated by oligochaete and insect larvae, particularly limnodrilid oligochaetes (*Limnodrilus* spp.) and chironomid insect larvae, and the bivalve *Corbicula fluminea* can also be abundant. Oligohaline habitats are dominated by the polychaete *M. viridis*, the amphipod *Leptocheirus plumulosus*, and the isopod *Cyathura polita*, although each of these species can be common in low mesohaline regions. Additional taxa that can dominate oligohaline habitats are the polychaete *Hobsonia florida*, tubificid oligochaetes (particularly *Tubificoides* spp.), the bivalve *Rangia cuneata*, and the amphipod *Lepidactylus dytiscus*. Throughout the mesohaline and polyhaline regions euryhaline polychaetes can be dominants, including *Neanthes succinea*, *P. pinnata*, *S. benedicti*, *G. solitaria*, and *H. filiformis*. Additional taxa that can dominate in mesohaline habitats are the bivalves *M. mitchelli*, *Macoma balthica*, and *M. lateralis*, and the cumacean *Leucon americanus*. In polyhaline regions, in addition to the previously listed euryhaline polychaetes species, dominant species may include the polychaetes *Clymenella torquata* and *Glycera dibranchiata*. In both polyhaline and euhaline regions dominants include the capitellid polychaete *M. ambiseta*, the gastropod *A. canaliculata*, the bivalve *T. agilis*, ampeliscid amphipods, phoronids, and nemerteans. In these regions, the bivalve *Gemma gemma* is usually concentrated in high densities in a small number of sites. In polyhaline and euhaline regions with coarse sediments (usually indicative of a high energy habitat), additional dominants are phoxocephalid and haustoriid amphipods, the bivalve *S. solidissima*, and a variety of polychaetes such as nephyds, paraonids, and syllids.

In the above studies, as well as in the present study, both sediment type and salinity were major determinants in species distributional patterns. Sediment type was a minor factor compared to salinity. Some species were clearly restricted to groups of sites with sediments containing a high percentage of sand (Tourtellotte and Dauer 1983; this study), but the dominant species of other site groups were widespread across sediment types.

In addition to salinity and sediment type, our classification analysis separated sites by geographical location. Most polyhaline and euhaline sites in North Carolina estuaries (Albemarle and Pamlico Sounds) were separated from all others (see Fig. 5). These results are consistent with those of Van Dolah et al. (1999), who found that stations in the Environmental Monitoring and Assessment Carolinian Province (inclusive of North Carolina estuaries) separated by latitude in polyhaline-euhaline habitats and were less distinguishable geographi-

cally at lower salinities. The differences observed in the present study between the higher salinity North Carolina sites and those of similar salinity to the north could have been attributed to biotic discontinuities between the Carolinian and Virginian zoogeographic provinces. We were unable to identify any group of species unique to North Carolina estuaries. Differences in assemblages between North Carolina and Delaware-Chesapeake estuaries were related to the relative abundance contributions of species and not to differences in species composition resulting in geographic discontinuities. The North Carolina species assemblages identified in this study appear to be typical of estuarine habitats of other mid-Atlantic estuaries. Suggestions of a zoogeographic barrier associated with a thermal gradient may exist for coastal shelf waters off North Carolina (Cutler 1975), but our results do not provide evidence for a barrier within the estuary. Engle and Summers (1999b), who observed evidence of latitudinal gradients in the estuarine benthos along the U.S. Atlantic coast (Cape Cod to Biscayne Bay), also found that the benthic fauna of Albemarle-Pamlico Sound were more similar to the Virginian Province fauna than to the fauna of the more southern portions of the Carolinian Province. The Albemarle-Pamlico Sound is usually considered part of the Carolinian Province. Engle and Summers (1999b) suggested that the boundary between the Virginian and Carolinian Provinces, with respect to estuarine benthic assemblages, be moved to a new location south of Albemarle-Pamlico Sound.

In this study, benthic samples from non-degraded reference sites throughout the MAIA region have been classified into habitat categories of similar biological composition. Results provide a basis for understanding how benthic assemblages of the region are distributed in relation to natural environmental factors and for beginning to define reference-range conditions in key benthic attributes by habitat type. This understanding is an essential step in the process of developing a benthic index that can be used region-wide to assess effects of pollution reliably and that takes into account variability related to natural habitat factors.

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