

Quality Assurance Documentation Plan for the  
Microzooplankton Component of the  
Chesapeake Bay Water Quality Monitoring Program

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## Summary

The Plankton component of the Chesapeake Bay Water Quality Monitoring Program has been conducted by the Academy of Natural Sciences' Estuarine Research Center (ANSERC) since August, 1984. The Academy's plankton component includes several elements: microzooplankton and, in two separate quality assurance plans, phytoplankton and horizontal distributions of chlorophyll *a* in the upper Potomac River. The 2000 microzooplankton study is designed to include measurements of the dominant microzooplankton in various layers of the water column and in various size fractions at sampling stations in the Maryland portion of Chesapeake Bay and its tributaries (Figure 1). Techniques for microzooplankton collections and analyses are described below.

## Project Objective

The objective of this component of the Chesapeake Bay Water Quality Monitoring Program is to characterize and enumerate the microzooplankton community of the Bay and its tributaries. The specific parameters measured include qualitative aspects of the microzooplankton communities (species composition) and quantitative measures (species abundance). From the species abundance data, microzooplankton biomass or the amount of carbon in the system which comes from this plankton component, can be calculated. The microzooplankton are the small animals in the plankton that measure between 20 and 200  $\mu\text{m}$  in size. This portion of the plankton is important in linking phytoplankton to the higher trophic levels in the food web. As part of the plankton community, the microzooplankton, along with the phytoplankton, are extremely important because they should be the first communities within the biota to respond to changes in nutrient loads. These responses at the base of the food web could ultimately affect the food availability to higher organisms such as larval stages of fish and shellfish. The Maryland microzooplankton program data are conceivably comparable to the data collected by the Virginia microzooplankton component of the Monitoring Program. The Maryland microzooplankton are also used in the larval striped bass food availability index, devised by Versar, Inc.

## Field Sampling

Samples are collected at 13 stations over the Maryland portion of the Chesapeake Bay and the Patuxent, Potomac, Choptank and Patapsco Rivers in conjunction with sampling programs by MDDNR and Versar, Inc. staff (Figure 1). From April - June, four additional fish spawning stations are sampled as part of the fish larval food availability analysis.

## Net Samples

From January-June, microzooplankton samples are collected with a strategy

designed to maximize sampling during the peak period of fish spawning. Temporally, this translates into a concentrated sampling effort from late-April through mid-June (Table 1). Spatially, this means the addition of four secondary sampling stations, CB2.1, TF2.4, TF1.6 and ET5.0A which are located in historically prolific fish spawning areas (Figure 1). One sampling occurs in the Patuxent River in either January or February. The thirteen primary plankton stations are sampled in March, while all seventeen sites are sampled once a month during April-June, with the tidal-fresh, oligohaline and secondary stations sampled on a second occasion in April and May. The stations which are sampled bi-monthly during this period are: TF2.3, TF1.5, ET5.1, RET2.2, TF1.7, CB2.2, TF2.4, TF1.6, ET5.0A and CB2.1. From July-December, microzooplankton samples are collected monthly (except for during November when no field sampling takes place) at 13 primary stations in the monitoring program.

At stratified mesohaline stations (CB3.3C, CB4.3C, CB5.2, LE1.1, LE2.2, ET5.2 and WT5.1), and tidal fresh Potomac station TF2.3, water from 5 depths above the pycnocline are sampled. At the stations typified by shallow depths and/or generally homogeneous water columns (CB2.1, CB2.2, TF1.5, TF1.6, TF1.7, TF2.4, RET2.2, ET5.0A and ET5.1, ), water is pumped from each of 10 depths over the entire water column into the nets.

Samples are collected using a small diaphragm pump and hose connected to a sampling missile. Water is pumped from depth through a 44  $\mu$ m mesh net.

All samples collected in 44  $\mu$ m mesh nets are decanted into a jar containing buffered formaldehyde (final concentration of 2%) and transferred to the laboratory. Field collections result in 162 samples (including QA/QC samples) enumerated (Table 1).

In the laboratory, the dominant taxa (greater than 44 $\mu$ m in size) in each sample are identified and enumerated using a Leitz Laborlux compound microscope and a total magnification of 100x. Counting continues until at least 250 individuals have been enumerated, yielding a counting error of 10-15% (Venrick, 1978). Meroplankton and mesozooplankton forms are also noted.

### Whole Water Samples

The ciliates are an important component of the microzooplankton assemblage in Chesapeake Bay. The net sampling is an inappropriate means of identifying and quantifying this important taxonomic group because of their size (often < 44 $\mu$ m) and their fragile nature. Therefore, beginning in March, 1998, whole water microzooplankton samples are taken at the mesohaline stations between March - September, in order to quantify the ciliates. The mesohaline stations samples are CB3.3C, CB4.3C, CB5.2, LE1.1, LE2.2, AND ET5.2. Beginning in March, 2000 samples are also taken at the tidal fresh stations TF2.3 and TF1.5. Samples are decanted from the replicate carboys that are collected at five discrete depths above the pycnocline. The whole water microzooplankton samples are preserved with acid Lugol's solution to a final concentration of 2 %. Field collections will result in 42 samples and 2 QA/QC



samples (Table 1).

In the laboratory, subsamples are added to settling chambers and examined on a Leitz Diavert inverted microscope at 200X magnification. A minimum of 100 individual cells or the individuals in 2 settling chambers are enumerated. Tintinnids are identified to the lowest possible taxonomic level based on lorica shape. The soft-bodied ciliates are identified to the lowest possible taxonomic level, based on morphology and size.

### Zebra Mussel Samples

Water samples from CB1.1 are concentrated in a 44  $\mu\text{m}$  mesh net for enumerating and identifying juvenile zebra mussels. Samples are collected once in April and September and bimonthly in May - August. One preserved and one viable sample (kept cool on ice) is returned to the laboratory. The live sample is scanned, using a polarized filter, within 24 h for identification of planktonic zebra mussel veliger larvae.

### Laboratory Methods

#### Net Samples

Each sample is gently mixed and a 1 ml aliquot removed with a Stempel pipet and put into a Sedgwick-Rafter cell for enumeration with a compound microscope at 100X magnification. Beginning with samples collected in April, 1986, a small drop of concentrated Rose Bengal stain is added to the cell prior to addition of the sample. The sample is allowed to set for 10 minutes before counting. At least one chamber (1 ml) is counted for each sample and if the total count does not reach 250 organisms, subsequent 1 ml aliquots are also enumerated until a count of 250 or more organisms is obtained or 3 ml are examined. If a certain organism is abundant (more than 60 per chamber), it is not counted in the subsequent 1 ml aliquots for a given sample. For extremely abundant taxa, less than one ml could be counted. A count program which generates normalized counts of numbers per liter for individual species as well as groups (see section on formulas, calculations and conversions for more information) is used. The NODC taxonomic codes are used for the taxa that are enumerated. Microzooplankton smaller than 44  $\mu\text{m}$  are noted on the original data sheet but not enumerated since estimates are not quantitative.

#### Whole Water Samples

In the lab, 5-25 ml are subsampled from the sample jar for settling. This amount depends on how much detritus and plankton are in the sample. If 25 ml are used, the bottle is shaken gently (slowly inverted 5 times) and 25 ml poured into a graduated cylinder. This is put into a 50 ml settling chamber and the graduated cylinder rinsed 3X. The sample is allowed to settle 48 h before being counted. If less than 25 ml aliquots are used, these are poured into 25 ml settling

chambers which settle for 24 hr before counting.

To count, the entire chamber is examined at 200X with an inverted microscope to obtain a minimum count of 100 organisms. If 100 organisms are not counted, another subsample is settled. Any organism that is abundant in the first aliquot (more than 60) is not counted. The count program used for the net samples (see above) is currently being adapted for use with whole water counts.

### Zebra Mussel Samples

Upon returning to the lab, the viable zebra mussel sample is placed in the refrigerator. Prior to examination for the presence of zebra mussel veliger larvae, the sample is concentrated on a 44µm mesh screen and subsamples of the concentrate are rinsed into a petri dish with filtered station water. A dissecting microscope with polarizing filters is used to screen the entire sample for the presence of veliger larvae. If the fixed sample is to be examined, the sample is allowed to settle for 24 h and the supernatant is drawn off. The sample is then examined using the net sample procedure except 5 separate 1 ml samples are scanned for the larvae. The presence of zebra mussel larvae will be immediately reported to Renee Karrh at MDDNR.

### QA/QC

Random sample recounts of previously counted microzooplankton samples are undertaken in order to determine counting error. One sample/20 samples is blindly selected and recounted. The recount total cell density must fall within 10 % of the total for the original count or the sample is counted again until 2 samples' total densities are within 10 % of one another. The recount and original sample data sheets are stored in a binder in the microscope laboratory at ANSERC.

### Split Samples

Three split samples are taken to compare the techniques used for counting by ANSERC and ODU taxonomists. Net and whole water samples are taken (described above) at the selected stations to be counted by ANSERC. From the same carboys that the whole water samples for ANSERC are taken, 2 one liter samples are taken to be sent to ODU. These are preserved in acid Lugol's solution for a final concentration of 1%. The sample bottle lids are taped and each bottle is put in a separate plastic bag. The bottles are then packed in a box and are hand carried or shipped second day air to ODU.

### Deliverables

Microzooplankton data are delivered to MDDNR in two ways, electronically by April 15, and October 15, and in reports due May 15, and November 15. The data analyzed for these two deliverables cover the period July-December, and January-June, respectively.

## Literature Cited

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Brownlee, D. C. and F. Jacobs. 1987. Mesozooplankton and microzooplankton in the Chesapeake Bay. In *Contaminant Problems of Living Chesapeake Bay Resources*. Edited by S. K. Majumdar, L. W. Hall and H. M. Austin. The Pennsylvania Academy of Sciences. (Pages 245-263)

Dolan, J. R. 1989. The ecology of ciliate microzooplankton in the Chesapeake Bay. PhD Dissertation. University of Maryland, College Park, MD. (Pages 20-21)

Stoecker, D. K., D. J. Gifford and M. Putt. 1994. Preservation of marine planktonic ciliates: losses and cell shrinkage during fixation. *Mar. Ecol. Prog. Ser.* 110:293-299.

Venrick, E.L. 1978. How many cells to count? Pages 167-180 in A. Sournia (ed.), *Phytoplankton manual*. UNESCO, Paris.

## Data Deliverable Information:

### Formulas, Calculations and Conversions

The following equation is used to convert **net sample** raw counts to density (#/L) for each taxon identified:

$$\text{DENSITY} = ((\text{RAWCT} / \text{MLSCNT}) * \text{CONCENTRATE VOLUME}) / \text{TOTAL SPLE VOL}$$

where RAWCT= raw count of the organism

MLSCNT=number of ml of the concentrated sample counted

CONCENTRATE VOLUME=concentration of the sample after it is drawn down in the lab

TOTAL SPLE VOL=number of liters pumped through the net

The following equation is used to convert **whole water sample** counts to density (#/L) for each taxon identified:

$$\text{DENSITY} = (\text{RAWCNT} / \# \text{MLSETTLED}) * 1000 * 1.02 = \# / \text{L} \quad \text{where } 1.02 \text{ is the dilution factor because of the addition of Lugol's solution.}$$

where RAWCNT= raw count of the organism

#MLSETTLED= number of ml of the sample put into the settling chamber

#/L=abundance of the organism in numbers per liter

### Species Inhouse Code, Reference Code, and Scientific Name

The inhouse code is the same as the reference code with the addition of an inhouse modifier in column 13. The modifier in the 13<sup>th</sup> column has different meanings depending on the taxa being considered. For larger organisms, a one in this column indicates that the organism is in its larval or naupliar stage. For smaller microzooplankton such as rotifers and non-loricate ciliates, this modifier may indicate a size category within a taxon. The NODC Code does not distinguish between life history stages. For tintinnids (3540...) a 0,1,2 (or 3, 4, 5) in the 13<sup>th</sup> column indicates that it was not or could not be determined if the lorica contained a cell, that the lorica did contain a cell (full), or that the lorica did not contain a cell (empty), respectively. For an undescribed species, the code is given down to the lowest possible taxonomic level, the unknown levels are given zeros and a number is assigned in the subspecies columns (11-12) (e.g. *Brachionus A* = 4506010400010).

Only organisms greater than 44µm and less than 200 µm in their smallest dimension are included in the species list. The microzooplankton less than 44µm are noted and recorded as either rare, common, abundant or dominant.

NODC Code with in-house modifications (13th column) and scientific name:

#### DIGIT REPRESENTS

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- 1-2 Phylum
- 3-4 Class and Order
- 5-6 Family
- 7-8 Genus
- 9-10 Species
- 11-12 Subspecies
- 13 Special in-house modifier (for explanation, refer to data dictionary)

#### CODE

#### SPECIES

0000000000001 UNIDED LARVAE  
0000000000002 UNIDED TROCHOPHORE LARVAE  
3438000000000 SARCODINA-UNIDED SARCODINID  
3442010000000 DIFFLUGIIDAE

3442010100000 DIFFLUGIA SPP.  
3442010201000 LESQUEREUSIA GIBBOSA  
3442020100000 ARCELLA SP.  
3442030000000 CENTROPYXIDAE  
3442030100000 CENTROPYXIS SP.  
3442030101000 CENTROPYXIS ACULEATA  
3442040000000 PARAQUADRULIDAE  
3442040200000 QUADRULELLA SP.  
3442050000000 HYALOSPHEINIIDAE-TESTATED AMOEBA  
3445020000000 EUGLYPHA SP.  
3445040000000 CYPHODERIIDAE  
3445040100000 CYPHODERIA SP.  
3448000000000 FORAMINIFERIDA  
3512000000000 CILIOPHORA-UNIDED CILIATE  
3516000000000 HAPTORIDA  
3516010100000 DIDINIUM SP.  
3516010202000 MESODINIUM RUBRUM  
3516020000001 TRACHELOCERCIDAE-LARGE  
3517000000001 CYRTOPHORIDA-LARGE  
3530000000001 PERITRICHIDA-LARGE  
3531000000000 SESSILINA-UNIDED SESSILINE PERITRICH  
3532000000000 MOBILINA-UNIDED MOBILINE PERITRICH  
3533000000000 SUCTORIA-UNIDED SUCTORIAN  
3534010100000 ACINETA SP.  
3534030700000 STAUROPHRYA SP.  
3534040100000 EPHELOTA SP.  
3537000000000 HETEROTRICHINA-UNIDED HETEROTRICH  
3539000000000 OLIGOTRICHINA-UNIDED OLIGOTRICH  
3539020100000 STENTOR SP.  
3539030000000 STROMBIDIDAE  
3539030100000 STROMBIDIUM SP.  
3539030200000 TONTONIA SP.  
3540000000000 TINTINNINA-UNIDED TINTINNID  
3540010100050 TINTINNIDIUM SP.-LARGE  
3540010100051 TINTINNIDIUM SP.-LARGE-FULL  
3540010100052 TINTINNIDIUM SP.-LARGE-EMPTY  
3540020100000 TINTINNOPSIS SP.  
3540020100001 TINTINNOPSIS SP.-FULL  
3540020100002 TINTINNOPSIS SP.-EMPTY  
3540020100030 TINTINNOPSIS SUBACUTA-HUGE  
3540020100031 TINTINNOPSIS SUBACUTA-HUGE-FULL  
3540020100032 TINTINNOPSIS SUBACUTA-HUGE-EMPTY  
3540020100050 TINTINNOPSIS SP. A

3540020100051 TINTINNOPSIS SP. A-FULL  
3540020100052 TINTINNOPSIS SP. A-EMPTY  
3540020105000 TINTINNOPSIS DADAYI  
3540020105001 TINTINNOPSIS DADAYI-FULL  
3540020105002 TINTINNOPSIS DADAYI-EMPTY  
3540020123000 TINTINNOPSIS FIMBRIATA  
3540020123001 TINTINNOPSIS FIMBRIATA-FULL  
3540020123002 TINTINNOPSIS FIMBRIATA-EMPTY  
3540020123003 TINTINNOPSIS FIMBRIATA-MEUNIERI GRP  
3540020123004 TINTINNOPSIS FIMBRIATA-MEUNIERI GRP-FULL  
3540020123005 TINTINNOPSIS FIMBRIATA-MEUNIERI GRP-EMPTY  
3540020129000 TINTINNOPSIS RADIX  
3540020129001 TINTINNOPSIS RADIX-FULL  
3540020129002 TINTINNOPSIS RADIX-EMPTY  
3540020133000 TINTINNOPSIS SUBACUTA  
3540020133001 TINTINNOPSIS SUBACUTA-FULL  
3540020133002 TINTINNOPSIS SUBACUTA-EMPTY  
3540020136000 TINTINNOPSIS MEUNIERI  
3540020136001 TINTINNOPSIS MEUNIERI-FULL  
3540020136002 TINTINNOPSIS MEUNIERI-EMPTY  
3540020137000 TINTINNOPSIS KARAJACENSIS  
3540020137001 TINTINNOPSIS KARAJACENSIS-FULL  
3540020137002 TINTINNOPSIS KARAJACENSIS-EMPTY  
3540020138000 TINTINNOPSIS NITIDA  
3540020138001 TINTINNOPSIS NITIDA-FULL  
3540020138002 TINTINNOPSIS NITIDA-EMPTY  
3540030100010 STENOSEMELLA SP.A  
3540030100011 STENOSEMELLA SP.A-FULL  
3540030100012 STENOSEMELLA SP.A-EMPTY  
3540050100020 METACYLIS SP. B  
3540050100021 METACYLIS SP. B-FULL  
3540050100022 METACYLIS SP. B-EMPTY  
3540050100030 METACYLIS SP. C  
3540050100031 METACYLIS SP. C-FULL  
3540050100032 METACYLIS SP. C-EMPTY  
3540050400000 CLIMACOCYLIS SP.  
3540050400001 CLIMACOCYLIS SP.-FULL  
3540050400002 CLIMACOCYLIS SP.-EMPTY  
3540050501000 STYLICAUDA PLATENSIS  
3540050501001 STYLICAUDA PLATENSIS-FULL  
3540050501002 STYLICAUDA PLATENSIS-EMPTY  
3540070100000 FAVELLA SP.  
3540070100001 FAVELLA SP.-FULL

3540070100002 FAVELLA SP.-EMPTY  
3540130100010 EUTINTINNUS SP.A  
3540130100011 EUTINTINNUS SP.A-FULL  
3540130100012 EUTINTINNUS SP.A-EMPTY  
3543000000000 HYPOTRICHIDA  
3545010000000 EUPLOTIDAE  
3545010100001 EUPLOTES SPP.-LARGE  
3545010100020 EUPLOTES SP. A  
3545010100030 EUPLOTES SP. B  
4400000000000 GASTROTRICHA  
4500000000000 ROTIFERA-UNIDED ROTIFER  
4500000000010 ROTIFER A  
4500000000020 ROTIFER B  
4500000000030 ROTIFER C  
4500000000040 ROTIFER D  
4500000000050 ROTIFER E  
4500000000060 ROTIFER F  
4504000000000 BDELLOIDA-UNIDED BDELLOID ROTIFER  
4504020100000 ROTARIA SP.  
4504020103000 ROTARIA CITRINUS  
4504020104000 ROTARIA NEPTUNIA  
4504020300000 MACROTRACHELA SP.  
4506010100000 KERATELLA SP.  
4506010100010 KERATELLA SP. A  
4506010102000 KERATELLA QUADRATA  
4506010103000 KERATELLA COCHLEARIS  
4506010103020 KERATELLA COCHLEARIS COCHLEARIS  
4506010103030 KERATELLA COCHLEARIS HISPIDA  
4506010103040 KERATELLA COCHLEARIS MICRACANTHA  
4506010103050 KERATELLA COCHLEARIS ROBUSTA  
4506010103060 KERATELLA COCHLEARIS TECTA  
4506010104000 KERATELLA CRASSA  
4506010105000 KERATELLA EARLINAE  
4506010106000 KERATELLA VALGA  
4506010200000 NOTHOLCA SP.  
4506010203000 NOTHOLCA ACUMINATA  
4506010300000 COLURELLA SP.  
4506010400000 BRACHIONUS SP.  
4506010400010 BRACHIONUS SP. A  
4506010400020 BRACHIONUS SP. B  
4506010401000 BRACHIONUS PLICATILIS  
4506010402000 BRACHIONUS CALYCIFLORUS  
4506010403000 BRACHIONUS HAVANAENSIS

4506010404000 BRACHIONUS PTERODINOIDES  
4506010405000 BRACHIONUS URCEOLARIS  
4506010406000 BRACHIONUS ANGULARIS  
4506010407000 BRACHIONUS BIDENTATA  
4506010408000 BRACHIONUS BUDAPESTINENSIS  
4506010409000 BRACHIONUS CAUDATUS  
4506010410000 BRACHIONUS DIVERSICORNIS  
4506010411000 BRACHIONUS QUADRIDENTATUS  
4506010412000 BRACHIONUS RUBENS  
4506010413000 BRACHIONUS VARIABILIS  
4506010500000 KELLICOTTIA SP.  
4506010501000 KELLICOTTIA LONGISPINA  
4506010502000 KELLICOTTIA BOSTONIENSIS  
4506010700000 LEPADELLA SP.  
4506010704000 LEPADELLA PATELLA  
4506010800000 ANURAEOPSIS SP.  
4506010801000 ANURAEOPSIS FISSA  
4506010900000 EPIPHANES SP.  
4506011000000 EUCHLANIS SP.  
4506011001000 EUCHLANIS DILATATA  
4506011100000 LOPHOCHARIS SP.  
4506011101000 LOPHOCHARIS SALPINA  
4506011200000 MACROCHAETUS SP.  
4506011300000 MYTILINA SP.  
4506011400000 PLATYIAS SP.  
4506011401000 PLATYIAS PATULUS  
4506011402000 PLATYIAS QUADRICORNIS  
4506011500000 TRICHOTRIA SP.  
4506011501000 TRICHOTRIA TTRACTIS  
4506020100000 LECANE SP.  
4506020200000 MONOSTYLA SP.  
4506020201000 MONOSTYLA BULLA  
4506020202000 MONOSTYLA CLOSTEROCERCA  
4506020203000 MONOSTYLA QUADRIDENTATA  
4506040100000 ENCENTRUM SP.  
4506040200000 PROALES SP.  
4506040300000 CEPHALODELLA SP.  
4506040302000 CEPHALODELLA GIBBA  
4506040400000 NOTOMMATA SP.  
4506040500000 MONOMMATA SP.  
4506040600000 EOSPHORA SP.  
4506070100000 TRICHOCERCA SP.  
4506070102000 TRICHOCERCA CYLINDRICA



4506070103000 TRICHOCERCA LONGISETA  
4506070104000 TRICHOCERCA MULTICRINIS  
4506070105000 TRICHOCERCA SIMILIS  
4506080100000 ASCOMORPHA SP.  
4506080101000 ASCOMORPHA OVALIS  
4506080102000 ASCOMORPHA SALTANS  
4506080200000 GASTROPUS SP.  
4506080201000 GASTROPUS MINOR  
4506120100000 ASPLANCHNA SP.  
4506120101000 ASPLANCHNA BRIGHTWELLI  
4506120102000 ASPLANCHNA HERRICKI  
4506120103000 ASPLANCHNA PRIODONTA  
4506130200000 SYNCHAETA SP.  
4506130200001 SYNCHAETA SPP. L-LARGE  
4506130200002 SYNCHAETA SPP. M-MEDIUM  
4506130200003 SYNCHAETA SPP. S-SMALL  
4506130200010 SYNCHAETA BICORNIS  
4506130200020 SYNCHAETA BALTICA  
4506130204000 SYNCHAETA PECTINATA  
4506130206000 SYNCHAETA OBLONGA  
4506130207000 SYNCHAETA STYLATA  
4506130300000 POLYARTHRA SP.  
4506130302000 POLYARTHRA DISSIMULANS  
4506130303000 POLYARTHRA DOLICHOPTERA  
4506130304000 POLYARTHRA EURYPTERA  
4506130305000 POLYARTHRA MAJOR  
4506130306000 POLYARTHRA REMATA  
4506130307000 POLYARTHRA VULGARIS  
4506130400000 PLOESOMA SP.  
4506130401000 PLOESOMA HUDSONI  
4506130402000 PLOESOMA TRUNCATUM  
4507010100000 TESTUDINELLA SP.  
4507010101000 TESTUDINELLA PATINA  
4507020100000 HEXARTHRA SP.  
4507020101000 HEXARTHRA MIRA  
4507040100000 CONOCHILOIDES SP.  
4507040102000 CONOCHILOIDES DOSSUARIUS  
4507040103000 CONOCHILOIDES NATANS  
4507040200000 CONOCHILUS SP.  
4507040201000 CONOCHILUS HIPPOCREPIS  
4507040202000 CONOCHILUS UNICORNIS  
4507050100000 FILINIA SP.  
4507050101000 FILINIA LONGISETA

4507050102000 FILINIA BRACHIATA  
 4507050103000 FILINIA TERMINALIS  
 4508010100000 COLLOTHECA SP.  
 4508010101000 COLLOTHECA MUTABILIS  
 4508010102000 COLLOTHECA PELAGICA  
 4700000000000 NEMATODA  
 5100000000001 GASTROPODA-LARVAE  
 5500000000001 PELECYPODA-LARVAE  
 5515370301000 DREISSENA POLYMORPHA  
 5922000000000 ACARINA-MITE  
 6117000000001 COPEPOD NAUPLII  
 6117000000005 COPEPOD NAUPLII+PERITRICHIS  
 6118180100001 DIAPTOMUS -NAUPLII  
 6118190200001 PSEUDODIAPTOMUS SP.-NAUPLII  
 6118200200001 EURYTEMORA SP.-NAUPLII  
 6118290100001 ACARTIA SP.-NAUPLII  
 6119050200001 SCOTTOLANA SP.-NAUPLII  
 6120080200001 CYCLOPS SP.-NAUPLII  
 6120080300001 MESOCYCLOPS SP.-NAUPLII  
 6120090100001 OITHONA SP.-NAUPLII  
 6120260100001 HEMICYCLOPS SP.-NAUPLII  
 7500000000000 TARTIGRADA

Numerical Variable Names

VARIABLE	VALID RANGE
-----	-----
DATE	840802-971218
TIME	0651-1935
TDEPTH	1.8-33
PDEPTH	0.5-22.0
VOLPDEPT	2-20
TOTVCOMP	12-200
SER_NUM	01001-240039
A_DEPTH1	0.2- 1.0
A_DEPTH2	0.4- 5.0
A_DEPTH3	0.5-10.0
A_DEPTH4	0.5-15.0

A\_DEPTH5 0.5-21.5  
 B\_DEPTH1 1-22.5  
 B\_DEPTH2 1-25.0  
 B\_DEPTH3 1-27.0  
 B\_DEPTH4 1-29.0  
 B\_DEPTH5 1-32.0  
 CONCENT 10-500  
 MLSCNT 0.01-10  
 RAWCT 1-9999  
 DENSITYV 0.05-10000.00  
 TDENSITY 1-999999.99

# CHARACTER VARIABLES - VALID VALUES

VARIABLE      VALID VALUES

-----      -----

REPNUM

1  
 2  
 3  
 4  
 T  
 B  
 W

TRIB\_COD

BAY  
 PAX  
 POT  
 PAT  
 CHS  
 CHP

SPECCODE

0000000000001-7500000000000

STATION

CB1.1  
 CB2.1  
 CB2.2  
 CB3.3C  
 CB4.3C

CB5.2  
LE1.1  
TF1.7  
TF1.6  
TF1.5  
TF2.3  
TF2.4  
RET2.2  
LE2.2  
ET4.2  
ET5.1  
ET5.0A  
ET5.2  
EE3.1  
WT5.1

#### Station Names, Numbers and Descriptions

CB1.1 01 Mouth of Susquehanna River, main Bay  
CB2.1 02 SW of Turkey Point, main Bay  
CB2.2 03 W of Still Pond near Buoy R 50, main Bay  
CB3.3C 07 N of Bay Bridge, main Bay  
CB4.3C 16 E of Dares Beach near Buoy R 78, main Bay  
CB5.2 20 E of Point No Point, main Bay  
LE1.1 28 Between Jack Bay sandspit and Sandgates in mid channel, Patuxent River  
TF1.7 30 ESE of Jacks Creek in mid channel, Patuxent River  
TF1.6 31 Midchannel off the wharf at Lower Marlboro  
TF1.5 32 At Nottingham in mid channel, Patuxent River  
TF2.3 35 Off Indian Head at Buoy N 54, Potomac River  
TF2.4 48 Buoy 44 between Possum Point and Moss Point  
RET2.2 34 Off Maryland Point at Buoy 19, Potomac River  
LE2.2 33 Off Ragged Point at buoy BW 51B, Potomac River (Prior to October 1988 data  
tape, this station was designated XBE9541)  
ET4.2 37 South of Eastern Neck Island at Buoy 9, lower Chester River  
ET5.1 38 Downstream of confluence with Tuckahoe Creek, upper Choptank River  
ET5.0A 49 Midchannel off the mouth of King's Creek  
ET5.2 39 Near Rt 50 bridge at Cambridge, lower Choptank River  
EE3.1 40 North Tangier Sound North of Buoy R 16, main Bay  
WT5.1 36 East of Hawkins Point at Buoy 5M, Patapsco River (Baltimore Harbor)

#### Station names, Latitudes, Longitudes and Total Depths (meters)

CB1.1	39-32.7	76-04.9	6.1
CB2.1	39-26.4	76-01.5	6.2
CB2.2	39-20.8	76-10.5	12.1
CB3.3C	38-59.7	76-21.6	23.7
CB4.3C	38-33.4	76-26.1	26.1
CB5.2	38-08.2	76-13.7	30.1
LE1.1	38-25.5	76-36.1	12.0
TF1.7	38-35.0	76-40.7	3.0
TF1.6	38-39.5	76-41.1	6.2
TF1.5	38-42.7	76-42.2	10.3
TF2.3	38-36.3	77-10.3	12.7
TF2.4	38-31.5	77-16.6	9.0
RET2.2	38-21.1	77-12.2	9.5
LE2.2	38-09.6	76-35.8	11.0
ET4.2	38-59	76-13	14.6
ET5.1	38-48.4	75-54.4	5.3
ET5.0A	38-46.5	75-58.1	11.0
ET5.2	38-34.8	76-02.5	12.3
EE3.1	38-12	75-58	13.7
WT5.1	39-12.7	76-31.4	15.7

#### Data Entry Method

Computer keyboard to disk from data sheets for sampling trips prior to April 1985, and keyboard to disk for trips from April 1985 through December 1994. Starting in January 1995, computer keyboard to inhouse computer network.

#### Data Verification

Visual inspection and computer verification program.

#### Variable Names, SAS Labels and Descriptions

ANSMZyyH.SD2 - common sample information may be merged with counts using SER\_NUM where yy is year of data set

#### List of Variables

VARIABLE	TYPE	LENGTH	POSITION	FORMAT	LABEL
TRIB_COD	CHAR	3	0		SPECIFIES TRIB. SAMPLED (OR MAIN BAY)
SER_NUM	NUM	8	3		SERIAL NUM. INCL. SAMPLING EFFORT NUM.
STATION	CHAR	7	11		STATION NAME
DATE	NUM	8	18	DATE7.	SAMPLE DATE
TIME	NUM	8	26	HHMM5.	SAMPLE TIME
GMETHOD	CHAR	1	34		GEAR METHOD CODE
TDEPTH	NUM	8	35		TOTAL DEPTH
PDEPTH	NUM	8	43		DEPTH INCLUDED IN SURFACE COMPOSITE
VOLPDEPT	NUM	8	51		VOLUME OF SAMPLE PER DEPTH
TOTVCOMP	NUM	8	59		TOTAL VOLUME OF COMPOSITE SAMPLE
A_DEPTH1	NUM	8	67		1st SAMPLE DEPTH ABOVE PDEPTH
A_DEPTH2	NUM	8	75		2nd SAMPLE DEPTH ABOVE PDEPTH
A_DEPTH3	NUM	8	83		3rd SAMPLE DEPTH ABOVE PDEPTH
A_DEPTH4	NUM	8	91		4th SAMPLE DEPTH ABOVE PDEPTH
A_DEPTH5	NUM	8	99		5th SAMPLE DEPTH ABOVE PDEPTH
B_DEPTH1	NUM	8	107		1st SAMPLE DEPTH BELOW PDEPTH
B_DEPTH2	NUM	8	115		2nd SAMPLE DEPTH BELOW PDEPTH
B_DEPTH3	NUM	8	123		3rd SAMPLE DEPTH BELOW PDEPTH
B_DEPTH4	NUM	8	131		4th SAMPLE DEPTH BELOW PDEPTH
B_DEPTH5	NUM	8	139		5th SAMPLE DEPTH BELOW PDEPTH
FC_INIT	CHAR	3	147		INITIALS OF FIELD COLLECTOR
LI_INIT	CHAR	3	150		INITIALS OF INDIVID. PERFORMING LAB ID.

ANSMZyyD.SD2 - counts data set

LIST OF VARIABLES AND ATTRIBUTES BY POSITION

VARIABLE	TYPE	LENGTH	POSITION	FORMAT	LABEL
SER_NUM	NUM	8	0		SERIAL NUM. INCL. SAMPLING EFFORT NUM.
STATION	CHAR	7	8		STATION NAME
DATE	NUM	8	15	DATE7.	SAMPLE DATE
TRIB_COD	CHAR	3	23		SPECIFIES TRIB. SAMPLED (OR MAIN BAY)
LAYER	CHAR	2	26		AP=ABOVE PYC,BP=BELOW PYC, WC= WATER COL
REPNUM	CHAR	1	28		REPLICATE NUMBER
CONCENT	NUM	8	29		VOLUME OF CONCENTRATED SAMPLE
MLSCNT	NUM	8	37		MILLILITERS COUNTED
SPECCODE	CHAR	13	45		SPECIES CODE
RAWCT	NUM	8	58		RAW COUNT
DENSITYV	NUM	8	66		DENSITY (# INDIV./L) OF GIVEN TAXON
TDENSITY	NUM	8	74		DENSITY (# INDIV./L) OF ALL ORGANISM

## **Appendix**

Figure 1. Map of Station Sites

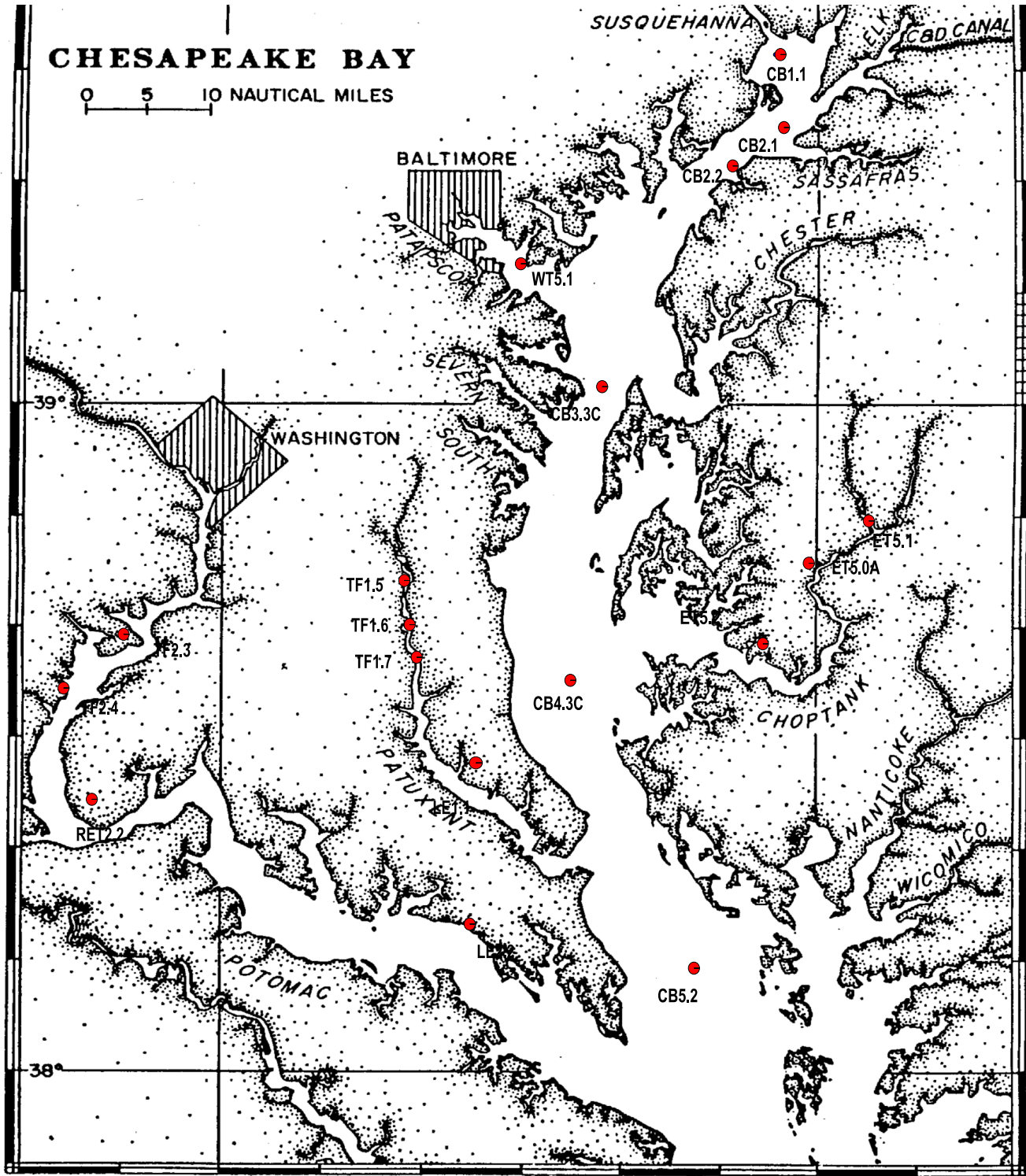




Table 1. Total Sample Numbers for Microzooplankton for the ANSERC Plankton Program

Measurements	J	A	S	O	N	D	J	F	M	A	M	J	QA/QC	Total
Net Samples	13	13	13	13	--	13	--	03	13	27	27	17	8	160
Whole Water Samples	08	08	08	--	--	--	--	--	08	08	08	08	3	59
Viable Zebra Mussels	02	02	01	--	--	--	--	--	--	01	02	01	--	9

Net Samples are taken from the following stations: CB2.2, CB3.3C, CB4.3C, CB5.2, WT5.2, ET5.1, ET5.2, TF1.5, TF1.7, LE1.1, TF2.3, RET2.2, LE2.2 and CB2.1, ET5.0A, TF2.4, TF1.6 during fish spawning season.

Whole water samples are taken from the following stations: CB3.3C, CB4.3C, CB5.2, ET5.2, LE1.1, LE2..2, TF2.3, TF1.5.

Zebra mussel samples are taken from station CB1.1.

## Examples of Laboratory Data Sheet with Original Counts and Recounts

249003 MCB2.2 980708 MZP W

MICROZOOPLANKTON > 44 UM

NUMBER OF LITERS FILTERED 100.0  
CONCENTRATE VOLUME (ML) 44.0

SPECIES CODE	SPECIES NAME	RAW CNT	#MLS CNTD	NORM CNT (#/L)
3540020123003	TINTINNOPSIS FIMBRIATA-MEUNIERI GRP	195	0.50	171.60
6118290100001	ACARTIA SP.-NAUPLII	41	1.00	18.04
4506130200003	SYNCHAETA SPP. S-SMALL	10	1.00	4.40
5500000000001	PELECYPODA-LARVAE	7	1.00	3.08
4506010103060	KERATELLA COCHLEARIS TECTA	6	1.00	2.64
4506130300000	POLYARTHRA SP.	6	1.00	2.64
6117000000001	COPEPOD NAUPLII	4	1.00	1.76
4506070100000	TRICHOCERCA SP.	3	1.00	1.32
4506010406000	BRACHIONUS ANGULARIS	3	1.00	1.32
4506130200010	SYNCHAETA BICORNIS	2	1.00	0.88
4506010103020	KERATELLA COCHLEARIS COCHLEARIS	2	1.00	0.88
4506010402000	BRACHIONUS CALYCIFLORUS	1	1.00	0.44

MICROZOOPLANKTON > 44UM RAW COUNT FOR ENTIRE SAMPLE 280

MICROZOOPLANKTON >44UM (#/L) FOR ENTIRE SAMPLE 209.0

249003 MCB2.2 980708 MZP W RECOUNT

MICROZOOPLANKTON > 44 UM

NUMBER OF LITERS FILTERED 100.0  
CONCENTRATE VOLUME (ML) 44.0

SPECIES CODE	SPECIES NAME	RAW CNT	#MLS CNTD	NORM CNT (#/L)
3540020123003	TINTINNOPSIS FIMBRIATA-MEUNIERI GRP	171	0.50	150.48
6118290100001	ACARTIA SP.-NAUPLII	46	1.00	20.24
4506130200003	SYNCHAETA SPP. S-SMALL	18	1.00	7.92
4506010103060	KERATELLA COCHLEARIS TECTA	14	1.00	6.16
5500000000001	PELECYPODA-LARVAE	12	1.00	5.28
6117000000001	COPEPOD NAUPLII	12	1.00	5.28
4506010103020	KERATELLA COCHLEARIS COCHLEARIS	6	1.00	2.64
4506070100000	TRICHOCERCA SP.	6	1.00	2.64
4506010400000	BRACHIONUS SP.	3	1.00	1.32
4506130300000	POLYARTHRA SP.	2	1.00	0.88
4506010406000	BRACHIONUS ANGULARIS	2	1.00	0.88
4506130200010	SYNCHAETA BICORNIS	1	1.00	0.44
4506010409000	BRACHIONUS CAUDATUS	1	1.00	0.44

MICROZOOPLANKTON > 44UM RAW COUNT FOR ENTIRE SAMPLE 294

MICROZOOPLANKTON >44UM (#/L) FOR ENTIRE SAMPLE 204.6

## Microzooplankton Data Dictionary

# NAME OF DATA FILE DESCRIBED BY THIS DATA DICTIONARY FILE

ANSMZ84H.SD2, ANSMZ84D.SD2

ANSMZ85H.SD2, ANSMZ85D.SD2

ANSMZ86H.SD2, ANSMZ86D.SD2

ANSMZ87H.SD2, ANSMZ87D.SD2

ANSMZ88H.SD2, ANSMZ88D.SD2

ANSMZ89H.SD2, ANSMZ89D.SD2

ANSMZ90H.SD2, ANSMZ90D.SD2

ANSMZ91H.SD2, ANSMZ91D.SD2

ANSMZ92H.SD2, ANSMZ92D.SD2

ANSMZ93H.SD2, ANSMZ93D.SD2

ANSMZ94H.SD2, ANSMZ94D.SD2

ANSMZ95H.SD2, ANSMZ95D.SD2

ANSMZ96H.SD2, ANSMZ96D.SD2

ANSMZ97H.SD2, ANSMZ97D.SD2

ANSMZ28H.SD2, ANSMZ28D.SD2

MMZP.SD2

# NAMES AND DESCRIPTIONS OF ASSOCIATED DATA DICTIONARY FILES

MZPDAT4.DIC, MZPDAT5.DIC, MZPDAT6.DIC, MZPDAT7.DIC

MZPDAT8.DIC, MZPDAT9.DIC, MZPDAT10.DIC, MZPDAT11.DIC

MZPDAT12.DIC, MZPDAT13.DIC, MZPDAT14.DIC, ANSMZP15.DIC

ANSMZP16.DIC, ANSMZP17.DIC, ANSMZP18.DIC, ANSMZP19.DIC

ANSMZP20.DIC, ANSMZP21.DIC, ANSMZP22.DIC, ANSMZP23.DIC

ANSMZP24.DIC, ANSMZP25.DIC, ANSMZP26.DIC, ANSMZP27.DIC

ANSMZP28.DIC

# PROJECT TITLE

Chesapeake Bay Monitoring Program

Microzooplankton Study from August 1984 through December 1999

Field Collection by ANSERC staff.

Sample analysis by R. E. Jacobsen, S. G. Sellner (Brownlee), S. S. Hedrick,

B. B. Wagoner, James H. Sniezek, Kimberly M. Burke, Ralph Matos, Jr.

Data Files Verified by S. G. Sellner (Brownlee)

ANSERC

# PRINCIPAL INVESTIGATORS

>PROGRAM MANAGER: R. V. Lacouture / K. G. Sellner

>MICROZOOPLANKTON: R. V. Lacouture/K. G. Sellner/D. C. Brownlee

>STATISTICIAN:

>PROGRAMMER/ANALYST: A. L. Imirie/ M. C. Marsh / S. E. Gunsalus

>DATA COORDINATOR: S. G. Sellner (Brownlee)

# FUNDING AGENCIES

Maryland Department of Natural Resources

# PROJECT COST

\$292,430 (July 1,1999 - June 30, 2000)

# QA/QC OFFICER

None

# LOCATION OF STUDY

Chesapeake Bay and tributaries in State of Maryland

# DATE INTERVALS:

840802 - 991213

# ABSTRACT

Beginning in August, 1984, field collections of 2 surface and 2 bottom replicate samples of microzooplankton were made monthly (usually excluding February) at 16 stations in conjunction with 3 other plankton elements of ANS portion of the DNR Water Quality Program. From August of 1984 until November 1984, 25 L of water were filtered through the net per replicate. After this time, usually, for each surface and bottom replicate 10 l of water at each of 5 depths were pumped through a 44 um mesh net, resulting in 50 l being filtered for each sample. After June, 1986, stations MET4.2 and MEE3.1 were no longer sampled. After March, 1985, replicate surface samples were combined for each station yielding 1 surface composite sample that had 20 l of water filtered for each of 5 depths resulting in 100 l filtered for each sample; bottom replicates were also combined. Beginning July, 1989 entire water column samples of 10 l from each of 10 depths (total of 100 l) were collected for stations XDA1177, XED4892, PXT0402, MET5.1, MCB1.1 and MCB2.2. Between August, 1984 and September, 1985, 1 ml of 1% neosyneprine was added to each concentrated sample. The sample was allowed to set for about 30 min then formaldehyde was added. Following a study which showed no significant difference in contraction between microzooplankton treated or not treated with neosyneprine, the neosyneprine step was eliminated. Instead buffered formaldehyde (final concentration approximately 2.5%) was added to each sample jar prior to the addition of the sample. Numbers and species identifications were subsequently made using repeated counts on 1 ml aliquots in Sedgewick-Rafter cells and a compound microscope (total mag = 100X). Beginning with samples collected in April 1986, a small drop of concentrated Rose Bengal in formaldehyde was added to the Sedgewick-Rafter cell before adding the sample. The counting cell was allowed to set for 10 minutes before counting. The NODC species code was employed. Microzooplankton smaller than 44 um were noted but not enumerated in counts after March 1985 since estimates would be non-quantitative. In May, 1992, 1993 & 1994 microzooplankton samples for stations MCB1.1, MCB2.2, XED4892, PXT0402, XDA1177, XEA6596, MET5.1 and MET5.2 were sampled twice to coincide with white perch and striped bass spawning periods. From April, 1993 through June, 1993 and again in 1994 and 1995 for the same time period, an additional station, MCB2.1, in the upper Chesapeake Bay was also sampled to coincide with the spawning periods.

In April, 1996, 3 more tidal fresh stations XEA1840 in the Potomac River, XED9490 in the Patuxent River, and MET5.0 in the Choptank River were added for microzooplankton sampling in April, May, and June. Stations MCB2.2, MCB2.1, XEA6596, XEA1840, XDA1177, PXT0402, XED9490, XED4892, MET5.1, and MET5.0 were sampled twice in April and May, again to coincide with white perch and striped bass spawning periods. Main Bay stations MCB1.1 and MCB5.2 were no longer sampled as of March, 1996. Sampling in November was discontinued in 1996.

# STATION NAMES NUMBERS AND DESCRIPTIONS

MCB1.1 01 Mouth of Susquehanna River, main Bay

MCB2.1 02 SW of Turkey Point, main Bay

MCB2.2 03 W of Still Pond near Buoy R 50, main Bay  
 MCB3.3C 07 N of Bay Bridge, main Bay  
 MCB4.3C 16 E of Dares Beach near Buoy R 78, main Bay  
 MCB5.2 20 E of Point No Point, main Bay  
 XDE5339 28 Between Jack Bay sandspit and Sandgates in mid channel, Patuxent River  
 XED4892 30 ESE of Jacks Creek in mid channel, Patuxent River  
 XED9490 31 Midchannel off the wharf at Lower Marlboro  
 PXT0402 32 At Nottingham in mid channel, Patuxent River  
 XEA6596 35 Off Indian Head at Buoy N 54, Potomac River  
 XEA1840 48 Buoy 44 between Possum Point and Moss Point  
 XDA1177 34 Off Maryland Point at Buoy 19, Potomac River  
 MLE2.2 33 Off Ragged Point at buoy BW 51B, Potomac River (Prior to October 1988 data  
 tape, this station was designated XBE9541)  
 MET4.2 37 South of Eastern Neck Island at Buoy 9, lower Chester River  
 MET5.1 38 Downstream of confluence with Tuckahoe Creek, upper Choptank River  
 MET5.0 49 Midchannel off the mouth of King's Creek  
 MET5.2 39 Near Rt 50 bridge at Cambridge, lower Choptank River  
 MEE3.1 40 North Tangier Sound North of Buoy R 16, main Bay  
 MWT5.1 36 East of Hawkins Point at Buoy 5M, Patapsco River (Baltimore Harbor)

# STATION NAMES, LATITUDES, LONGITUDES, AND TOTAL DEPTHS (METERS)

MCB1.1	39-32.7	76-04.9	6.1
MCB2.1	39-26.4	76-01.5	6.2
MCB2.2	39-20.8	76-10.5	12.1
MCB3.3C	38-59.7	76-21.6	23.7
MCB4.3C	38-33.4	76-26.1	26.1
MCB5.2	38-08.2	76-13.7	30.1
XDE5339	38-25.5	76-36.1	12.0
XED4892	38-35.0	76-40.7	3.0
XED9490	38-39.5	76-41.1	6.2
PXT0402	38-42.7	76-42.2	10.3
XEA6596	38-36.3	77-10.3	12.7
XEA1840	38-31.5	77-16.6	9.0
XDA1177	38-21.1	77-12.2	9.5
MLE2.2	38-09.6	76-35.8	11.0
MET4.2	38-59	76-13	14.6
MET5.1	38-48.4	75-54.4	5.3
MET5.0	38-46.5	75-58.1	11.0
MET5.2	38-34.8	76-02.5	12.3
MEE3.1	38-12	75-58	13.7
MWT5.1	39-12.7	76-31.4	15.7

# METHODOLOGY DESCRIBING CHAIN OF CUSTODY FOR LAB SAMPLES

Microzooplankton samples were collected by a member of the Benedict Estuarine Research Center biomonitoring section. On return to the laboratory, the samples were transferred to the BERC plankton laboratory. Counts and identifications were then made and sample concentrates were subsequently archived.



# MONITORING QA/QC PLAN FOR PROJECT

For each monthly microzooplankton collection, one sample was randomly selected as the QA/QC sample. Two separate counts were performed using the same enumeration techniques on 2 individual aliquots removed from the same jar.

>PARAMETER: Microzooplankton counts

- COLLECTION METHODS: Samples were pumped from 5 depths in the surface and 5 depths in the bottom layers and each composite was filtered through a 44 um mesh net and rinsed into a jar. After February, 1985, the 2 top surface layer replicates and 2 bottom layer replicates were each combined in the lab and counted as one sample. Beginning July, 1985, replicate samples were combined in the field with the five depths for the surface and bottom being pumped through the net and rinsed into their respective jars 2X.

Beginning July, 1989 entire water column samples from 10 depths were collected from stations XDA1177, XED4892, PXT0402, MET5.1, MCB1.1, MCB2.2, MCB2.1, XEA1840, XED9490, and MET5.0 (when sampled).

- SAMPLE PRESERVATIVES: Between August, 1984 and September, 1985, 1 ml of neosynephrine was added to each concentrated sample. The sample was allowed to set for 30 minutes and then buffered formaldehyde was added. The neosynephrine step was eliminated after this time and buffered formaldehyde was added to each sample jar prior to the addition of the sample (final concentration of fixative was approximately 2.5%).

-SAMPLE STORAGE ENVIRONMENT: Laboratory

-TIME IN STORAGE: Indefinite

-LAB TECHNIQUES WITH REFERENCES:

Standard Methods

>DATA ENTRY METHOD: Computer keyboard to disk from data sheets for sampling trips prior to April 1985, and keyboard to disk for trips from April 1985 through December 1994. Starting in January 1995, computer keyboard to inhouse computer network.

>DATA VERIFICATION: Visual inspection and computer verification program.

# VARIABLE NAMES, SAS LABELS, AND DESCRIPTIONS

where yy is yy of data set

>ANSMZyyH.SD2 - common sample information maybe merged with counts using SER\_NUM

----LIST OF VARIABLES AND ATTRIBUTES BY POSITION----

VARIABLE	TYPE	LENGTH	POSITION	FORMAT	LABEL
TRIB_COD	CHAR	3	0		SPECIFIES TRIB. SAMPLED (OR MAIN BAY)
SER_NUM	NUM	8	3		SERIAL NUM. INCL. SAMPLING EFFORT NUM.
STATION	CHAR	7	11		STATION NAME
DATE	NUM	8	18		DATE7. SAMPLE DATE
TIME	NUM	8	26		HHMM5. SAMPLE TIME
GMETHOD	CHAR	1	34		GEAR METHOD CODE
TDEPTH	NUM	8	35		TOTAL DEPTH
PDEPTH	NUM	8	43		DEPTH INCLUDED IN SURFACE COMPOSITE
VOLPDEPT	NUM	8	51		VOLUME OF SAMPLE PER DEPTH
TOTVCOMP	NUM	8	59		TOTAL VOLUME OF COMPOSITE SAMPLE

A_DEPTH1 NUM	8	67	1st SAMPLE DEPTH ABOVE PDEPTH
A_DEPTH2 NUM	8	75	2nd SAMPLE DEPTH ABOVE PDEPTH
A_DEPTH3 NUM	8	83	3rd SAMPLE DEPTH ABOVE PDEPTH
A_DEPTH4 NUM	8	91	4th SAMPLE DEPTH ABOVE PDEPTH
A_DEPTH5 NUM	8	99	5th SAMPLE DEPTH ABOVE PDEPTH
B_DEPTH1 NUM	8	107	1st SAMPLE DEPTH BELOW PDEPTH
B_DEPTH2 NUM	8	115	2nd SAMPLE DEPTH BELOW PDEPTH
B_DEPTH3 NUM	8	123	3rd SAMPLE DEPTH BELOW PDEPTH
B_DEPTH4 NUM	8	131	4th SAMPLE DEPTH BELOW PDEPTH
B_DEPTH5 NUM	8	139	5th SAMPLE DEPTH BELOW PDEPTH
FC_INIT CHAR	3	147	INITIALS OF FIELD COLLECTOR
LI_INIT CHAR	3	150	INITIALS OF INDIVID. PERFORMING LAB ID.

> ANSMZyyD.SD2 - counts data set

----LIST OF VARIABLES AND ATTRIBUTES BY POSITION----

VARIABLE	TYPE	LENGTH	POSITION	FORMAT	LABEL
SER_NUM	NUM	8	0		SERIAL NUM. INCL. SAMPLING EFFORT NUM.
STATION	CHAR	7	8		STATION NAME
DATE	NUM	8	15		DATE7. SAMPLE DATE
TRIB_COD	CHAR	3	23		SPECIFIES TRIB. SAMPLED (OR MAIN BAY)
LAYER	CHAR	2	26		AP=ABOVE PYC,BP=BELOW PYC,WC=WATER COL
REPNUM	CHAR	1	28		REPLICATE NUMBER
CONCENT	NUM	8	29		VOLUME OF CONCENTRATED SAMPLE
MLSCNT	NUM	8	37		MILLILITERS COUNTED
SPECCODE	CHAR	13	45		SPECIES CODE
RAWCT	NUM	8	58		RAW COUNT
DENSITYV	NUM	8	66		DENSITY (# INDIV./L) OF GIVEN TAXON
TDENSITY	NUM	8	74		DENSITY (# INDIV./L) OF ALL ORGANISMS

# PHYSICOCHEMICAL VARIABLE NAMES, MEASUREMENT UNITS, AND DESCRIPTIONS

None

# BIOLOGICAL ENUMERATION TECHNIQUES: Each sample was gently mixed and a 1 ml aliquot was removed with a Stempel pipet and put into a Sedgewick-Rafter cell for enumeration with a compound microscope at 100X magnification. Beginning with samples collected in April, 1986, a small drop of concentrated Rose Bengal stain was added to the cell prior to addition of the sample. The sample was allowed to set for 10 minutes before counting. At least one chamber (1 ml) was counted for each sample and if the total count did not reach 250 organisms, subsequent 1 ml aliquots were also enumerated until a count of 250 or more organisms was obtained or 3 ml were examined. If a certain organism was abundant (more than 60 per chamber), it was not counted in the subsequent 1 ml aliquots for a given sample. For extremely abundant taxa, less than one ml could be counted. Species identification was made using the NODC species code. Microzooplankton smaller than 44 um were noted on the original data sheet but not enumerated since estimates would not be quantitative.

#FORMULAS, CALCULATIONS, AND CONVERSIONS:

The following equation was used to convert raw counts to density (#/L) for each taxon identified:  
$$\text{DENSITYV} = ((\text{RAWCT}/\text{MLSCNT}) * \text{CONCENT}) / \text{TOTVCOMP}$$

If the sample was counted by rows MLSCNT is determined by dividing the number of rows by 28.4.

#### # SPECIES CODE USED

NODC CODE WITH INHOUSE MODIFICATIONS (13th column) AND  
SCIENTIFIC NAME:

#### DIGIT REPRESENTS

-----

- 1-2 Phylum
- 3-4 Class and Order
- 5-6 Family
- 7-8 Genus
- 9-10 Species
- 11-12 Subspecies
- 13 Special inhouse modifier

#### # SPECIES INHOUSE CODE, REFERENCE CODE, AND SCIENTIFIC NAME

INHOUSE CODE IS SAME AS REFERENCE CODE WITH THE ADDITION OF AN INHOUSE MODIFIER IN COLUMN 13. THE MODIFIER IN THE 13TH COLUMN HAS DIFFERENT MEANINGS DEPENDING ON THE TAXA BEING CONSIDERED. FOR LARGER METAZOA, A ONE IN THIS COLUMN INDICATES THAT THE ORGANISM IS IN ITS LARVAL OR NAUPLIAR STAGE. FOR SMALLER METAZOA SUCH AS ROTIFERS AND FOR NON-TINTINNINE CILIATE PROTOZOA, THIS MODIFIER MAY INDICATE A SIZE CATEGORY WITHIN A TAXA. THE NODC CODE DOES NOT DISTINGUISH BETWEEN LIFE HISTORY STAGES. FOR TINTINNINE CILIATES (3540...), A 0, 1, 2 (OR 3, 4, 5)

IN THE 13TH COLUMN INDICATES THAT IT WAS NOT OR COULD NOT BE DETERMINED IF THE LORICA CONTAINED A CELL, THAT THE LORICA DID CONTAIN A CELL (FULL), OR THAT THE LORICA DID NOT CONTAIN A CELL (EMPTY), RESPECTIVELY. FOR AN UNDESCRIBED SPECIES, THE CODE IS GIVEN DOWN TO THE LOWEST POSSIBLE TAXONOMIC LEVEL, THE UNKNOWN LEVELS ARE GIVEN ZEROS AND A NUMBER IS ASSIGNED IN THE SUBSPECIES COLUMNS (11-12) (E.G. BRACHIONUS A = 4506010400010; ROTIFER A = 45000000010). THESE NUMBERS WILL BE COMPLETED WHEN MORE DETAILED TAXONOMIC INFORMATION IS OBTAINED.

ONLY THE ORGANISMS GREATER THAN 44 AND LESS THAN 200 UM IN SMALLEST DIMENSION ARE INCLUDED IN THE SPECIES LIST. THE MICROZOOPLANKTON LESS THAN 44 UM ARE NOTED AND RECORDED AS EITHER RARE, COMMON, ABUNDANT, OR DOMINANT. THE MACROZOOPLANKTON ARE ENUMERATED AND RECORDED SEPARATELY. HARD COPIES AND COPIES ON DISKETTES OF THESE DATA AS WELL AS THE SPECIES LISTS FOR THESE

GROUPS ARE AVAILABLE FROM S.G. BROWNLEE AT THE BENEDICT LAB.

CODE	SPECIES
0000000000001	UNIDED LARVAE
0000000000002	UNIDED TROCHOPHORE LARVAE
3438000000000	SARCODINA-UNIDED SARCODINID
3442010000000	DIFFLUGIIDAE
3442010100000	DIFFLUGIA SPP.
3442010201000	LESQUEREUSIA GIBBOSA
3442020100000	ARCELLA SP.
3442030000000	CENTROPYXIDAE
3442030100000	CENTROPYXIS SP.
3442030101000	CENTROPYXIS ACULEATA
3442040000000	PARAQUADRULIDAE
3442040200000	QUADRULELLA SP.
3442050000000	HYALOSPHEINIIDAE-TESTATED AMOEBA
3445020000000	EUGLYPHA SP.
3445040000000	CYPHODERIIDAE
3445040100000	CYPHODERIA SP.
3448000000000	FORAMINIFERIDA
3512000000000	CILIOPHORA-UNIDED CILIATE
3516000000000	HAPTORIDA
3516010100000	DIDINIUM SP.
3516010202000	MESODINIUM RUBRUM
3516020000001	TRACHELOCERCIDAE-LARGE
3517000000001	CYRTOPHORIDA-LARGE
3530000000001	PERITRICHIDA-LARGE
3531000000000	SESSILINA-UNIDED SESSILINE PERITRICH
3532000000000	MOBILINA-UNIDED MOBILINE PERITRICH
3533000000000	SUCTORIA-UNIDED SUCTORIAN
3534010100000	ACINETA SP.
3534030700000	STAUROPHRYA SP.
3534040100000	EPHELOTA SP.
3537000000000	HETEROTRICHINA-UNIDED HETEROTRICH
3539000000000	OLIGOTRICHINA-UNIDED OLIGOTRICH
3539020100000	STENTOR SP.
3539030000000	STROMBIDIDAE
3539030100000	STROMBIDIUM SP.
3539030200000	TONTONIA SP.
3540000000000	TINTINNINA-UNIDED TINTINNID
3540010100050	TINTINNIDIUM SP.-LARGE
3540010100051	TINTINNIDIUM SP.-LARGE-FULL
3540010100052	TINTINNIDIUM SP.-LARGE-EMPTY

3540020100000 TINTINNOPSIS SP.  
3540020100001 TINTINNOPSIS SP.-FULL  
3540020100002 TINTINNOPSIS SP.-EMPTY  
3540020100030 TINTINNOPSIS SUBACUTA-HUGE  
3540020100031 TINTINNOPSIS SUBACUTA-HUGE-FULL  
3540020100032 TINTINNOPSIS SUBACUTA-HUGE-EMPTY  
3540020100050 TINTINNOPSIS SP. A  
3540020100051 TINTINNOPSIS SP. A-FULL  
3540020100052 TINTINNOPSIS SP. A-EMPTY  
3540020105000 TINTINNOPSIS DADAYI  
3540020105001 TINTINNOPSIS DADAYI-FULL  
3540020105002 TINTINNOPSIS DADAYI-EMPTY  
3540020123000 TINTINNOPSIS FIMBRIATA  
3540020123001 TINTINNOPSIS FIMBRIATA-FULL  
3540020123002 TINTINNOPSIS FIMBRIATA-EMPTY  
3540020123003 TINTINNOPSIS FIMBRIATA-MEUNIERI GRP  
3540020123004 TINTINNOPSIS FIMBRIATA-MEUNIERI GRP-FULL  
3540020123005 TINTINNOPSIS FIMBRIATA-MEUNIERI GRP-EMPTY  
3540020129000 TINTINNOPSIS RADIX  
3540020129001 TINTINNOPSIS RADIX-FULL  
3540020129002 TINTINNOPSIS RADIX-EMPTY  
3540020133000 TINTINNOPSIS SUBACUTA  
3540020133001 TINTINNOPSIS SUBACUTA-FULL  
3540020133002 TINTINNOPSIS SUBACUTA-EMPTY  
3540020136000 TINTINNOPSIS MEUNIERI  
3540020136001 TINTINNOPSIS MEUNIERI-FULL  
3540020136002 TINTINNOPSIS MEUNIERI-EMPTY  
3540020137000 TINTINNOPSIS KARAJACENSIS  
3540020137001 TINTINNOPSIS KARAJACENSIS-FULL  
3540020137002 TINTINNOPSIS KARAJACENSIS-EMPTY  
3540020138000 TINTINNOPSIS NITIDA  
3540020138001 TINTINNOPSIS NITIDA-FULL  
3540020138002 TINTINNOPSIS NITIDA-EMPTY  
3540030100010 STENOSEMELLA SP.A  
3540030100011 STENOSEMELLA SP.A-FULL  
3540030100012 STENOSEMELLA SP.A-EMPTY  
3540050100020 METACYLIS SP. B  
3540050100021 METACYLIS SP. B-FULL  
3540050100022 METACYLIS SP. B-EMPTY  
3540050100030 METACYLIS SP. C  
3540050100031 METACYLIS SP. C-FULL  
3540050100032 METACYLIS SP. C-EMPTY  
3540050400000 CLIMACOCYLIS SP.  
3540050400001 CLIMACOCYLIS SP.-FULL  
3540050400002 CLIMACOCYLIS SP.-EMPTY

3540050501000 STYLICAUDA PLATENSIS  
3540050501001 STYLICAUDA PLATENSIS-FULL  
3540050501002 STYLICAUDA PLATENSIS-EMPTY  
3540070100000 FAVELLA SP.  
3540070100001 FAVELLA SP.-FULL  
3540070100002 FAVELLA SP.-EMPTY  
3540130100010 EUTINTINNUS SP.A  
3540130100011 EUTINTINNUS SP.A-FULL  
3540130100012 EUTINTINNUS SP.A-EMPTY  
3543000000000 HYPOTRICHIDA  
3545010000000 EUPLOTIDAE  
3545010100001 EUPLOTES SPP.-LARGE  
3545010100020 EUPLOTES SP. A  
3545010100030 EUPLOTES SP. B  
4400000000000 GASTROTRICHA  
4500000000000 ROTIFERA-UNIDED ROTIFER  
4500000000010 ROTIFER A  
4500000000020 ROTIFER B  
4500000000030 ROTIFER C  
4500000000040 ROTIFER D  
4500000000050 ROTIFER E  
4500000000060 ROTIFER F  
4504000000000 BDELLOIDA-UNIDED BDELLOID ROTIFER  
4504020100000 ROTARIA SP.  
4504020103000 ROTARIA CITRINUS  
4504020104000 ROTARIA NEPTUNIA  
4504020300000 MACROTRACHELA SP.  
4506010100000 KERATELLA SP.  
4506010100010 KERATELLA SP. A  
4506010102000 KERATELLA QUADRATA  
4506010103000 KERATELLA COCHLEARIS  
4506010103020 KERATELLA COCHLEARIS COCHLEARIS  
4506010103030 KERATELLA COCHLEARIS HISPIDA  
4506010103040 KERATELLA COCHLEARIS MICRACANTHA  
4506010103050 KERATELLA COCHLEARIS ROBUSTA  
4506010103060 KERATELLA COCHLEARIS TECTA  
4506010104000 KERATELLA CRASSA  
4506010105000 KERATELLA EARLINAE  
4506010106000 KERATELLA VALGA  
4506010200000 NOTHOLCA SP.  
4506010203000 NOTHOLCA ACUMINATA  
4506010300000 COLURELLA SP.  
4506010400000 BRACHIONUS SP.  
4506010400010 BRACHIONUS SP. A  
4506010400020 BRACHIONUS SP. B

4506010401000 BRACHIONUS PLICATILIS  
4506010402000 BRACHIONUS CALYCIFLORUS  
4506010403000 BRACHIONUS HAVANAENSIS  
4506010404000 BRACHIONUS PTERODINOIDES  
4506010405000 BRACHIONUS URCEOLARIS  
4506010406000 BRACHIONUS ANGULARIS  
4506010407000 BRACHIONUS BIDENTATA  
4506010408000 BRACHIONUS BUDAPESTINENSIS  
4506010409000 BRACHIONUS CAUDATUS  
4506010410000 BRACHIONUS DIVERSICORNIS  
4506010411000 BRACHIONUS QUADRIDENTATUS  
4506010412000 BRACHIONUS RUBENS  
4506010413000 BRACHIONUS VARIABILIS  
4506010500000 KELLICOTTIA SP.  
4506010501000 KELLICOTTIA LONGISPINA  
4506010502000 KELLICOTTIA BOSTONIENSIS  
4506010700000 LEPADELLA SP.  
4506010704000 LEPADELLA PATELLA  
4506010800000 ANURAEOPSIS SP.  
4506010801000 ANURAEOPSIS FISSA  
4506010900000 EPIPHANES SP.  
4506011000000 EUCHLANIS SP.  
4506011001000 EUCHLANIS DILATATA  
4506011100000 LOPHOCHARIS SP.  
4506011101000 LOPHOCHARIS SALPINA  
4506011200000 MACROCHAETUS SP.  
4506011300000 MYTILINA SP.  
4506011400000 PLATYIAS SP.  
4506011401000 PLATYIAS PATULUS  
4506011402000 PLATYIAS QUADRICORNIS  
4506011500000 TRICHOTRIA SP.  
4506011501000 TRICHOTRIA TETRACTIS  
4506020100000 LECANE SP.  
4506020200000 MONOSTYLA SP.  
4506020201000 MONOSTYLA BULLA  
4506020202000 MONOSTYLA CLOSTROCERCA  
4506020203000 MONOSTYLA QUADRIDENTATA  
4506040100000 ENCENTRUM SP.  
4506040200000 PROALES SP.  
4506040300000 CEPHALODELLA SP.  
4506040302000 CEPHALODELLA GIBBA  
4506040400000 NOTOMMATA SP.  
4506040500000 MONOMMATA SP.  
4506040600000 EOSPHORA SP.  
4506070100000 TRICHOCERCA SP.

4506070102000 TRICHOCERCA CYLINDRICA  
4506070103000 TRICHOCERCA LONGISETA  
4506070104000 TRICHOCERCA MULTICRINIS  
4506070105000 TRICHOCERCA SIMILIS  
4506080100000 ASCOMORPHA SP.  
4506080101000 ASCOMORPHA OVALIS  
4506080102000 ASCOMORPHA SALTANS  
4506080200000 GASTROPUS SP.  
4506080201000 GASTROPUS MINOR  
4506120100000 ASPLANCHNA SP.  
4506120101000 ASPLANCHNA BRIGHTWELLI  
4506120102000 ASPLANCHNA HERRICKI  
4506120103000 ASPLANCHNA PRIODONTA  
4506130200000 SYNCHAETA SP.  
4506130200001 SYNCHAETA SPP. L-LARGE  
4506130200002 SYNCHAETA SPP. M-MEDIUM  
4506130200003 SYNCHAETA SPP. S-SMALL  
4506130200010 SYNCHAETA BICORNIS  
4506130200020 SYNCHAETA BALTICA  
4506130204000 SYNCHAETA PECTINATA  
4506130206000 SYNCHAETA OBLONGA  
4506130207000 SYNCHAETA STYLATA  
4506130300000 POLYARTHRA SP.  
4506130302000 POLYARTHRA DISSIMULANS  
4506130303000 POLYARTHRA DOLICHOPTERA  
4506130304000 POLYARTHRA EURYPTERA  
4506130305000 POLYARTHRA MAJOR  
4506130306000 POLYARTHRA REMATA  
4506130307000 POLYARTHRA VULGARIS  
4506130400000 PLOESOMA SP.  
4506130401000 PLOESOMA HUDSONI  
4506130402000 PLOESOMA TRUNCATUM  
4507010100000 TESTUDINELLA SP.  
4507010101000 TESTUDINELLA PATINA  
4507020100000 HEXARTHRA SP.  
4507020101000 HEXARTHRA MIRA  
4507040100000 CONOCHILOIDES SP.  
4507040102000 CONOCHILOIDES DOSSUARIUS  
4507040103000 CONOCHILOIDES NATANS  
4507040200000 CONOCHILUS SP.  
4507040201000 CONOCHILUS HIPPOCREPIS  
4507040202000 CONOCHILUS UNICORNIS  
4507050100000 FILINIA SP.  
4507050101000 FILINIA LONGISETA  
4507050102000 FILINIA BRACHIATA



4507050103000 FILINIA TERMINALIS  
 4508010100000 COLLOTHECA SP.  
 4508010101000 COLLOTHECA MUTABILIS  
 4508010102000 COLLOTHECA PELAGICA  
 4700000000000 NEMATODA  
 5100000000001 GASTROPODA-LARVAE  
 5500000000001 PELECYPODA-LARVAE  
 5515370301000 DREISSENA POLYMORPHA  
 5922000000000 ACARINA-MITE  
 6117000000001 COPEPOD NAUPLII  
 6117000000005 COPEPOD NAUPLII+PERITRICHS  
 6118180100001 DIAPTOMUS -NAUPLII  
 6118190200001 PSEUDODIAPTOMUS SP.-NAUPLII  
 6118200200001 EURYTEMORA SP.-NAUPLII  
 6118290100001 ACARTIA SP.-NAUPLII  
 6119050200001 SCOTTOLANA SP.-NAUPLII  
 6120080200001 CYCLOPS SP.-NAUPLII  
 6120080300001 MESOCYCLOPS SP.-NAUPLII  
 6120090100001 OITHONA SP.-NAUPLII  
 6120260100001 HEMICYCLOPS SP.-NAUPLII  
 7500000000000 TARTIGRADA

THE ABOVE SPECIES LIST HAS BEEN INCLUDED ON THE DATA TAPE AS FILE MMZP.SD2.

4/15/92 - THE NAME OF SPECIES #4506130200010 HAS CHANGED FROM SYNCHAETA SP. A-LONG HORNS TO SYNCHAETA BICORNIS AND FOR SPECIES #4506130200020, THE NAME HAS CHANGED FROM SYNCHAETA SP. B-'PORKER' TO SYNCHAETA BALTICA.

# NUMERICAL VARIABLE NAMES - WARNING AND ERROR BOUNDS

VARIABLE	VALID RANGE
-----	-----
DATE	840802-971218
TIME	0651-1935
TDEPTH	1.8-33
PDEPTH	0.5-22.0
VOLPDEPT	2-20
TOTVCOMP	12-200
SER_NUM	01001-240039
A_DEPTH1	0.2- 1.0
A_DEPTH2	0.4- 5.0
A_DEPTH3	0.5-10.0

A_DEPTH4	0.5-15.0
A_DEPTH5	0.5-21.5
B_DEPTH1	1-22.5
B_DEPTH2	1-25.0
B_DEPTH3	1-27.0
B_DEPTH4	1-29.0
B_DEPTH5	1-32.0
CONCENT	10-500
MLSCNT	0.01-10
RAWCT	1-9999
DENSITYV	0.05-10000.00
TDENSITY	1-999999.99

# CHARACTER VARIABLES - VALID VALUES

VARIABLE	VALID VALUES
-----	-----

REPNUM

1  
2  
3  
4  
T  
B  
W

TRIB\_COD

BAY  
PAX  
POT  
PAT  
CHS  
CHP

SPECCODE 000000000001-750000000000

STATION

MCB1.1  
MCB2.1  
MCB2.2  
MCB3.3C  
MCB4.3C  
MCB5.2  
XDE5339  
XED4892  
XED9490  
PXT0402

XEA6596  
XEA1840  
XDA1177  
MLE2.2  
MET4.2  
MET5.1  
MET5.0  
MET5.2  
MEE3.1  
MWT5.1

GMETHOD P

FC\_INIT

CVW  
REJ  
RVL  
KGS  
SGB  
SSH  
CRP  
KRB  
DCB  
CF  
MHB  
ABT  
BBW  
JSH  
ALI  
DET  
PHD  
DCN  
AMH  
KAM  
KMB  
ALM  
RMJ  
SGS  
KAR  
SGS  
JLG

LI\_INIT

BBW  
REJ  
SGB  
SSH  
JHS

KAM  
KMB  
RMJ  
KAR  
SGS  
JLG

# KEY WORDS (EXCLUDING VARIABLE NAMES)

# INITIALS OF SCIENTISTS IN DATA SET

CVW Carolyn V. Watson  
REJ Richard E. Jacobsen  
RVL Richard V. Lacouture  
KGS Kevin G. Sellner  
SGB Stella G. Brownlee  
SGS Stella G. Sellner (Brownlee)  
SH Sharyn S. Hedrick  
SSH Sharyn S. Hedrick  
CRP Charles R. Parrish  
KB Kevin R. Braun  
KRB Kevin R. Braun  
DCB David C. Brownlee  
CF Chris Frye  
MHB Marie H. Bundy  
ABT Allison B. Tate  
BBW Bruce B. Wagoner  
JSH Jeffrey S. Handen  
ALI Amy L. Imirie  
PHD Paul H. DiNunno  
DET Douglas E. Talaber  
DCN Donna C. Nicholson  
AMH Ann Marie Hartsig  
JHS James H. Sniezek  
KAM Kimberly A. Morcom  
KMB Kimberly Morcom Burke  
ALM Andrea L. Morcom  
RMJ Ralph Matos, Jr.  
KAR Karen A. Rota  
JLG Jennifer L. Gronefeld