

DYNAMIC SYSTEMS ANALYSIS OF FOSSIL DINOFLAGELLATES  
FROM THE ATLANTIC COASTAL PLAIN, USA.

by

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# **DYNAMIC SYSTEMS ANALYSIS OF FOSSIL DINOFLAGELLATES FROM THE ATLANTIC COASTAL PLAIN, USA.**

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Geological Sciences

## **ABSTRACT**

Dynamic Systems modeling suggests that complex coastal dinoflagellate bio-systems can be modeled using environmental parameters such as temperature, salinity, and bulk nutrient levels. The former Salisbury Embayment of northern Virginia and Maryland is modeled here, using STELLA II and FORTRAN models based on physical oceanography and temperature, salinity, and nutrient conditions of the modern Yellow Sea. In these models, dinoflagellate assemblages are predicted based on environmental conditions associated with depth.

Cluster analyses of fossil dinoflagellate frequency data from Tertiary Pamunky Group (Aquia and Nanjemoy Formations) of the Salisbury Embayment produce 17 discrete groupings. Samples within the Salisbury fossil cluster groups are statistically similar (via ANOVA analysis), but not the same. Therefore they represent paleocommunity types rather than paleocommunities. Although individual dinoflagellate species recur in similar environmental settings, the paleocommunity types do not appear to repeat.

In the past, such associations have been used as depth indicators. It is suggested here that they relate to estuarine, nearshore, and offshore coastal regions because of the temperature, salinity, and nutrient conditions of each.

In the modern Yellow Sea, nearshore and offshore regions are separated by discrete lateral fronts in some areas, and by gradational regions of mixing in others. Both types of watermass boundaries are modeled in this study. Results suggest that evidences that discrete watermass boundaries might have occurred between some fossil dinoflagellate associations. Circulation patterns of the Salisbury Embayment may have been roughly similar to those of the modern Yellow Sea.

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## **INTRODUCTION**

### **General Comments**

Fossil marine dinoflagellates (microscopic phytoplankton belonging to the Division Pyrrophyta) are used extensively for age dating, correlation, and paleoenvironmental analysis. However, they have never been studied via system dynamics analyses of their community structure. System dynamics are used routinely in many fields of science to study complex multivariate systems (Roberts, 1983; Rose, 1987). Analyses of living systems and their environment involve complex interactions (May, 1981; May, 1993). This study is the first to use system dynamics to study fossil marine dinoflagellate assemblages.

The Virginia Tech Palynology Program produced numerous dissertations and theses on the taxonomy, morphology, and biostratigraphy of Cretaceous and Tertiary dinoflagellates from the Atlantic Coastal Plain. This current study uses data from previous studies of the Aquia (Paleocene) and Nanjemoy (Eocene) Formations of the Virginia Coastal Plain (McLean, 1969; McLean, 1972; Witmer, 1975; Goodman, 1975; and Witmer, 1987). The Aquia and Nanjemoy Formations were deposited in an ancient embayment known as the Salisbury Embayment located geographically in the region of the present Chesapeake Bay and Atlantic Coastal Plain in the states of Virginia, Maryland, and Delaware (Figure 1). The localities studied herein are from Virginia, in the central and southern parts of the Salisbury Embayment.

### **Objectives of this Study**

The primary objective of this study is to examine fossil dinoflagellate frequency data from the Salisbury Embayment for paleocommunity structure and environmental association using system dynamics and paleocommunity analysis. A second objective of this study is to test the potential of using system dynamics FORTRAN modeling in describing coastal plankton systems.

### **Methodology**

Data from the Tertiary Pamunky group was used in this project for three main reasons. This stratigraphic interval provides a relatively continuous record of changing local environmental conditions (a regression followed by a transgression). The depositional record for this interval is assumed to be relatively constant (as opposed to K/T boundary sediment below, or upper Eocene possibly meteor impact-disturbed sediment above (Ward, 1993)). In addition, significant fossil dinoflagellate frequency data exist for this interval.

The frequency data used are derived from previous taxonomic studies of this interval.<sup>2</sup>

Environmental association is explored by modeling paleoenvironmental conditions of the Salisbury Embayment during the Tertiary based on modern analogs of the Yellow Sea and the Gulf and Atlantic Seaboard. The Yellow Sea was chosen as the primary physical analog because of its similarity in size, extent, geographical orientation, and possibly similar circulation patterns to the Salisbury Embayment. The modern analogs are used to construct both dynamic systems STELLA II and FORTRAN computer models, based on biological and environmental information, by which possible dinoflagellate environments and interactions may be modeled. STELLA II is a model construction program which uses an iterative approach to exploring dynamic processes. STELLA II is used to design and test structural diagrams (see Richmond, et al., 1987). FORTRAN modelling, however is more mathematically rigorous. Data sets for both modeling approaches were manipulated using EXCEL spreadsheets (See Cobb, 1985).

Potential paleocommunity structure is tested for using Q-mode cluster analysis and analysis of variance techniques. Cluster analysis and comparative sorting and graphing techniques are used to delineate separate species groups based on similarity coefficients.

Groups produced by cluster analysis appear preferentially associated with particular coastal regions and their physical conditions.

## **I. GEOGRAPHICAL AND GEOLOGICAL SETTING**

### **The Salisbury Embayment**

The Atlantic Coastal Plain is made up of several ancient embayments separated from one another by tectonic arches (Figure 1). The Salisbury Embayment (Figure 2) was one of these ancient embayments. It existed mostly during Tertiary time, covering areas of what are now Virginia, Maryland, and Delaware (Glaser, 1968; Gallagher, 1984; Gibson, 1989). The modern Chesapeake Bay, centered over the old Salisbury Embayment is several times smaller than its ancient counterpart.

The Salisbury Embayment was continuously connected to the open Atlantic Ocean. Water level in the Salisbury was a function of tectonic and eustatic sea level changes. At its maximum, it covered up to 20 thousand km<sup>2</sup> at depths up to 150 meters in its eastern deepest part. Sediments deposited in the Salisbury Embayment indicate sediment-starved (glauconite-rich) nearshore marine to brackish-water conditions (Ward and Krafft, 1985).

The Norfolk Arch provided a primary structural control for the Salisbury Embayment during this time by affecting the shape and orientation of the embayment (Figure 2). Aquia and Nanjemoy sediments contain mica flakes in sands and coarse material which match with Roanoke River group sediments, derived primarily from the Virginia Piedmont (Ward and Krafft, 1985). The Piedmont region lies immediately southwest of the embayment. The Norfolk Arch structure to the south diverted freshwater Piedmont drainage northward and into the embayment. Also Norfolk Arch controlled both access from the open sea and circulation within the embayment. Throughout the Paleocene and early Eocene, normal marine flow entered the embayment primarily as bottom waters but was controlled by arch and bottom topography. The wide mouth of the embayment may have been protected to varying degree by banks or bars, and/or the topographically high remains of a Jurassic aged reef complex just to the northeast of the embayment mouth (Poag, 1993). Topographic submarine highs may have acted as partial sediment dams, keeping sand and fine clay within the embayment.

The presence of glauconite in the sediments and marine macro-fossils, particularly bivalves (Ward and Krafft, 1985; Edwards, 1989) indicate marine conditions. Glauconite, along with phosphate pellets and local carbonate, may reflect sediment starvation, and the

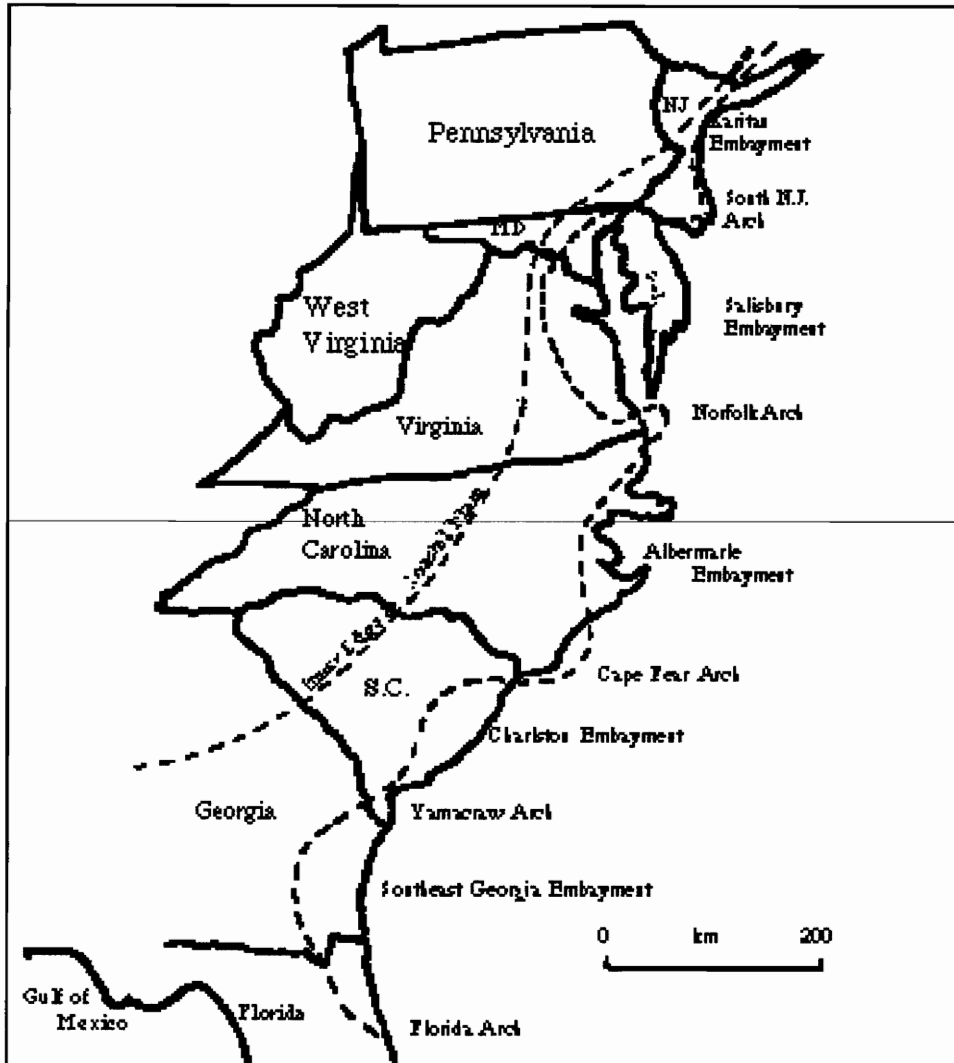


Figure 1. East Coast of US. showing Arches and Embayments.

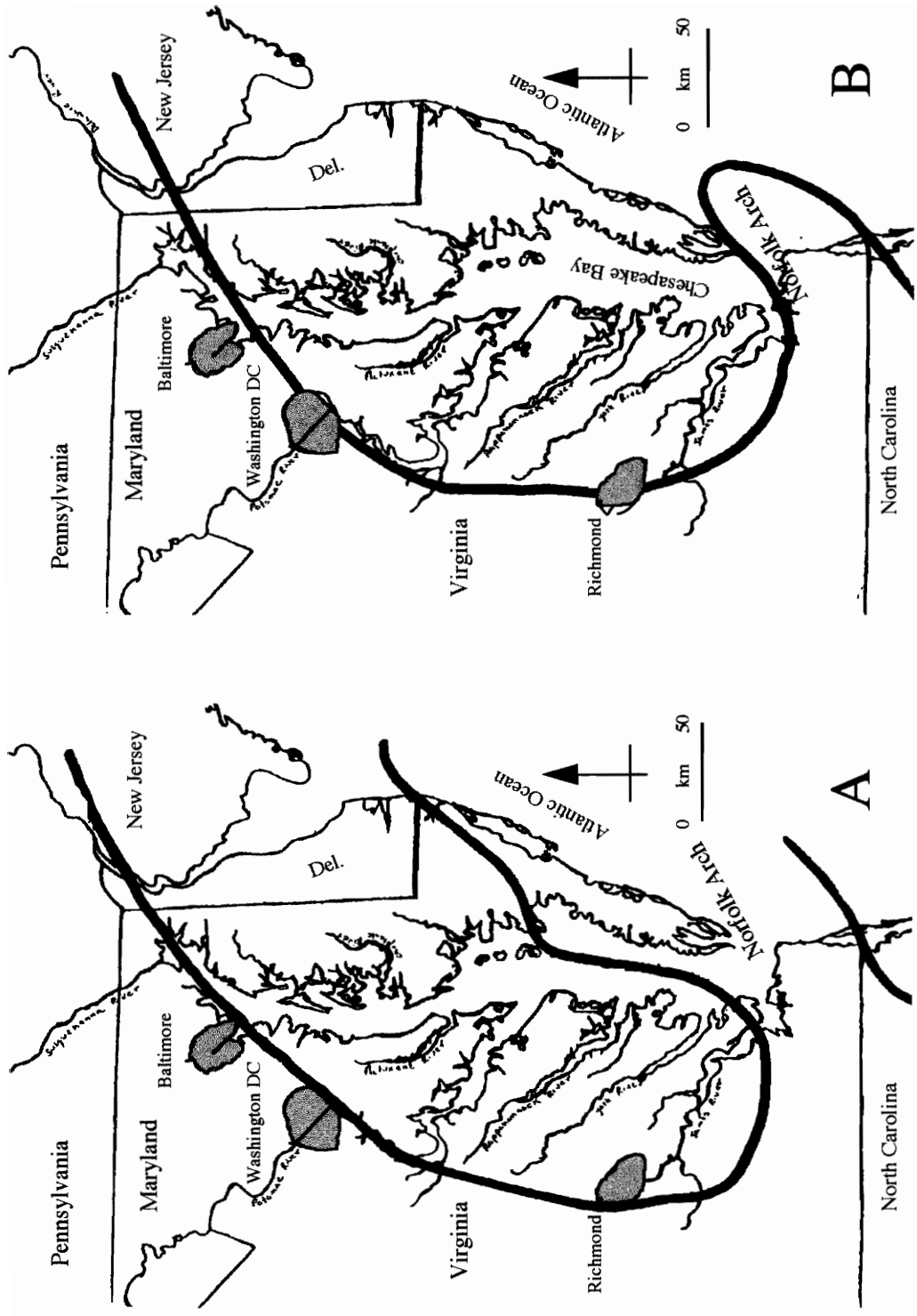


Figure 2. Approximate areal extent of the Salisbury Embayment during A) Aquia Deposition, B) Nanjemoy Deposition (after Ward and Krafft, 1985.)

presence of fine terrestrial clays in contact with warm marine water (Mallinson and Lee, 1986). Glauconite has been associated with local marine flooding events (Tenison, 1989). Percentages of glauconite have been used to estimate depth and/or amount of marine influence within the Salisbury Embayment (Porrenga, 1967).

Salisbury Embayment sediments represent a series of unconformity bounded cyclic pulses of sediment during the times of transgression (Baum, 1986; Harris, 1993). Ward and Krafft (1985) note that, "the Atlantic basins...were characterized by relatively thin deposits, principally marine, with only remnants of nearshore facies. It is clear that large volumes of sediment were not being transported to the Atlantic Coastal Plain during the Tertiary."

Samples for this study are from the marine Pamunkey Group (Paleocene and Eocene) from the central and southwestern portions of the Salisbury Embayment.

### **The Pamunkey Group**

The Pamunkey Group (Figure 3) is the basal Tertiary unit of the Virginia-Maryland Coastal Plain. It is subdivided into the Aquia (Paleocene) and Nanjemoy (Eocene) Formations (Figure 3).

The Pamunkey Group was described and named as a formation by Darton (1891). Clark (1895) subdivided the Pamunkey Group into Aquia Creek and Woodstock "Stages." Clark and Martin (1901) divided the group into the Aquia and Nanjemoy Formations which were each subdivided into two members (Figure 3), and into several finer scale "zones" which were based on lithology and fossils.

The "zones" of Clark and Martin do not correspond with the modern definition of a "zone" as set by the Code of Stratigraphic Nomenclature (1970). The term "unit" will be used in this work (after McLean, 1969).

The Marlboro clay, a meter thick clay unit elevated to formational status by Glaser (1971), separates the Aquia and Nanjemoy Formations.

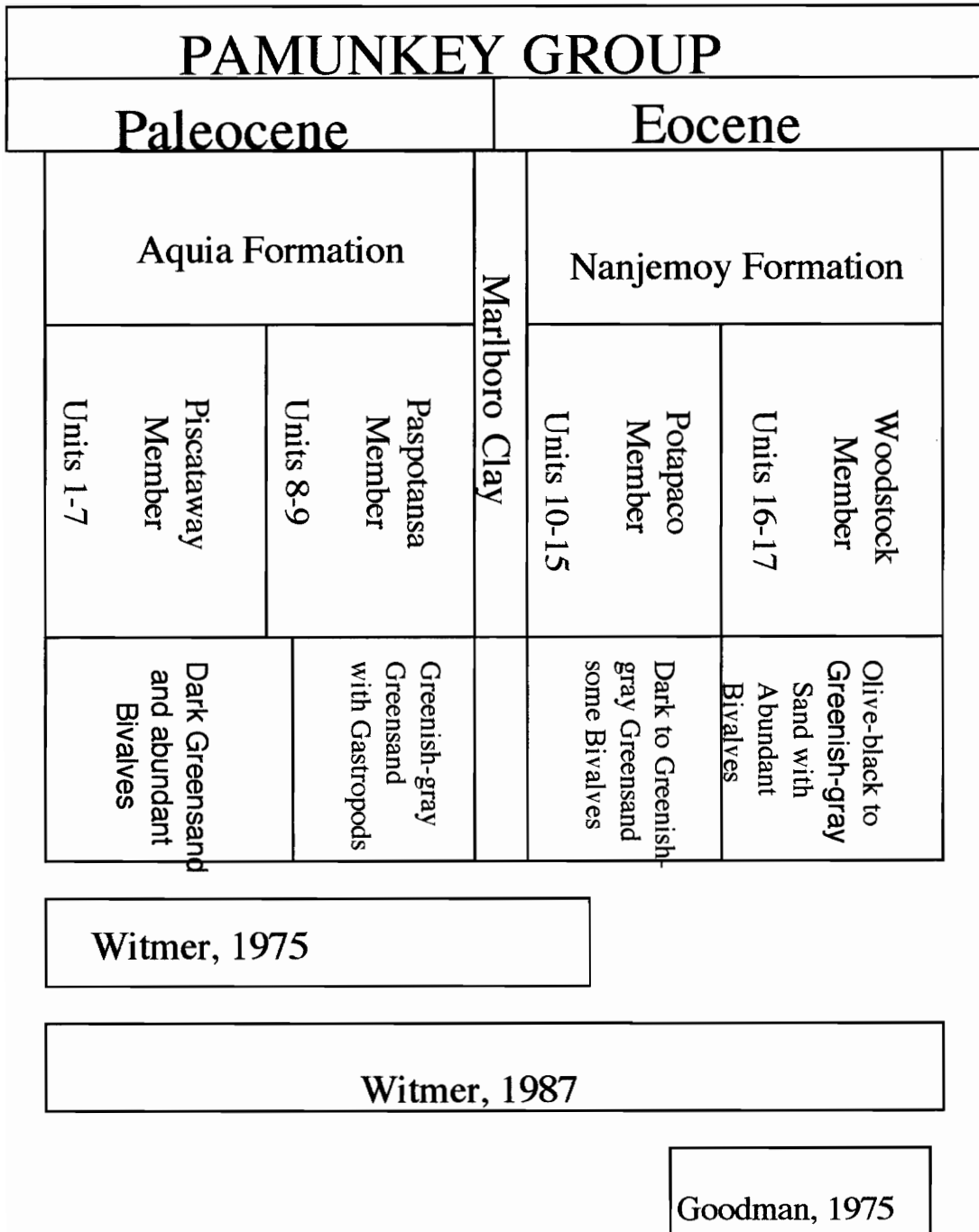


Figure 3. Pamunkey Group geologic column. (After Witmer, 1987)

### **The Aquia Formation**

The Aquia Formation is the basal unit of the Pamunkey Group. It rests unconformably upon Late Cretaceous marine sediments over most of its geographical extent.

The type locality of the Aquia Formation is at the mouth of the Aquia Creek in Stafford County, Virginia. Its thickness ranges from about 30 meters at its type locality to a maximum of 76 meters at its eastern-most extent in Delaware and New Jersey. It consists primarily of glauconitic quartz sands. The Aquia contains fossil remains of tropical reptiles, mammals, and birds including pelican and albatross (Olsen, 1985; Weems, 1985). It is subdivided into the Piscataway (basal) and Paspotansa (overlying) members (Figure 3).

The Piscataway Member is a glauconitic (20 to 70 percent) clay rich and silty quartz sand transgressive marine unit (Ward and Krafft, 1985) deposited at a water depth of about 100 meters (Nogan, 1964). Common macrofossils include large bivalves and occasional *Turritella* gastropods (see Ward and Krafft, 1985, for details). The Piscataway Member contains units 1-7 of Clark and Martin.

The Paspotansa Member consists of glauconitic micaceous silty quartz sands somewhat better sorted than those of the Piscataway Member, and may represent a higher energy depositional environment (Ward and Krafft, 1985). A few beds contain bivalve macrofossils (see Ward and Krafft, 1985, for details).

The Paspotansa also is a transgressive unit deposited under progressively shoaling conditions. It includes units 8 and 9 of Clark and Martin.

### **The Marlboro Formation**

The Marlboro Formation is a roughly meter-thick silty clay unit that separates the Aquia and Nanjemoy Formations. It varies in thickness, and is missing at some localities. It rests unconformably upon the Aquia Formation and is in turn overlain unconformably by the Nanjemoy Formation.

Interpretation of the depositional environment has been controversial. Ward and Krafft interpret the Marlboro as tidal flat deposition (Ward and Krafft, 1985), based partially on ripple marks within siltier portions of the unit. Others have evoked deposition in quiet waters of a protected embayment (Roger Cuffey, personal communication, 1996.)



The Marlboro is not included in the present study because of poor preservation, paucity of dinoflagellates, and reworking (Dewey McLean, personal communication, 1996).

### **The Nanjemoy Formation**

The Nanjemoy sediment is composed of fine glauconitic quartz sands which are often argillaceous and sometimes calcareous with some beds containing abundant gypsum crystals. At the type locality along Nanjemoy Creek, Charles County, Maryland it is about 38 meters thick. The beds thicken to the east and northeast. The Nanjemoy Formation is divided into the Potapaco and overlying Woodstock members.

The Potapaco Member includes units 10-15 of Clark and Martin. Lower portions of the Potapaco consist of black to pink clay-rich sands containing (often detrital) glauconite. Some layers contain gypsum crystals and concretions. Worn and broken small bivalve fossils, scaphopods, *Calianassa* burrows and other bioturbation indicate more energetic conditions, possibly because of shoaling. Upper portions of the member contain higher amounts of glauconite, along with phosphate pebbles and wood fragments. This suggests a transgressive pulse during which sedimentation rates were low (Ward and Krafft, 1985).

The Woodstock Member lies unconformably upon the Potapaco Member. It includes units 16 and 17 of Clark and Martin. The basal Woodstock is marked by a pebble concentration, burrows and wood fragments. The Woodstock consists of very fine, well-sorted, silty glauconitic sands. Glauconite increases upward through the member whereas wood fragments decrease. The Woodstock contains diverse molluscan fossils (Ward and Krafft, 1985). It was deposited during transgression which returned the embayment from relatively nearshore to offshore marine shelf conditions. Woodstock sediments at Popes Creek, Maryland contain fossil fruits of *Wetherellia marylandica*. (Tiffney, 1985). These, and similar tropical mangrove-like fossil fruits from time-equivalent beds in Mississippi (Call, 1993), suggest that tropical climate existed throughout the region of the Salisbury Embayment.

## II. THE DINOFLAGELLATES

R. H. Whittaker divided life into 5 kingdoms: Animalia, Plantae, Monera, Fungi, and Protoctista. The dinoflagellates belong to the latter, and more specifically, to the Division Pyrrophyta (Pascher, 1913) and the Class Dinophyceae (Fritch, 1935). Lee (1992) recognizes six orders of modern Dinophyceae: Prorocentrales, Dinophysiales, Peridinales, Dinocapsales, Dinococcales, and Dinotrichales. Of these, only the Peridinales (Heackel, 1894) are important to this study.

Dinoflagellates are eukaryotes, but because of their nuclear organization, they are best described as mesokaryotic (between prokaryotic and eukaryotic). Most are single celled, but some form colonies. Thousands of species, both marine and freshwater, exist. Most are marine and are most abundant in warm waters. Most are heterotrophic, but some are phagotropic and predatory (Burkholder, 1992; 1995; Steidinger, 1996). This study deals with single celled marine dinoflagellates.

### **Dinoflagellate Life Cycles: Theca Versus Cyst Forms**

Dinoflagellates are microscopic single celled organisms with complex life cycles many of which include both asexual and sexual reproduction, and both motile and resting stages (see figure 4 for generalized life cycle and description). What is known of fossil dinoflagellate life cycles is based on studies of modern dinoflagellates. It is assumed that many fossil forms utilized similar strategies.

Living motile dinoflagellates range in size from 20 to over 200 micrometers. Many species encase the protoplasm within a cellulosic shell or theca. Unfortunately, since cellulose is not well-preserved in the fossil record, our knowledge of fossil dinoflagellates is based on the study of resting cysts.

Cysts are composed of a complex chemically resistant carotenoid-like organic polymer known as sporopollenin, which is preserved in the fossil record. Dinoflagellate cysts often exist in sediments in which calcareous and silicious microfossils have been destroyed. Many cyst forms reflect faithfully the tabulation pattern of its corresponding thecate form. Dinoflagellate thecae are made up of numerous tiny plates, each of which has a unique shape and position in the theca (figure 5). These plates are arranged into series (See figure 5). The overall arrangement of plates is referred to as tabulation. Tabulation is an important criterion in dinoflagellate classification. The resulting pattern provides diagnostic cyst morphotypes by which we can identify modern or fossil cysts. In

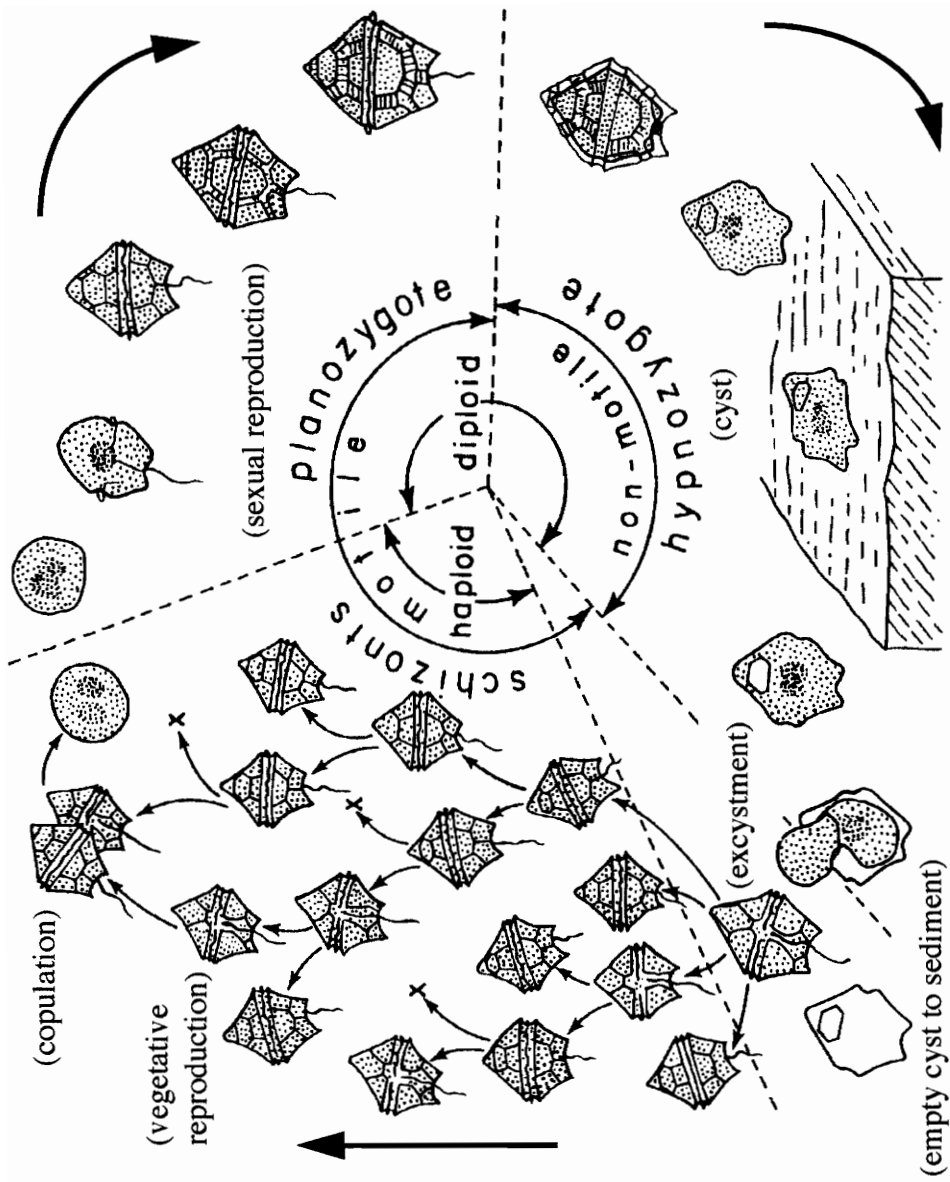


Figure 4. Dinoflagellate Life Cycle. After Evitt, 1985.

accordance with Evitt (1976; 1985), such cyst features are referred to with the prefix para-attached. Thus, cyst features become paraplates, paratabulation, parasulcus, and paracingulum (See figure 4).

There are two primary architectural styles of tabulation/ paratabulation in the peridinialean dinoflagellates. Those whose thecal plate arrangement resembles that of living *Gonyaulax* are placed in the "group" Gonyaulacaceae. Those whose plate pattern approximates that of modern *Peridinium* are referred to "group" Peridiniaceae.

In some cysts the cyst wall is deposited within, and next to the thecal wall. These are known as proximate cysts. Choratae cysts are similar, but have well-developed ornamentation or processes caused by shrinkage of the inner cyst from the inside of the thecal wall. Gonyaulacoid cysts tend to be chorate. In cavate cysts, space exists between layers of the cyst wall, and layers are separated by often intricate perpendicular walls or columns. Peridinioid cysts are often cavate.

Other criteria used for cyst classification include shape, ornamentation such as processes or spines, etc., cyst wall structure, and archeopyle type (opening in the cyst wall(s) through which the protoplasm escapes the cyst). See figure 5.

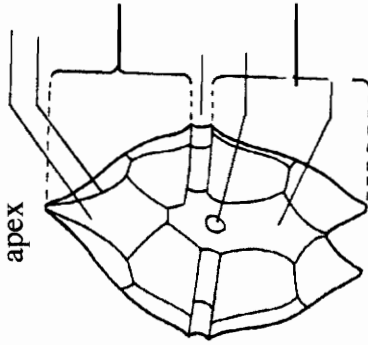
#### FIGURE 5. Dinoflagellate Morphology

The sulcus is a longitudinal furrow located on the ventral portion of the theca. A smooth (acronematic) longitudinal flagellum is attached to the sulcus. It is responsible for both steering and forward movement of the dinoflagellate. The cingulum (or girdle) is an equatorial furrow which divides the theca into upper and lower hemispheres. The ends of the cingulum meet the sulcus, and may vertically offset one another. A transverse flagellum lies coiled helically within the cingulum. This transverse flagellum is usually covered with mastigonemes (small fibrillar hairs). A rhythmic beating of this flagellum causes both forward movement of the dinoflagellate and also a rotational movement of the cell through the water (Thomas, 1995).

The archeopyle is the aperture formed by the loss of either a single paraplate or group of paraplates (plate field) which allows the protoplasm to escape from the cyst during excystment (Evitt, 1976). The archeopyle of a given species is uniform in shape, precisely located and oriented, and can be used in determinations of cyst morphospecies. The paraplate or paraplate field which is opened or detached in the formation of an archeopyle is called the operculum.

Thecate dinoflagellates reproduce both sexually and asexually, depending on season or as environmental conditions begin to change (Lee, 1992). When sexual reproduction occurs, a planozygote or zygospore is formed. This is usually (though not always) in the form of an organic walled (sporopollenin) resting cyst. The tabulation pattern of the cyst is directly related to the thecal tabulation pattern, although the cyst may or may not reflect the original theca. After an inert period, spent either within the water column or within bottom sediments, the zygocyst germinates and reestablishes a motile phase. Some dinoflagellate cyst morphologies, such as spines and appendages, retard sinking (Parsons 1977). Because excystment is controlled by both water temperature and cyst age, cysts falling below about 200 meters rarely germinate (Lentin 1980).

theca  
 plate  
 suture  
 (tabulation  
 epitheca



cyst  
 paraplate  
 parasuture  
 paratabulation)  
 epicyst

cingulum  
 flagellar  
 pore  
 hypotheca  
 sulcus

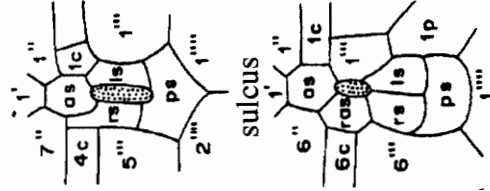
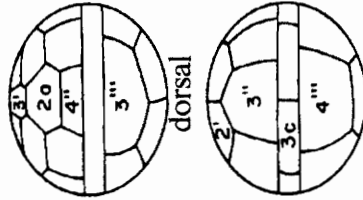
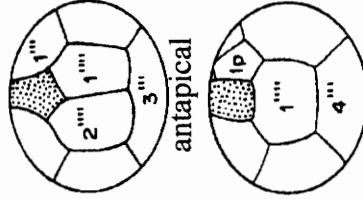
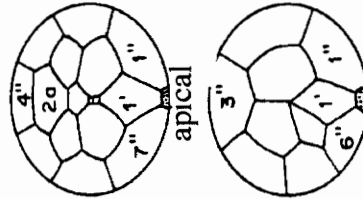
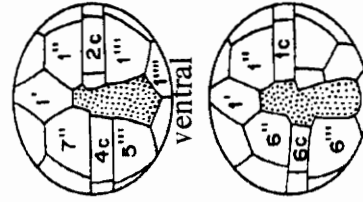
antapex

archeopyle = excystment feature

(After Evitt, 1985)

## Peridinium

2 pr, 4', 3a, 7", 4c, 5", Op, 2'''



Opr, 4', 0a, 6'', 6c, 6''', 1p, 1''''

## Gonyaulax

## **DINOFLAGELLATES: FACTORS CONTROLLING ABUNDANCE, DIVERSITY AND PRESERVATION**

### **General Statement**

Many factors control dinoflagellate abundance in living populations and in the fossil record. As microscopic phytoplankton, living dinoflagellates respond to environmental conditions of the photic zone. Some modern species encyst during unfavorable conditions. Others encyst as part of a seasonal life cycle, while others, especially open marine forms, do not produce cysts (Dodge, 1995). When produced, cysts must often sink through great water depths to become preserved in the fossil record. Many factors, such as feeding, breakage or chemical dissolution may destroy dinoflagellate cysts before they reach the sea floor.

Even after cysts reach the sea bottom, other factors such as bioturbation or diagenesis can still selectively destroy them. Because of both encystment patterns and taphonomic factors, fossil cyst assemblages may not reflect the composition of living assemblages.

### **Physical Factors Controlling Abundance and Diversity**

#### **Temperature**

Ambient temperatures and temperature fluctuation influence dinoflagellate abundance via effects on metabolism and photosynthesis (Loeblich, 1967; Parsons, 1977). In addition, temperature difference is often a cause of water column stratification which may encourage dinoflagellates. Dinoflagellates exist at wide ranges of temperatures, but generally prefer warm waters. Maximum reproduction rates often occur near upper temperature tolerance levels (Taylor, 1987). Reproductive optimum temperatures for nearshore species are from 15° to 25° C., and for oceanic species occur at 20° C. (Loeblich, 1967).

Dinoflagellate populations are sparse in cold seasons or colder waters. Diatoms tend to replace dinoflagellates in cooler conditions (Guillard and Kilham, 1977).

#### **Salinity**

Marine surficial salinities vary with mixing, upwelling, and evaporation. In

nearshore environments, freshwater flows out over more saline estuarine and coastal waters, maintaining a less saline surficial layer in which phytoplankton congregate (Thurman, 1994).

Nearshore and estuarine dinoflagellates range from obligate freshwater forms (such as *Peridinium limbatum*) to cosmopolitan euryhaline species. Terrestrial nutrients in estuarine environments correlate inversely with salinity (Mallin et al., 1991). Thompson (1987) notes many “stenohaline” associations may be dependent on, or intolerant of, runoff related constituents other than salt. Generally, peridiniacean dinoflagellates prefer near-shore environments. The presence of proximate and peridinioid cysts suggests marginal marine or reduced salinity conditions (Harker et al., 1990).

Offshore waters are relatively stable salinity-wise, and contain mixed cosmopolitan and stenohaline (or runoff-intolerant) dinoflagellate species, including many gonyaulacoid types. In the fossil record, the presence of mainly spiny chorate and gonyaulacoid or gymnodinioid cysts reflects normal marine salinity (Harker et al., 1990).

### **Eddies, Fronts and Upwelling**

Flow of deep marine water into shallow coastal areas promotes mixing or upwelling. Fronts, however, occur at the contact between two water masses having different densities, due to temperature, salinity, or other factors. In these situations, a relatively sharp boundary occurs between the two water masses instead of a well-mixed zone. Fronts are often zones of convergence, capturing foam, floating wood and detritus, and near-surface phytoplankton (Bowden, 1983). Eddies occur when a portion of an intruding watermass is “spun off” into dissimilar coastal waters.

Eddies, fronts and upwelling are often associated with high nutrient conditions, and are often sites of highly productive, though sometimes patchy, phytoplankton growth (Tappan, 1985). Phytoplankton productivity in response to front-induced conditions may continue at times when frontal conditions are no longer present (Barry and Dayton, 1991). Where stratification or convergence occurs, dinoflagellates may have functional advantage over other phytoplankton, and occur in high numbers and/or diversity. Boundary regions too cold for dinoflagellates are occupied primarily by diatoms (Smayda, 1967; Guillard and Kilham, 1977).

Fronts often separate different floras, associated with the different water masses. Phytoplankton populations along fronts and upwelling zones often contain higher diversity,



and/or different species associations than do waters on either side of the boundary (Gray, 1981). Bottom sediments beneath the boundary may preserve records of phytoplankton from one or all flora, depending on sedimentation patterns, sediment mixing, and locational stability (or seasonal repeatability) of the boundary. Long-standing boundary conditions may locally affect the gradient of change of fossil phytoplankton. Even sharp boundaries, however, will produce a "blurred" record due to mixing.

### **Nutrients**

Dinoflagellates utilize nearshore (terrestrially-derived) nutrient sources more so than upwelled oceanic nutrients (Thompson, 1987). Nitrogen and phosphorus availability are primary limiting factors in near-shore conditions. Iron, organic nutrients, and vitamins (such as B12) are important limiting factors, particularly in open oceanic conditions (Parsons, 1977; Bonin and Maestrini, 1981; Thompson, 1987).

Nitrogen and phosphorus (usually at a ratio of 15N to 1P) tie linearly to productivity (Parsons, 1983). In summer, stratified near surface waters are often deficient in nitrogen and phosphorus, reducing dinoflagellate productivity (Lieberman, 1994). During dinoflagellate blooms, sexual reproduction (and encystment) occurs when nitrogen levels eventually drop, suggesting that it is nitrogen which becomes a limiting factor (Weeks, et al., 1993). Decrease in nitrogen available might instead trigger sexual reproduction by changing the local carbon/nitrogen ratio (Bruce Parker 1994, personal communication). In this case, algal productivity may become self-terminating before nitrogen becomes a true limiting nutritive factor.

Phytoplankton require iron for their respiration and photosynthesis reactions (Glover et al., 1978). In near-shore areas, iron is available from terrestrial runoff, though only some forms are chemically useful to algae (Parsons, 1977). Open ocean waters are generally iron deficient (Menzel, 1961; Tranter, 1963). Studies by Ryther and Kramer (1961) found that oceanic dinoflagellates (especially Gonyaulicoids) have lower iron requirements than do near-shore types (including the Peridineeaceans).

Organic nutrients and vitamins are often derived from terrestrial runoff, and are more available in nearshore, less saline waters (Gallegos, 1992; Carlsson and Graneli, 1993; Lopez, 1993).

## **Light**

Light levels vary with turbidity and season. Seasonal light variation operates in conjunction with temperature to trigger encystment and productivity changes (Lee, 1992). In turbid or eutrophic environments, some dinoflagellates ingest other plankton, including diatoms (Marshall and Alden, 1993; Mullin, 1995). Dinoflagellates are often replaced by diatoms in times or areas of high turbidity (Marshall and Alden, 1993).

## **Biological Responses to Physical Factors Succession**

Succession is an ongoing shift in community structure or makeup as it continuously readjusts to changing environmental conditions (Margalef, 1968; Drury, 1973). Phytoplankton often progresses through a seasonal succession beginning with early season pioneer species and progressing to later, stability-associated species such as dinoflagellates. Within each successional range, incumbency plays a significant role. Once a species group is established, it dominates resources until conditions change, and it is displaced by a different group (Rosenzweig, 1991).

Study of seasonal succession suggests that dinoflagellate species occurring together in a fossil assemblage may not have existed together or interacted significantly with each other while alive. For example, a dinoflagellate species dominant when the water is first warmed in the summer may not interact with one dominant only in the fall. The length of dominant periods also changes with conditions: a dinoflagellate may flourish in tropical environments over 8 months of the year, while the same species will be active only through July and August in colder waters.

The fossil record indicates that dinoflagellate species groups or associations also change with environmental conditions over longer periods of time. Offshore sediments have different associations or assemblages of dinoflagellate cysts than do sediments deposited in shallow estuarine or shallow coastal settings. Fossil dinoflagellate cysts change stratigraphically in sedimentary layers in a way which correlates with transgression/regression patterns of the paleocoastline (Habib, 1989). The fossil record may suggest persistence of general estuarine dinoflagellate "gene pools," sensu Buzas and Culver (1994) through time.

A species (or species association) may become dominant when environmental

conditions become optimal to its particular needs (Harrison and Quinn, 1989). During other times, it may remain present in the local environment, but remain encysted, suppressed, or as a minor fraction of the community. Oliver (1990) refers to continuous presence of many species, even when not dominant (in regard to forest stand dynamics) as an “initial floristics model.”

### **Inoculation**

Survival of a dinoflagellate species in a region depends on at least some cysts remaining in or returning to the local environment (Keafer et al., 1992). When cysts hatch under optimal conditions, they may remain rare, or they may reproduce asexually in great numbers, potentially allowing even a single cyst to produce a large population.

Cyst abundances less than 1% (“rare”) in rock samples are usually not considered to be statistically significant (Ludwig and Reynolds, 1988). Even rare cysts, however, are present in local sediments in large numbers, and represent potentially viable stock during that time. Even cyst types not encountered within sample counts may be present, although below a threshold where they are found in most individual samples. Cairns (1969) discusses the threshold density of finding a single specimen of a rare species within modern ecosystems, concluding that one almost never finds “rare” living species in samples, although their overall population density must be adequate to sustain them.

### **Dinoflagellate Blooms**

Optimal environmental conditions may give rise to large populations of dinoflagellates. These dinoflagellate blooms, or “red tides” can occur periodically due to seasonal or localized environmental conditions. Environmental factors conducive to dinoflagellate blooms include warm stratified surface water, gentle onshore winds, intense sunlight, and high nutrient levels (Lee, 1992). Nearshore environments and upwelling regions are often nutrient-rich, and often become sites of abundant dinoflagellate growth (Harris, 1986, Xiaohong et al., 1991). Dinoflagellate blooms often occur after cessation of upwelling, and/or directly diatom blooms. Diatoms “set up” specific nutrient and pH conditions amenable to dinoflagellate blooms (Smayda, 1967; Hinga, 1993) Dinoflagellate blooms are often toxic, producing extensive fish kills, poisoning shellfish, and destroying much of the local non-dinoflagellate phytoplankton (Prakash, 1967; Mullin, 1995).

During late stages of a bloom, mass encystment of dinoflagellates takes place (Heiskanen, 1993). Cysts remain inert within the water column or sediment, sometimes

for several years until bloom conditions return (Keafer, 1992). Those buried deeply may never hatch.

Dinoflagellate biomass is greatest in near-shore and/or estuarine environments. Dinoflagellate species diversity drops off from open shelf environments toward shore, roughly correlative with decreasing water depth (Hulbert, 1963; Wall, 1977). In estuarine environments the reduction may be so drastic that the dinoflagellate community is represented by only a few tolerant species.

Many living dinoflagellate species cannot cope effectively with rapid salinity changes, or with temperature changes greater than about 5° C at a time (Parsons, 1977). Nearshore dinoflagellates often migrate geographically within estuaries in response to drastically changing environmental conditions (Habib, 1989; Firth, 1993; Burkholder, 1995). Vertical diurnal migration strategies allow modern nearshore and estuarine dinoflagellates to exist in variable, or eutrophic settings (Santos and Carreto, 1992; Salonen, 1994). Some nearshore and estuary dinoflagellate species can encyst within minutes when exposed to unfavorable conditions. Such production of resting cysts seems a specific adaptation to unstable environments within estuaries (Burkholder, 1995).

Large numbers of fossil specimens representing a single species may indicate environments where blooms occurred repeatedly. Whereas diverse dinoflagellate fossil assemblages may reflect seaward paleoenvironments low diversity may indicate nearshore or estuarine conditions.

### **Factors Controlling Preservation**

Copepods and other zooplankton eat encysted dinoflagellates, breaking, and often destroying, the cysts. However, some cysts preferentially pass through the zooplankter gut intact and become incorporated into fecal pellets. The relatively large fecal pellets settle rapidly to the sea floor. Buck and Newton (1995) note that most dinoflagellate cysts become incorporated into the fossil record in this manner. Additionally, zooplankters prefer some prey species over others, introducing feeding bias into the record.

Within bottom sediment, changes in sedimentation rate can concentrate or dilute the relative numbers of cysts present (Rutherford, 1994).

Bioturbation and other time averaging obscures details and decrease resolution of

the dinoflagellate record (Walker and Bambach, 1971). Time averaging may limit dinoflagellate data resolution to thousands or tens of thousands of years (Flessa, 1993).

Diagenesis also influences dinoflagellate preservation. Cysts are damaged or destroyed by local pore fluid and sediment chemistry, especially where hot or basic. Most dinoflagellate cysts contain sporopollenin which is resistant to chemical breakdown (Evitt, 1985; Traverse, 1988) except to oxidation, such as in high pH conditions. In calcareous samples, for example, dinoflagellate recovery is often poor. In ferric sandstones and siltstones, as well as in close proximity to Jurassic and Cretaceous dinosaur bones in lakes and brackish seaways, general palynomorph recovery is often poor (Alfred Traverse, personal communication, 1995). Recovery in organic rich sediments, and in relatively acidic or reducing conditions is often high.

#### **IV. THE YELLOW SEA AS A MODERN PHYSICAL ANALOG FOR THE SALISBURY EMBAYMENT**

In developing a model of the Salisbury Embayment, The Yellow Sea is an appropriate modern analog. It (along with inner portions of the East China Sea) is similar in extent, depth and physical structure, latitude, and eastern continental location to the Salisbury Embayment (figure 6). It might share similar circulation and current structures with the former Salisbury Embayment.

The Yellow Sea is a shallow embayment bordered by China to the West and North, and by Korea to the East. It opens to the East China Sea to the south. The Yellow Sea bottom structure includes a north-south oriented deep trough (located somewhat nearer to Korea than to China) that provides the Yellow Sea with a core of warm, saline marine water derived from the Kuroshio, or regional boundary current.

Zhou (1989) has described the dynamic deposition systems of the Yellow (and East China) Sea. Even though influenced by human activities (Milliman et al., 1987), the depositional patterns and structures (Wang and Zhu, 1990; Park et al. 1992) may be analogous to those of the Aquia, Marlboro and Nanjemoy Formations of the Salisbury Embayment (See also Je et al. 1988). Glauconite is produced in deeper, offshore waters of the Yellow Sea Embayment and East China Sea (Lu-Xiaozhen, 1989).

Circulation within the Yellow Sea appears driven by wind-induced (i.e., Ekman upwelling) and tidal currents (figure 6). These currents wax and wane with seasonal monsoon winds and freshwater input (Park, 1986). Warm Kuroshio water enters the deepest portions of the embayment via the Yellow Sea Warm Current (Hwang and Choi, 1993).

In the northern half of the Yellow Sea Embayment, seasonal winds create an extensive surface zone of upwelling and mixing, forcing warm marine water to migrate northward as far as the Bohai Gulf (Tomczak and Godfrey, 1994) The marine influence varies seasonally as a function of wind strength and direction.

The China Coastal Current transports cool, low salinity water southward from northern portions of the Yellow Sea. A similar coastal current flows southward along the western coastline of Korea. These currents are strengthened seasonally by monsoonal freshwater influx into the margins of the Yellow Sea Embayment.

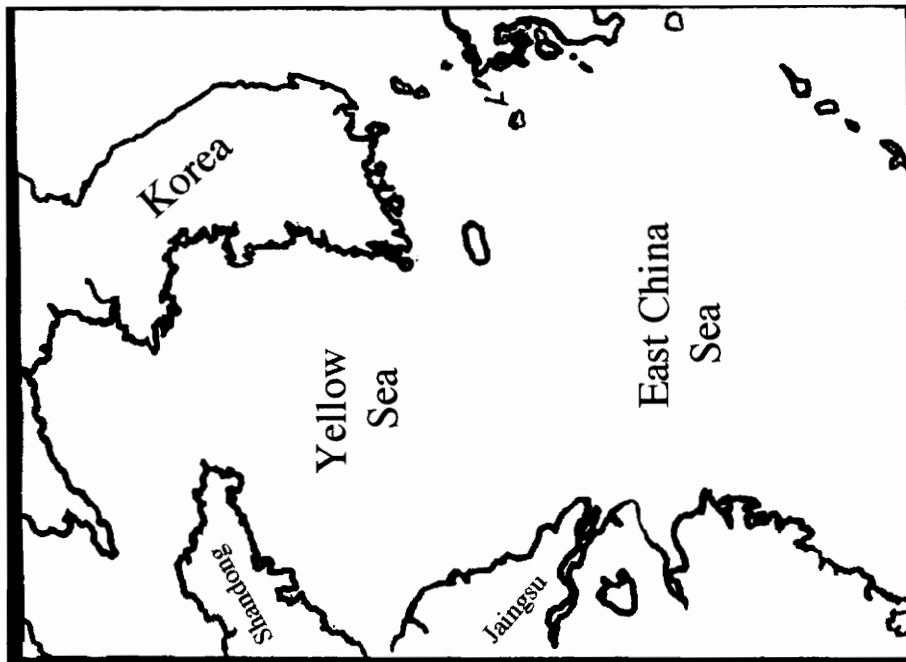
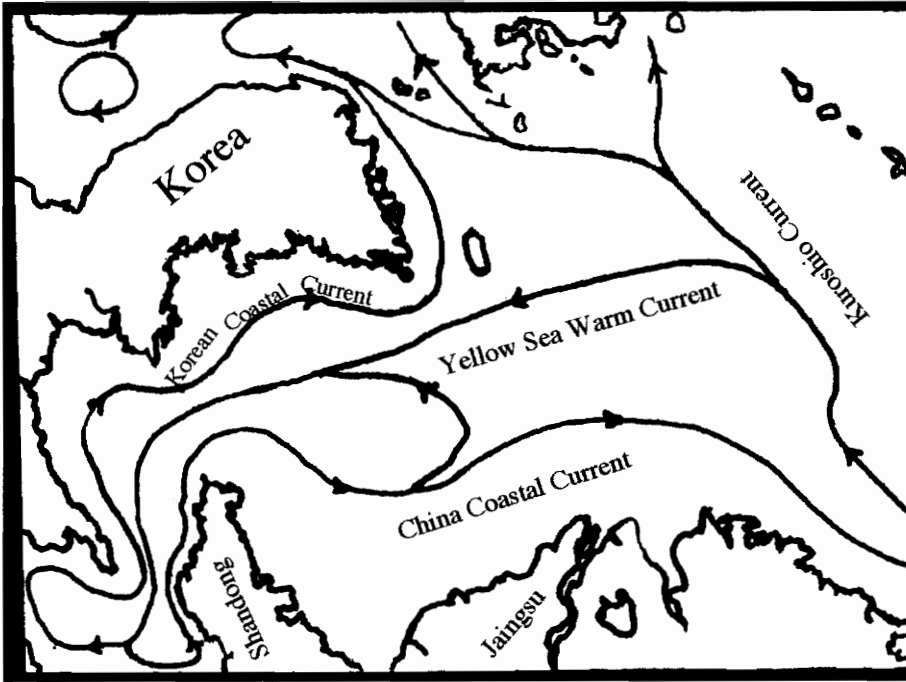


Figure 6. The Yellow Sea, showing coastline features and major currents.

The northward and southward flows are separated by lateral oceanic fronts (see figure 6). Here, the warm marine Kuroshio-influenced water mass meets with cooler less saline coastal water. Due to density differences, the two masses slide past each other without mixing significantly.

Freshwater drainage into the Yellow Sea Embayment creates local estuarine conditions, bounded by salinity and turbidity fronts, which separate them to varying degrees from nearshore coastal waters. These regions include portions of the Bohai (Yellow River) Gulf (Deng, 1988), the Yangtze River (Tomczak and Godfrey, 1994), the Kyongg Bay, the Kum River Estuary, and other areas (Yoo, 1986).

This physical circulation pattern produces a system of three relatively distinct water masses. They are: offshore, onshore, and estuarine. Offshore (warm, more marine) conditions are separated by an oceanic front from nearshore (cool, less marine) conditions. Salinity and/or turbidity fronts in turn separate nearshore conditions from more extreme and brackish conditions within estuaries (Seung et al. 1990; Shim et al., 1991)

In general, temperature and salinity in nearshore waters increase seaward. Offshore waters continue to show seaward temperature increases, although salinity there remains at about 34.4 o/oo (Kim et al. 1991; Cheong et al. 1992). Yellow Sea temperatures range from less than 2° C in nearshore winter waters, to about 27 °C in offshore summer waters (Kim et al. 1991; Tomczak and Godfrey, 1994). Temperatures of the warm Kuroshio Current range from 16 °C in winter to 29°C in summer. Its salinity ranges from 33.7 to 34.9 o/oo (Kim et al. 1991, Schweitzer, 1993).

During the summer season, thermoclines and stratification become well-developed in the Yellow Sea, especially in conjunction with front regions (Matsuda et al., 1989; Seung et al., 1990). Deep waters are characterized by high salinity. Surface waters are often nutrient poor (Chang et al., 1990; Yang and Kim, 1991). Primary productivity along the regional fronts is higher than that of either the outer stratified waters or the inner mixed coastal waters. This productivity is associated with increased light (decreased turbidity) and high nutrient levels along the fronts (Choi, 1991)

Because of the relatively cool temperatures and turbid conditions in shallow nearshore portions of the Yellow Sea, diatoms dominate the phytoplankton (Chang, 1986,



Choi and Shim, 1986, Lee et al., 1989). Dinoflagellate populations are sparse, making up portions of the phytoplankton primarily in late summer. The dinoflagellates occur primarily in warmer, more stratified waters, and in association with the regional fronts. Within the estuaries, phytoplankton is reduced by levels of turbidity. Dinoflagellates and other phytoplankton are sparse in highly turbid environments (Shim et al., 1991), with diatoms predominant.

## V. METHODS

### Sources of Data Sets

Samples for most data sets were collected by former graduate students in the Virginia Tech palynology program from surface outcrop localities within the former Salisbury Embayment (figure 7). Witmer collected samples from the Oak Grove core (Witmer, 1987). Samples were numbered stratigraphically upwards at each collecting site. Samples were prepared via standard palynological maceration techniques to recover the organic-walled dinoflagellates. The technique destroys calcareous and siliceous microfossils such as diatoms and foraminifera.

Taxonomic identification and specimen counts were made by the original workers. In this study, original data sets were transformed to percentage equivalent values to allow comparison between data sets. Transformed percentage data sets are presented in Appendix A.

The species names used in this study are those of the original workers. This is not a taxonomic project. The reader is referred to the original thesis sources for species descriptions (See also Barss and Williams, 1983).

### Correlation of Samples (Cluster analysis)

Cluster analysis is a straightforward form of multivariate analysis, whereby data are grouped into a hierarchical classification based on quantitative measure of their individual similarity of occurrence to one another. The output for a cluster analysis is a dendrogram, which groups and ranks the samples by similarity coefficient. Samples whose components are similar group together with high similarity coefficients, producing short dendrogram branches. In this study, cluster analysis is used to divide the combined Aquia-Nanjemoy dinoflagellate data set into dinoflagellate similarity clusters.

Cluster analysis begins with a comparison of (occurrence) similarity of all samples being tested. This produces a data matrix of similarity coefficients. From this matrix, the two samples which are most similar cluster first. The average value of this first cluster element is then compared with all other remaining values, and the next most similar sample is added to the next cluster level. This comparison and clustering continues until all samples are integrated (or forced) into the hierarchical dendrogram. After a sample is clustered, its coefficient value is averaged into the group value, and the sample is no longer

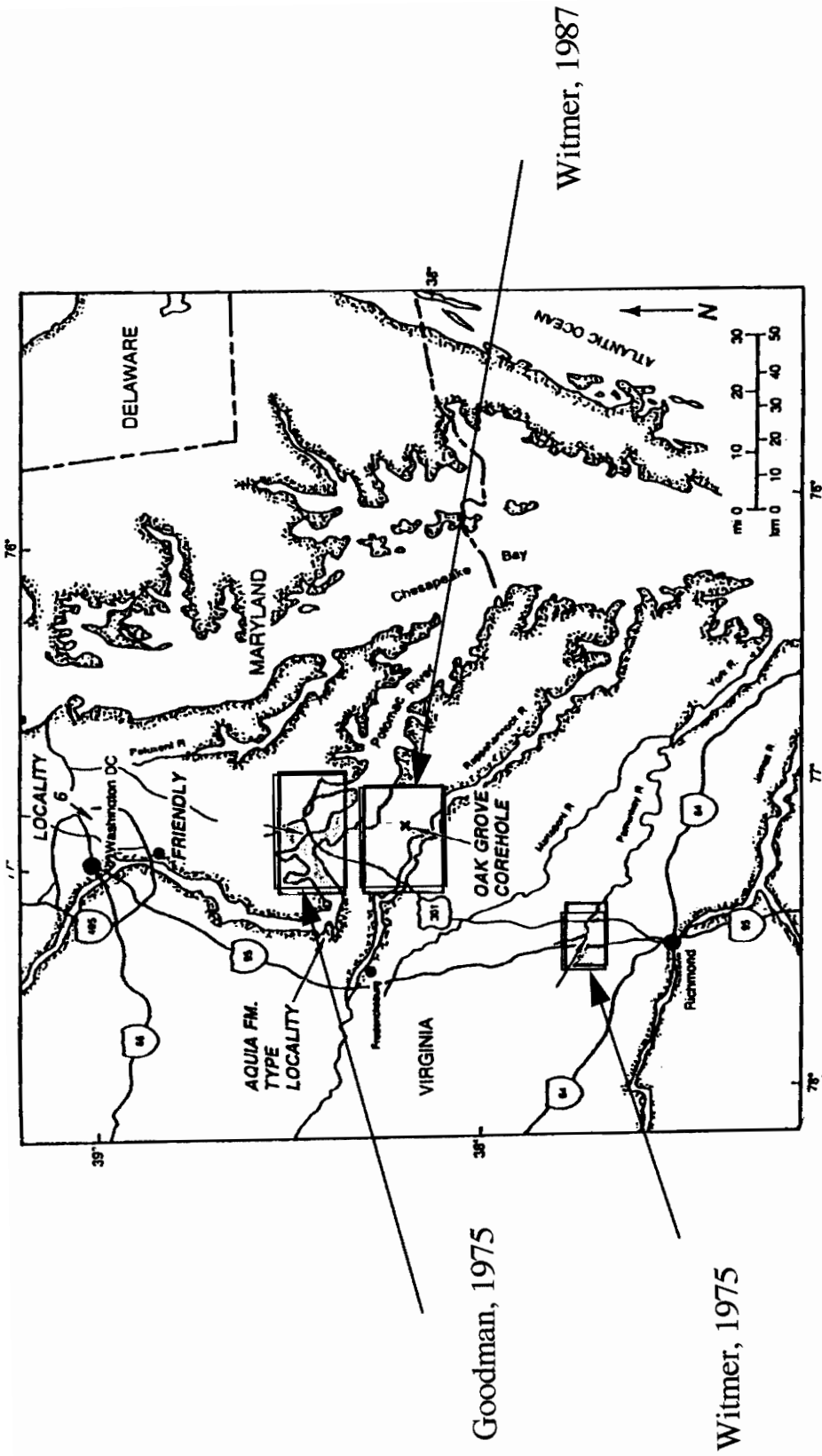


Figure 7. Site Locations

compared individually with any other sample in the data set. As new samples are clustered, their individual relationships become less distinct. Several results are possible. For one, the precise relationships of samples within cluster groups may not be accurate, although the cluster grouping is accurate. Clustering which occurs at low coefficients of similarity may not be as reliable as those grouping at higher similarities. Clusters at low similarities may be artifacts, associated only because the calculating engine joins together all loose dendrogram branches before terminating the analysis. Cluster analysis, however, is sufficient to quantitatively sort groups with similar characteristics, even though any suggested relationships between such clustered groups may be suspect (Size, 1987).

### **Q-mode Cluster Analysis**

Q-mode cluster analysis forms hierarchical clusters of samples by comparing the similarities of occurrence of the species in each. Thus, Q-mode clusters represent physical or geographical units of similarity. They serve to determine potential boundaries of environmental settings among samples. Misunderstanding exists on how Q-mode analysis produces similarity clusters, particularly in gradient situations. Given a constant environmental gradient, Q-mode analysis forces the data into arbitrary subdivisions across the gradient, generating boundary conditions where none exist. However, Q-mode analysis will place group boundaries along non-constant gradients preferentially at points where boundaries do exist, or where the slope of the gradient changes most significantly over a short distance. Because the effects of even relatively sharp environmental boundaries grade into "transition zones" (expressed as a change in gradient slope.), Q-mode analysis is an appropriate tool.

Clustering for this project was done via a FORTRAN program written by Robert Plants (1977) for the IBM370 mainframe, and modified by Tom Rounds and Arnie Miller (1981; see appendix C). This program provided consistency with earlier cluster analysis by Virginia Tech palynology students (See Welch, 1986). The Virginia Tech IBM 3090-300 E mainframe computer was used for this analysis. In this project, the Czekanowski (or "dice") coefficient was used with UPGWA for comparing and choosing similar sample values within the clustering program. See the Results section for the output dendrograms of these analyses.

Q-mode dendrograms were also produced from data subsets containing only Very Common (greater than 25% of the sample makeup), common (greater than 10% of the sample makeup), intermediate (between 10% and 1% of sample makeup) and rare (less than 1% of the sample makeup) species. These dendrograms provide comparison of

species within these general frequency groups in the “paleocommunity.” See the Results section for discussion of these separate dendrograms.

### **Fossil Community Definitions**

Coastal dinoflagellate associations change geographically according to environmental conditions from offshore to nearshore to estuarine regions (Steele, 1978). These regions may be gradational, or they may be separated by physical environmental boundaries, such as fronts (Springer and Bambach, 1985).

A sample of plankton consists of phytoplankton (including dinoflagellates, diatoms, etc.) and zooplankton (including copepods, etc.). In the living sample, the species and (potential) interactions between them form a local community. Often, only a subset of the overall local community is studied (Bennington, 1995); such as in this study the “dinoflagellate” local community is the unit of interest. An aggregate of statistically significantly similar living local communities in a region constitutes a community. The community boundaries can be defined by the degree of statistical similarity or difference between recurring local communities. In the case where local communities resemble one another, but are not statistically the “same,” then the term community type is used (Bennington and Bambach, 1996).

Fossil assemblages may not closely resemble the original living community. Bennington and Bambach (1996) suggested using fossil record definitions comparable to modern community definitions. An assemblage present in a fossil sample is thus termed a local paleocommunity. An aggregate of statistically identical local paleocommunities is a paleocommunity. Related sets of local paleocommunities that are similar, but not statistically the same, are designated as paleocommunity types. Paleocommunity types group together in statistical cluster analysis of fossil data sets (Bennington and Bambach, 1996). Generally, temporally or geographically similar samples should be similar (particularly when the environment appears similar), whereas samples separated by time, geography, or environmental conditions, are often more dissimilar.

Bennington (Bennington, 1995; Bennington and Bambach, 1996) used ANOVA (analysis of variance of means, as described below) for determining “sameness” between local paleocommunities in benthic marine assemblages (See also Zenetos, 1991). ANOVA analysis is used in this study to compare member samples of cluster groups.

### **ANOVA Analysis**

Analysis of Variance compares the means of individual sample groups. ANOVA calculates two components of variance (within-group variation, and between group variation) and then compares the ratio of the two components. The resulting number (or F-value) suggests the sameness or difference between members of the group. A probability (or P-value) is then calculated for the likelihood that the F-value will be more extreme than the one reported. Generally, a P-value of less than 0.05 indicates that one or more samples within the group are statistically dissimilar. See Sokal and Rohlf (1969; also Young, 1962; Fredericksen, 1974) for a description of ANOVA procedures.

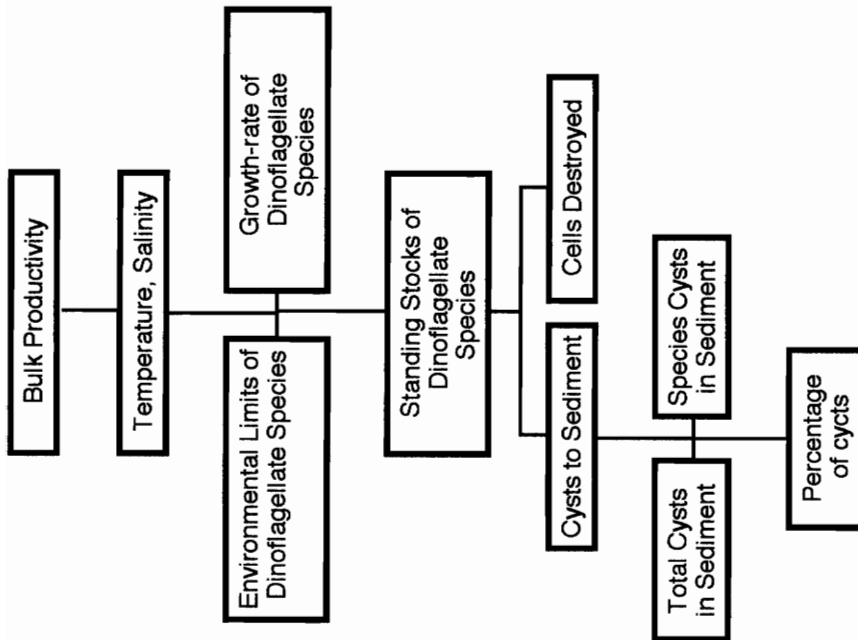
All ANOVA statistics for this project were run using the Macintosh program STATVIEW, on a Macintosh IIsi computer. See Appendix A for ANOVA results of individual cluster groups.

### **Dynamic Systems Modeling**

A major project goal was to construct a dynamic systems model of dinoflagellate species behavior using oceanographic and biologic data. Initial stages of this modeling used the dynamic systems modeling program STELLA II for the Macintosh computer. For a discussion of STELLA modeling, see Richmond (1987).

STELLA II modeling explored the correlation between dinoflagellate cyst distribution and paleobathymetry, as developed by others. In this model, six dinoflagellate species were created whose populations operated independently of each other and dependent only on depth. Two were “shallow water“ species, two were “deep water“ species, and two were “intermediate” depth species. Population growth rates differed between the species. No differentiation was made between living thecal populations and fossil cyst populations. Cyst counts for possibly analogous species from a portion of the Oak Grove core data set (Witmer, 1987) were entered into STELLA II format for direct comparison with the model species. Similar estimated depth (“shallow up”) curves were plotted for the data and run for the STELLA II model.

More complex modeling required use of FORTRAN programming. Code structures for the FORTRAN model could be specifically designed for the modern analog of the Yellow Sea. FORTRAN models for this project were designed and run on an IBM 486 personal computer, using KEDIT and a WATFOR87 compiler.



Physical Conditions are Based on the Yellow Sea

Productivity is Related to Depth

Ten Dinoflagellate Species are Used

Populations are Sampled Monthly

Cyst Stocks are Reset Periodically to Zero

$\% \text{ Cysts} = \frac{\text{Species Cyst Count}}{\text{Total Cyst Count}}$

Figure 8. FORTRAN Model Flowchart

FORTTRAN modeling for this project began with the construction of a simple linear-donor-controlled model, in which the results of each model step are controlled only by the previous step. The flow chart diagram of this structure is shown in figure 8. By successively adding complex relationships to the model code, three time varying non-linear Figure 8

models were created. The final FORTRAN model code is shown in Appendix B. Data for the final models were drawn from information discussed in the background sections of this report (Also including Balech, 1967; Atkinson et al., 1985; Begon et al., 1986; Brock and Madigan, 1991). The FORTRAN model is broken down into three separate portions. The first is a module describing environmental parameters and relationships. The second presents hypothetical dinoflagellate species and their environmental requirements, growth parameters, encystment parameters, and environmental limitations. The third contains the assumptions and procedures associated with dinoflagellate cysts added to the sediment, and with time averaging. Results of these FORTRAN models are given in the Results section.

### **Phase Plane Interactions**

Phase planes are a method for graphically testing possible species interactions. In this graphical technique, two species' population values are compared relative to each other, rather than as a time series. If one species is systematically affecting the other (Davidson, 1995), then the population values of both species may show the interaction geometrically when plotted in x-y phase plane space. When oscillations in population numbers of interacting species are stable with respect to each other through time (such as in a traditional Lynx versus Hares predation model), the phase diagram "stability field" produces a circular equilibrium figure, which repeats through time.

When the parameters of the system interact convergently toward an equilibrium condition, the stability field is expressed as an inward spiral, toward a circular equilibrium figure. When the parameters diverge from equilibrium, the spiral is outwards.

There are specific patterns which are stable and others which are not (figure 9). Interaction figures in real data may be overprinted by strong environmental signals, or interactions with more than one species at once. Two species reacting similarly to each other may both be reacting to a third over-riding factor. In such cases, the phase plane figures are erratic, and produce figures unlike those of the accepted archtypes. Interaction, where it exists, may be most apparent during times of overlap and replacement between populations, or may all be below the level of resolution of the data set.



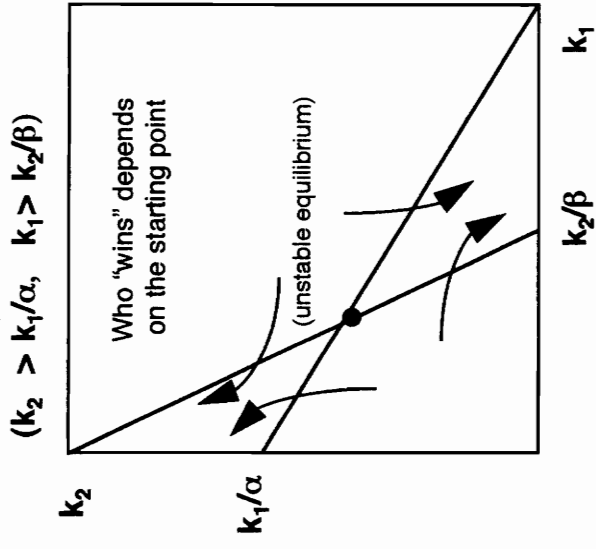
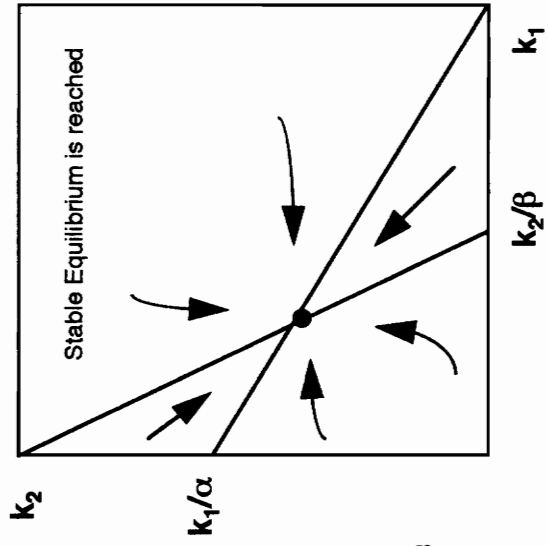
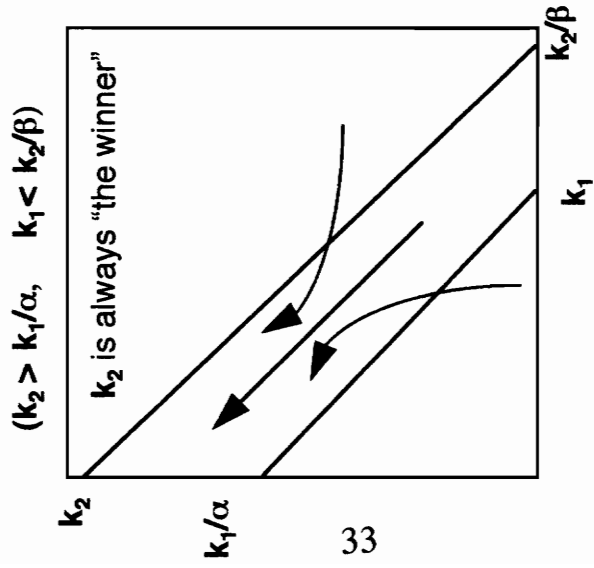


Figure 9. Phase Plane Archetypes

## VI. RESULTS AND DISCUSSION

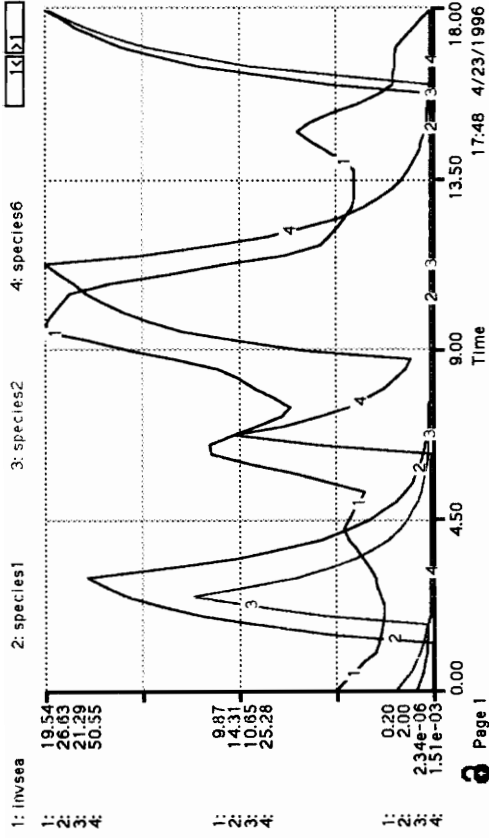
### Stella Dynamic Systems Model

The STELLA II model assumes that instability and variability increases with sea level shallowing. It plots species frequencies dependent on the sea level and according to simple growth rate equations in the model code. The resulting time series (figure 10a) is graphed against a “shallow-up” relative sea level curve from a portion of the Oak Grove Core (Witmer, 1987).

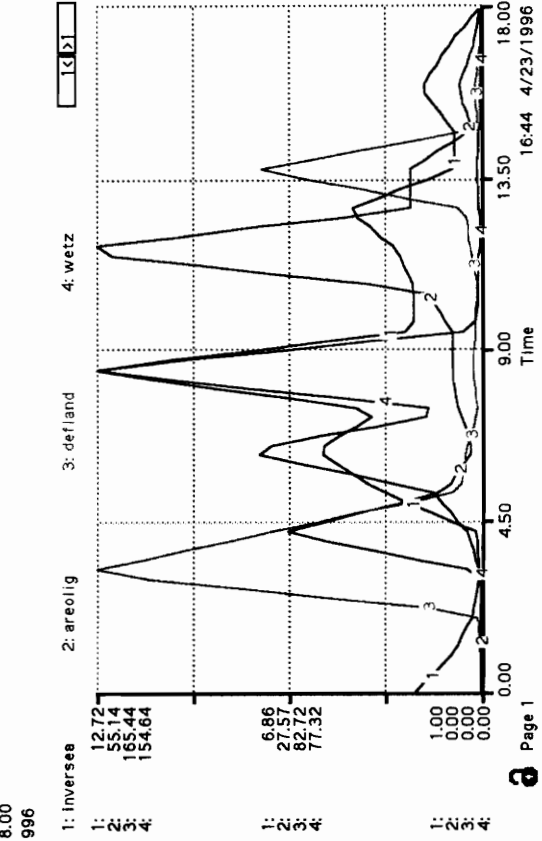
The model output can be compared with equivalent data plots of *Areoligera*, *Deflandria*, and *Wetzeliella* from the Oak Grove core (figure 10b). Line 1 (inverse) in both models represents the inverted sea level curve.

In both the model (top), and in the graphed data set (bottom), “deep” versus “shallow” species are present. The most noticeable contrast between the model and the plotted data is the finer scale shape and position of the frequency curves from the real data. In the model, there is a slow (mathematical) response time between sea level change and calculated dinoflagellate response. This is primarily due to the iteration period used within the model. Shapes of the modeled curves are simpler than those from data because they are defined by relatively simple mathematical modeling equations.

In general, the STELLA II model suggests that depth correlations might be quantifiable for some species, if precise dinoflagellate counts and detailed sea level information were available. Still, associations with bathymetry do not directly suggest causation.



- STELLA II model output of three dinoflagellate species and (inverse) water depth, plotted against time.



- STELLA II plot of actual dinoflagellate data from a portion of the Oak Grove core. (data from Witmer, 1987)

Figure 10. . STELLA II Model Time Series

### **Results of the FORTRAN Dynamic Systems Models**

The FORTRAN models X, Y, and Z predict dinoflagellate populations according to depth related changes in temperature, salinity, and nutrient based on conditions of the Yellow Sea. Preliminary runs of the FORTRAN models using (cool) Yellow Sea temperatures produced small dinoflagellate populations. Increasing the average model temperatures by 5° C produced more extensive dinoflagellate populations. The warmer model variations are presented here. Results of the modeling suggest that the Salisbury Embayment may have been somewhat warmer than the modern Yellow Sea.

Dinoflagellate Model X reflects mixed conditions between nearshore and offshore waters, such as occur in the northern region of the Yellow Sea. Table B-1 (Appendix B) contains output from this model. For regions of mixing, Model X suggests gradational change in dinoflagellate associations from nearshore to offshore. Individual dinoflagellate species fluctuate seasonally as temperature and salinity conditions change.

Dinoflagellate Model Y describes the effects of a front boundary separating nearshore from offshore waters, such is found in lateral regions of the Yellow Sea. Table B-2 contains output from the second model.

For front-bounded conditions, Model Y suggests disparity between the two watermasses, their environmental conditions and the dinoflagellates present. Nearshore forms remain more separate from offshore forms. The physical and geographical change between the two groups remains sharper even with seasonal changes. Both mixed and front conditions may be found within the modern Yellow Sea. Lateral fronts remain at relatively constant locations, while the large mixed zone migrates seasonally, and has no sharply definable boundaries. Similar conditions may have existed within the Salisbury Embayment.

Table B-3 contains output from the third dynamic system model version, Dinoflagellate Model Z. This version produces model cyst data based on the dinoflagellate populations described above. Model depth increases by 5 meters at each sampling event. After each sampling period, all cyst values (including total cysts present) are reset to zero. The model reports cyst data for each species as a percentage of the total number of cysts present. Cyst species 11 and 12 represent secondary morphotypes of species 3 and 4.

Cyst information produced by the model suggests that dinoflagellate changes across

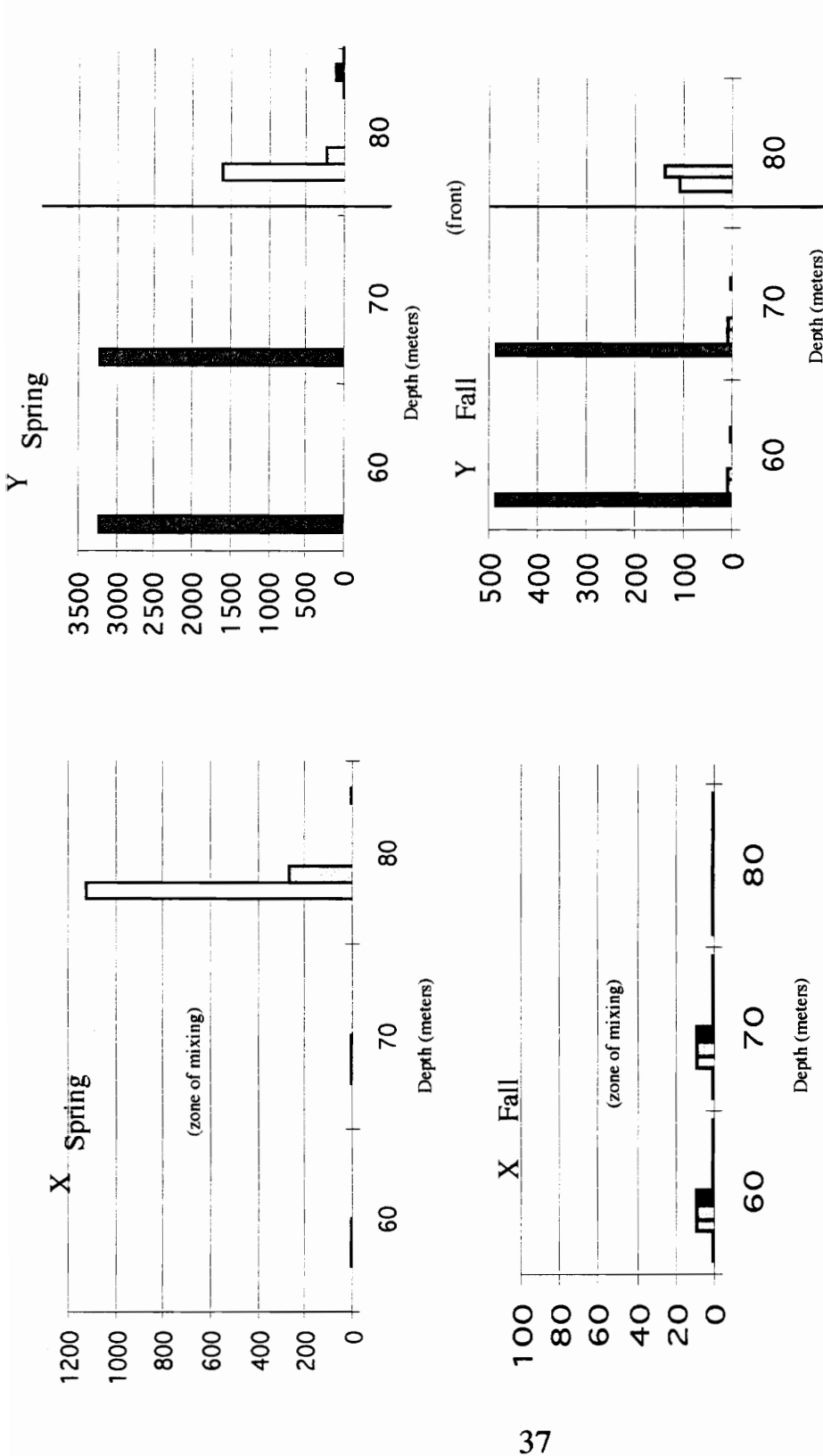


Figure 11. Representative model output for model X, showing effect of mixed conditions in Spring and Fall, and for Model Y, showing separation of nearshore and offshore conditions associated with a coastal front. (Bars represent species counts of model species 1 to 10.)

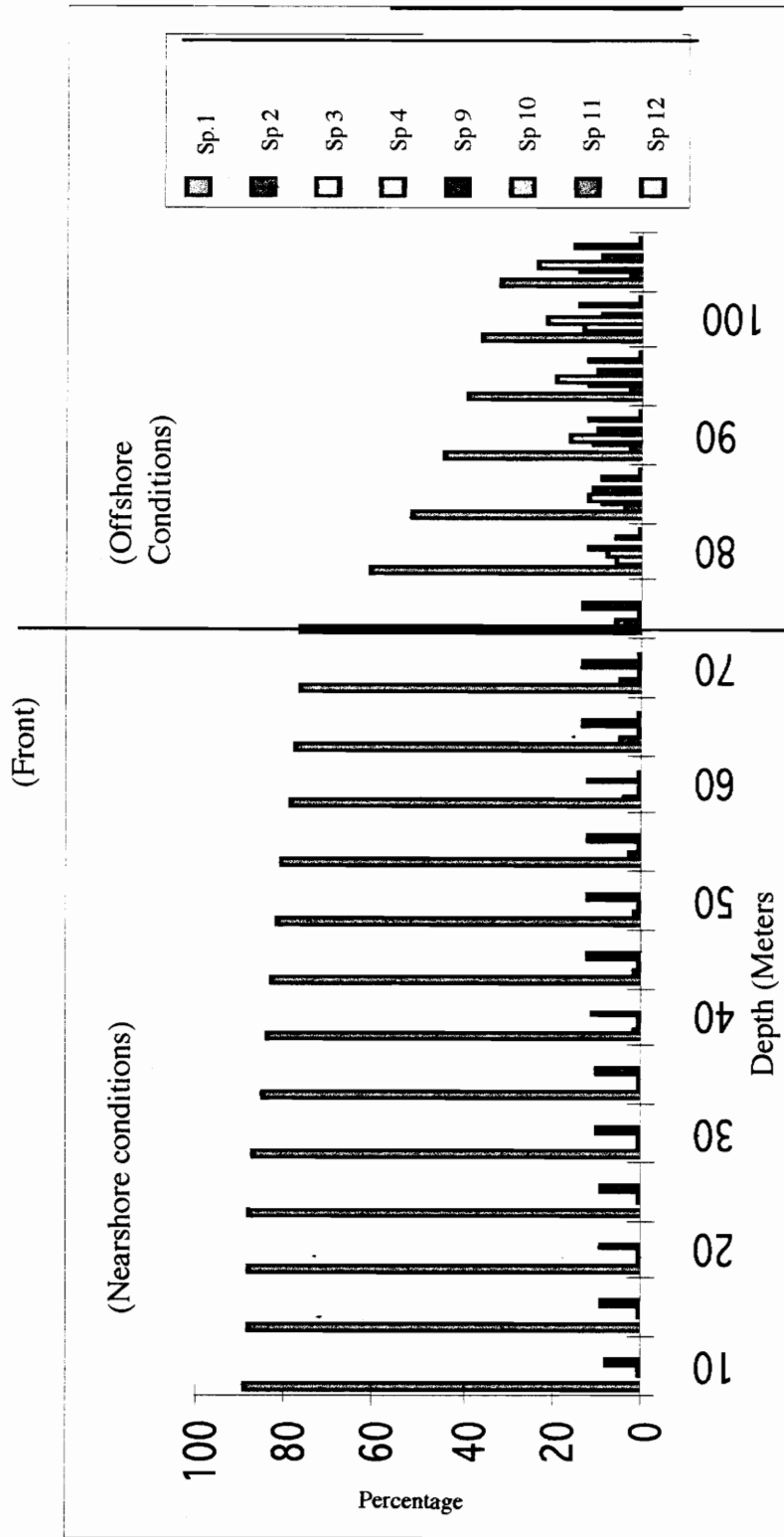


Figure 12. Model Z output, percentages of cyst types in sediment with depth. Modeled front is at 75 meters.

a front or boundary between coastal regions might be discernible in fossil assemblages (figure 11). This is also suggested by community analysis of the fossil dinoflagellate sets in this study. This assumes that effects of real front or boundary conditions would be significant enough to be recognized above background gradient effects (see figure 12).

The models, although preliminary, suggest that coastal dinoflagellate systems can effectively be described. Sets of potential conditions can be tested in a dynamic manner, provided that background modeling information is of good resolution, and modern analogs are carefully chosen.

### **Discussion of the FORTRAN Models**

Models simulate and simplify reality (See Caswell, 1972; Drew, 1985; Goodman, 1988). The modeler tries to reproduce dynamic patterns using the fewest parameters possible. He then considers additional factors individually, to add refinement and complexity to his model. The modeling is a method for discovering the relative importance of different processes in the dynamic system.

In this modeling, I began by addressing dinoflagellate species temperature thresholds. Modern dinoflagellates survive well across wide temperature ranges. Temperature affects dinoflagellate growth most acutely on the seasonal scale. Temperature fluctuation is a strong limiting factor in shallow water, particularly where low temperatures occur. Temperature alone, however, proved unsatisfactory as a limiting boundary condition in the models.

Salinity seems a more powerful control in coastal and near-shore environments. Low and variable near-shore salinity limits dinoflagellates. In other coastal environments dinoflagellates exist in conditions near their upper salinity limit. These upper (seaward) limits model well as range limits. It is difficult, though, to determine dinoflagellate responses to salinity from responses to terrestrially-derived nutrients and compounds.

Nutrient or potential productivity in the model was problematic. Dinoflagellate auxotrophy is complex, and all simplifications quickly became unsatisfactory. Gross potential productivity calculations for phytoplankton are not specific to dinoflagellates (Richards and McGavan, 1989; Kang, 1992). Photosynthetic rates are affected by temperature, turbidity, and diurnal migration patterns. Light plays significant (but not well explained) roles in cueing dinoflagellate activities and life cycle stages. These effects are difficult to model in a meaningful way.

It became obvious during model construction that near-shore dinoflagellates are tied closely to terrestrial nutrient input and terrestrial drainage patterns. Possibly, these terrestrial factors, along with salinity, largely control coastal dinoflagellates. Terrestrial nutrient input is difficult to quantify.

Nutrient control entered the models as a growth-limiting factor, as nitrogen or phosphorus depletion limits bloom conditions. While this does provide limitations on logistic growth in the models, it does not reflect the broader role of nutrients on real dinoflagellates.

When these parameters are integrated, a plausible preliminary model is produced. The models do appear to resemble their natural analog. Still, they lack refinement. Use and adjustment of the models is necessary. Additional parameters should be explored.

## **Results of Q-mode Cluster analysis**

### **Total Data set**

Cluster analysis of the combined fossil data set results in 17 clusters at a coefficient of similarity of 0.4 (figure 13). These groups reflect changes in dinoflagellate associations, controlled largely by environmental conditions. Clustering follows the biostratigraphic order of the samples, representing a shallowing trend from the lower to upper Aquia (from sample 174 to 197), through a shallow and variable period of the uppermost Aquia and lowermost Nanjemoy (samples 303 to 318), and then a deepening trend into the Woodstock Member of the Nanjemoy (samples from 324 to 339 and 239 to 271).

Low association between geographically different data sets occurs between samples 317 and 324, and between 201 and 303.

An *Areoligera*-dominated association occurs during two separate time intervals. Samples 247 to 249 of the Nanjemoy cluster with 174 through 178 of the lower Aquia. Although the two assemblages are similar, they are not statistically the same.

Clusters representing (regressive) Aquia samples are more similar to each other than are clusters of the (transgressive) Nanjemoy samples. This suggests a more set pattern of incumbent species in the regressive samples.



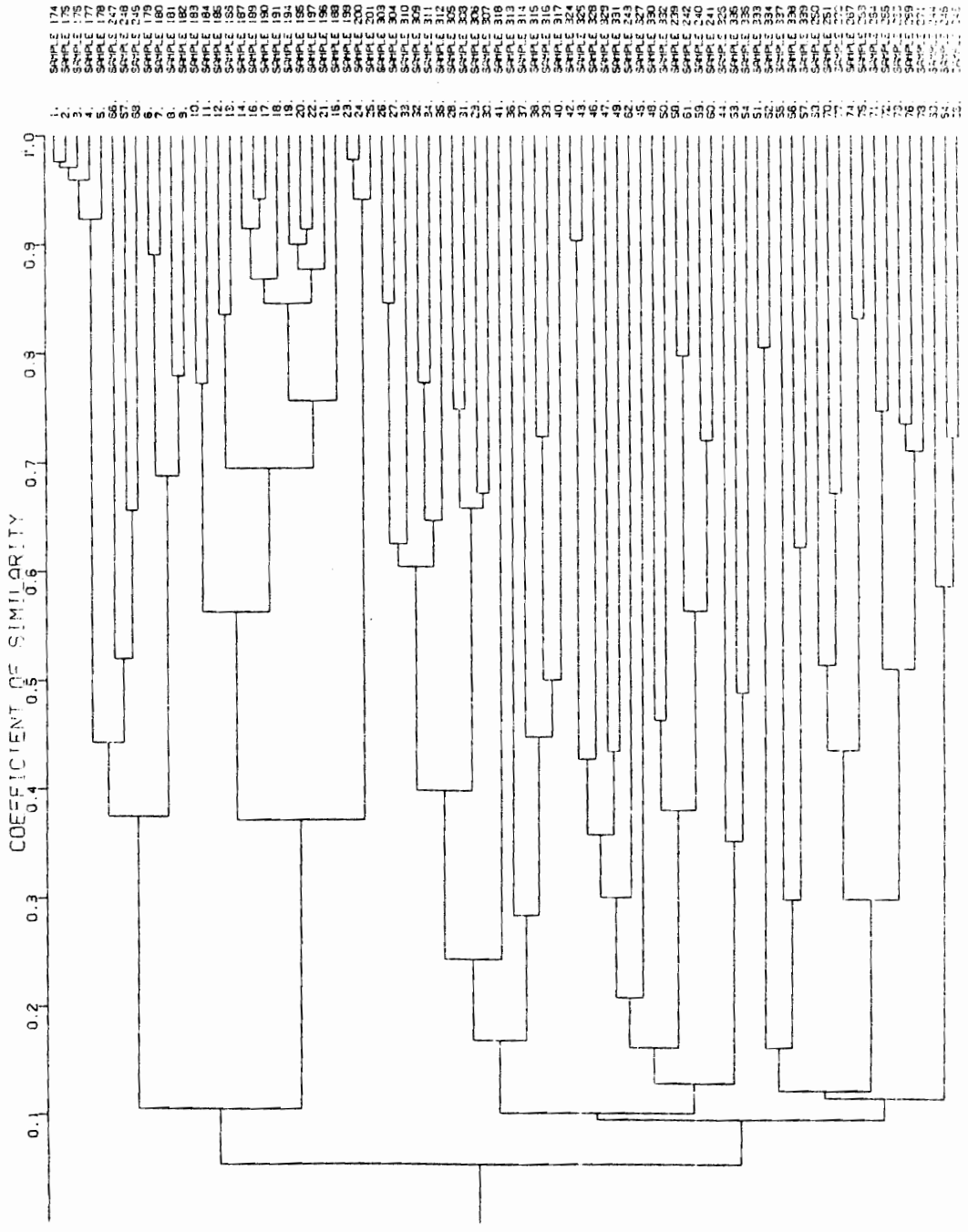


Figure 13. Total Data Set Q-mode Cluster Dendrogram.  
Cluster groups were chosen at the 0.4 similarity coefficient level.

### **Common Species Cluster Associations (>10%)**

The common species subset includes dinoflagellates composing more than 10% of any sample. These represent the largest species biomass, and largely determine the “structure” of the groups.

The Q-mode dendrogram for the common portion of the data set is similar to the pattern of the total data set dendrogram. Again, 17 groupings occur at or above the 0.4 similarity coefficient. Group makeup is similar, though not identical to that of the total data set. (See table 1)

Here, single, dissimilar samples become more apparent (i.e., samples 327, 313). These may represent transition or boundary conditions between larger, unlike groups. *Areoligera*-dominated samples within the Aquia cluster at high levels of similarity, highlighting the role of the most common species in forming the statistical basis of the cluster. Nanjemoy samples appear more evenly clustered between sample sets.

### **Intermediate Species Cluster Associations (1-10%)**

The second subset dendrogram includes intermediate species, defined as representing between 1% and 10% of any sample. Intermediate species occur in an order of magnitude less than the common species.

The intermediate dendrogram does not correspond well structurally to either the total or the common data. Individual species within the intermediate subset do not necessarily follow the boundaries defined by the ranges of more common species. When only clusters at or above the 0.4 similarity coefficient are considered, however, the result is more recognizable. Intermediate frequency species do respond in the same general pattern as the complete system.

### **Rare Species Cluster Associations (<1%)**

Rare species (less than 1% of any given sample) individually have very little statistical impact on the structure or grouping of the samples. The dendrogram for rare species follows environmental or stratigraphic order of the samples. The species within this subset occur in low numbers (by definition) and have low impact. In this dendrogram, the geographical boundaries between sample sets (between samples 201 and 303, and between 339 and 239) are evident. This dendrogram also produces 17 groupings at 0.4 similarity or above. When rare groupings are tabulated against those of common or intermediate sets (table 1), the similarity is apparent.

Table 1 numerical comparison of dendrogram groups

Q-mode Groupings  
(Paleocommunity types)

	All	common	Intermediate	Rare
	174:	174	174	174
	175:	175	175	175
	176:	176	176	176
PT 1	177:	177	177	177
	178:	178	312:	178
	247:	249:	314:	
	248:	179:		
	249:	180:	178:	
		247:		
		248:	181	
		181		179
	179:	182:	179:	180
PT 2	180:		180:	182
	182:		181:	183:
			182:	181
			183:	
			184:	
	183:	183:	316:	184
	184:	184:		185
	185:	185:	185:	
	186:	186:	186:	186
	187:	188:		187
PT 3	189:	187:	187:	188
	190:	195:	189:	189
	191:	189	190:	190
	194:	197	188:	191
	195:	190:	196:	194
	197:	194:	197:	196
	196:	196:	194:	197
	188:	191	195:	195
			191	
			329:	
			251	
			199	
	199:	199		199
PT 4	200:	200	200:	200
	201:	201:		201
			201	
				311
	303:	303:	303:	314
	304:	311	304:	312:
PT 5	310:	304	309:	
	309:	309	310	303
	311:	310	311	304
	312:	312:		309
				306
				307
	305:	305	305	308
PT 6	308:	308:	308	
	306:	306:	306:	305
	307:	307:	307	
	314:	314		315
PT 7	315	315	315	315

Q-mode Groupings  
(Paleocommunity types)

	316	316		
	317	317	317	317
				318
				319
(sample 313)	313	313	313	313
(sample 318)	318	318	318	318
	324	324	324	324
PT 8	325	325	325	325
	328	328		
				326
				327
	329	329	329	328
PT 9	331			329
		331	331	331
	330	330	330	330
PT 10	332			332
			332	
PT 11	333	333		333
	334	334		334
PT 12	335	335	335	335
	336	336		
				336
(sample 337)	337	337	337	337
	338	338	338	338
PT 13	339	339		
				339
		332		
	239	240	239	242
PT 14	242	241	242	240
	240	239	241	241
	241	242	240	243
				246
(sample 243)	243	243	243	245
(sample 247)	247		247	247
				244
	244	244		
PT 15	245	245	245	
	246	246		
			246	
				248
			249	249
				250

Figure 22  
 Q-mode Groupings  
 (Paleocommunity types)

			250	
	250	250	334	
	251	251	266	251
PT 16	270	267	271	
	267	268	336	
	268	266	270	
		270	267	
		269	268	
		271		
	264	264	264	264
PT 17	265	265	265	265
	266			
	269		269	266
	271			267
				268
				269
				270
				271

### Discussion of Paleocommunity Makeup

Q-mode clustering of the data sets all show approximately 17 coherent subgroupings, or paleocommunity types, in addition to individual samples which are consistently dissimilar to other samples. Both of these observations suggest real rather than arbitrary cluster boundaries. These boundaries likely represent changes in the physical environment through time. ANOVA analysis shows that, at the 0.4 similarity level, all grouped samples are statistically similar, but are not the same. Therefore, the cluster groups in this study are paleocommunity types. Only in *Areoligera*-dominated samples of the Aquia, where similarity is much higher between some samples might a paleocommunity occur.

There are 255 dinoflagellate species represented in this study. These species, and their general relative frequencies are shown in table 2. The first column contains a species identification number. The first 176 numbers correspond to species identification numbers assigned by Witmer (1987) to species of the Oak Grove core; succeeding numbers are additional species reported in other data sets.

Table 5 shows that a large number of species only ever occur as rare (R%=100). Such species represent no appreciable impact on community structure at any time. Where only a single cyst is reported across the entire data set, the percentage value does not record its presence (R%=0, S%=0, C%=0).

There are several species which are nearly always common. The most extreme case of this is *Spinidinium macmurdoensis*, which is present as a common and dominant species, or is not present at all. Within this extreme presence-absence pattern, a 'preferential prey' or preservation effect may be in operation. There is a trend of common species as preferentially "near-shore" types (although the case of *Areoligera* appears different), which might reflect bloom behavior.

Other species alternate between rare to intermediate, or from rare to common, in varying proportions.

Table 2. Dinoflagellate Master List and Count Tabulation.

	Count values			Percentage values			>25% values		
	ItI#	R#	S#	C#	R%	S%	C%	VC#	VC%
1: <i>Achieila bifurmoides</i>	25.1	7.2	18.9	0	27.5	72.4	0	0	0
2: <i>Adnatosphaeridium multispinosum</i>	195	3.47	42.3	150	1.77	21.5	76.5	114	58.2
3: <i>Adnatosphaeridium robustum</i>	73.3	1.33	38.5	33.9	1.3	52.3	45.9	0	0
4: <i>Aliscocysta cf. a. marganta</i>	2.67	1	1.67	0	37.5	62.5	0	0	0
5: <i>Andalusieilla rhomboneira</i>	4.33	1.67	2.67	0	38.5	51.5	0	0	0
6: <i>Apectodinium Homomorphum complex</i>	31.7	2.33	21.77	0	2.96	2.45	94.7	3	38
7: <i>Apectodinium sp. a</i>	0	0	0	0	0	0	0	0	0
8: <i>Apectodinium australiense</i>	5.33	0	5.33	0	0	100	0	0	0
9: <i>Apectodinium cribosum</i>	9.73	1.37	5.37	0	21.4	79.3	0	0	0
10: <i>Apectodinium apyrrinthum</i>	0.33	0.33	0	0	100	0	0	0	0
11: <i>Apectodinium retolium</i>	1.67	1.67	0	0	100	0	0	0	0
12: <i>Arcelligera sp.</i>	37.4	3.07	113	758	0.35	12.9	36.8	70	30.2
13: <i>Ascotomocystus hvana</i>	3.33	2.33	6	0	29	72	0	0	0
14: <i>Batacospaera microreticulata</i>	0	0	0	0	0	0	0	0	0
15: <i>Batacospaera sp.</i>	4	3	1	0	75	25	0	0	0
16: <i>Biconidium longissimum</i>	0	0	0	0	0	0	0	0	0
17: <i>Calogonidium amicum</i>	121.7	3	18.7	0	13.8	86.2	0	0	0
18: <i>Cassidium calceonicum</i>	21.7	2.67	2.33	16.7	12.3	10.9	76.9	0	0
19: <i>Chiroptendium partsonarium</i>	0	0	0	0	0	0	0	0	0
20: <i>Chlamydomonella cf. C. urna</i>	1.67	3.23	20.3	128	5.23	13.2	81.5	0	0
21: <i>Cladocyxidium saeptum</i>	4	4	0	0	100	0	0	0	0
22: <i>Cleistosphaeridium diversispinosum</i>	107	4.73	27.4	74.5	4.44	25.7	69.9	0	0
23: <i>Canniximura imbrata</i>	3	3	0	0	100	0	0	0	0
24: <i>Cardosphaeridium biarmatum</i>	2.67	1.67	1	0	52.5	37.5	0	0	0
25: <i>Cardosphaeridium fibrospinosum</i>	12.7	12.7	0	0	100	0	0	0	0
26: <i>Cardosphaeridium giganteum</i>	73.7	1	14	58.7	1.26	19	79.6	0	0
27: <i>Cardosphaeridium gracilis</i>	50.1	9.13	40.9	0	18.2	31.3	0	0	0
28: <i>Cardosphaeridium inodes</i>	31.5	17.7	53.3	0	21.7	78.3	0	0	0
29: <i>Cardosphaeridium inodes robustum</i>	1	1	0	0	100	0	0	0	0
30: <i>Cardosphaeridium multispinosum</i>	2.73	2.73	0	0	100	0	0	0	0
31: <i>Cardosphaeridium solaster</i>	0.33	0.33	0	0	100	0	0	0	0
32: <i>Cardosphaeridium amoulatospinosum</i>	0	0	0	0	0	0	0	0	0
33: <i>Cardosphaeridium calosum</i>	10.8	9.47	1.33	0	87.7	12.3	0	0	0
34: <i>Canea sp.</i>	14.3	1.33	13	0	9.3	90.7	0	0	0
35: <i>Capsulidium pseudocolligerum</i>	0	0	0	0	0	0	0	0	0
36: <i>Deliancia dartmora</i>	54.5	0.67	53.3	0	1.22	98.3	0	0	0
37: <i>Deliancia phosphonica</i>	208	5.27	41.3	160	2.02	20.1	76.9	160	75.3
38: <i>Deliancia wardenensis</i>	3.3	3.3	0	0	100	0	0	0	0
39: <i>Dinocervgium ciadides</i>	7.33	3.33	4	0	45.5	54.5	0	0	0
40: <i>Dinocervgium lenmarnense</i>	3	1	3	0	11.1	38.9	0	0	0
41: <i>Dionyes colligerum</i>	25.5	10.7	15.9	0	40.5	59.5	0	0	0
42: <i>Dionyes colligerum</i>	6	4.67	1.33	0	77.3	22.2	0	0	0
43: <i>Dionyes colligerum (sensu cockson 1985)</i>	5	5	0	0	100	0	0	0	0
44: <i>Distadodinium paradoxum</i>	0	0	0	0	0	0	0	0	0
45: <i>Electrocysta censoaculata</i>	43	7	20.3	15.7	16.3	47.3	36.4	0	0
46: <i>Electrocysta obscurotubulata</i>	137	0	46	38	2.19	33.5	64.2	29	21.2
47: <i>Eociadopyxis penicillatum</i>	242	9.03	20.7	212	3.74	3.56	87.7	60.3	25.2
48: <i>Exochosphaeridium bifidum</i>	4.33	4.33	0	0	100	0	0	0	0
49: <i>Fibracinium annetorbense</i>	181	1.83	10.7	168	1.01	5.9	93.1	64.3	35.5
50: <i>Fibrocysta bioflavore</i>	5	3	2	0	60	40	0	0	0
51: <i>Fibrocysta lappacea</i>	1.33	1.33	0	0	100	0	0	0	0
52: <i>Fibrocysta ocaitoscinosa</i>	2.67	1.67	1	0	52.5	37.5	0	0	0
53: <i>Fibrocysta radiata</i>	4	4	0	0	100	0	0	0	0
54: <i>Fibrocysta sp.</i>	1	1	0	0	100	0	0	0	0
55: <i>Florentinia ferox</i>	11.7	1.67	10	0	14.3	35.7	0	0	0
56: <i>Forma A</i>	4.33	3	1.33	0	69.2	30.8	0	0	0
57: <i>Forma B</i>	25.3	0.33	3.33	16.7	1.22	32.3	65.3	0	0
58: <i>Forma C</i>	6	1	5.67	53.3	1.64	10.3	87.4	0	0
59: <i>Forma D</i>	2	2	0	0	100	0	0	0	0

50	Forma E	0	0	0	0	0	0	0	0	0
51	Fromea fragilis	1	5	5	0	48.5	54.5	0	0	0
52	Fromea? laevigata	0.67	0.67	0	0	100	0	0	0	0
53	Glaucovocysta exuberans coix	322	2	17.7	302	0.62	5.49	93.9	248	77
54	Glaucovocysta pruinata	55.3	0.33	3.67	51.3	0.5	5.63	92.3	21.7	57.2
55	Glaucovocysta sp. A	1.69	1.67	27.0	140	0.99	16.2	82.3	140	82.3
56	Glaucovocysta sp. B	5	0.67	5.33	0	1	38.9	0	0	0
57	Hamiasphaera cf. h. septata	10.67	0.67	0	0	100	0	0	0	0
58	Hamiasphaera septata	1.9	5	12	0	33.3	56.7	0	0	0
59	Heteraulacocysta camoanula	7.5	5.4	2.2	0	71	29.9	0	0	0
70	Homotryblium calicum	1.33	1.33	0	0	100	0	0	0	0
71	Homotryblium tasmaniense	27.9	2.37	25.1	0	10.3	39.7	0	0	0
72	Homotryblium pallidum	57.3	2.33	5.67	48.3	4.07	11.5	84.3	27.3	47.7
73	Horologinella apiculata	1.67	1.67	0	0	100	0	0	0	0
74	Hystriochloidioma esenackii	2	2	0	0	100	0	0	0	0
75	Hystriochloidioma mentum	0.33	0.33	0	0	100	0	0	0	0
76	Hystriochloidioma nquade	18.2	3.2	10	0	45.1	54.9	0	0	0
77	Hystriochloidioma tumescens	5.73	4.43	2.3	0	55.8	34.2	0	0	0
78	Hystriochloidioma umsonum	2.1	2.1	0	0	100	0	0	0	0
79	Hystriochloidiomium cf. h. tubiferum	0.8	0.8	0	0	100	0	0	0	0
80	Hystriochloidiomium spp.	5.33	5.33	0	0	100	0	0	0	0
81	Hystriochloidiomium tubiferum	40.8	13.1	27.7	0	32	68	0	0	0
82	Hystriochloidiomium tubiferum brevispin	1.33	1.33	0	0	100	0	0	0	0
83	Hystriochloidiomium ovum	1.2	1.2	0	0	100	0	0	0	0
84	Imoagidinium cf. h. soeciosum	10.7	0.67	10	0	5.25	33.8	0	0	0
85	Imoagidinium sp. A	0.67	0.67	0	0	100	0	0	0	0
86	Imoagidinium sp. B	0	0	0	0	0	0	0	0	0
87	Imoagidinium soeciosum	3.67	0	3.67	0	0	100	0	0	0
88	Imoetosonaenidium kroemmerbeinii	0.67	0.67	0	0	100	0	0	0	0
89	Imoetosonaenidium rugosum	17.7	15.4	2.33	0	36.8	13.2	0	0	0
90	Imoetosonaenidium? sp. A	0	0	0	0	0	0	0	0	0
91	Inversidinium eximuram	5.2	3.37	2.33	0	52.4	37.5	0	0	0
92	Isaceliidinium cooksoniae	0.33	0.33	0	0	100	0	0	0	0
93	Kailosonaenidium brevisarbatum	37.9	12.3	25.6	0	32.4	67.6	0	0	0
94	Kisseiovia paleotrypta	2.67	1.67	0	0	52.5	37.5	0	0	0
95	Lateuncysta? sp. A	0	0	0	0	0	0	0	0	0
96	Lentonia ruginosa	10.3	3.33	7	0	32.3	57.7	0	0	0
97	Lentonia spinigera	14.3	4.67	9.67	0	32.5	57.4	0	0	0
98	Linguodinium macraeophorum	70.4	12	41.5	16.8	17	59	23.9	0	0
99	Melittosonaenidium pseudocurvatum	1.67	1.67	0	0	100	0	0	0	0
100	Memoranillarnaca leptoderma	12.1	5.8	6.53	0	46.2	53.8	0	0	0
101	Microdinium ornatum	39.9	10.5	11.6	57.9	11.5	12.9	75.4	55.8	52.1
102	Millicodinium gusevii maior	10	4	5	0	40	50	0	0	0
103	Muratodinium limonatum	11.5	4.33	30.5	30.5	3.75	26.5	69.7	30.5	69.7
104	Nematosonaeropsis cf. h. pertusa	0.73	0.73	0	0	100	0	0	0	0
105	Nematosonaeropsis pertusa	5.33	4.67	1.67	0	73.7	26.3	0	0	0
106	Nematosonaeropsis pulvisosa	1.33	0.33	1	0	25	75	0	0	0
107	Nematosonaeropsis rabeculata	16.3	3	3.33	0	49	51	0	0	0
108	Oligosonaeridium complex	3.33	1.33	2	0	40	60	0	0	0
109	Oercuodinium brevispinosum	2.67	1.67	1	0	52.5	37.5	0	0	0
110	Oercuodinium centrocarum	25.6	17	9.6	0	53.9	36.1	0	0	0
111	Oercuodinium cf. d. brevispinosum	0.33	0.33	0	0	100	0	0	0	0
112	Oercuodinium israelianum	10.4	2.4	3	0	23.1	76.9	0	0	0
113	Oercuodinium multispinosum	5.33	1.67	3.67	0	31.3	68.7	0	0	0
114	Oercuodinium vanespinosum	0	0	0	0	0	0	0	0	0
115	Paiaeocystodinium colzowense	29.8	14.2	14.6	0	49.4	50.6	0	0	0
116	Paiaeocystodinium pyrochorum	1.9	3.67	18.3	0	3.5	96.5	0	0	0
117	Paralecniella indentata	68.2	4.63	48.6	15	57.9	71.2	22	0	2
118	Paucosonaeridium inversibocanum	0	0	0	0	100	0	0	0	0
119	Pentadinium abactum granulatum	0	0	0	0	0	0	0	0	0
120	Pheodinium magnificum	5	3.67	1.33	0	73.3	26.7	0	0	0



21	Phthanocerium cf. p. ritonum	0.30	0.30	0	0	100	0	0	0	0
22	Phthanocerium echinatum	0	0	0	0	33.3	56.7	0	0	0
23	Polysphaeridium cf. p. zoharyi	1.67	1.67	0	0	100	0	0	0	0
24	Polysphaeridium zoharyi	4	2.33	1.67	0	58.3	41.7	0	0	0
25	Renidium membraniferum	6.33	6.33	0	0	34.2	15.8	0	0	0
26	Renidium sp. A	0	0	0	0	100	0	0	0	0
27	Rothesia corussica	6.33	4.33	0	0	31.3	18.3	0	0	0
29	Samiandia chlamydochora	2.2	2.2	0	0	100	0	0	0	0
29	Samiandia reticulifera	6.33	1.33	5	0	21.1	78.9	0	0	0
30	Senecalinium? obscurum	10.3	0	7.33	0	29	71	0	0	0
31	Senecalinium? asymmetricum	1.33	1.33	0	0	100	0	0	0	0
32	Senecalinium? diivynense	49.7	6.67	12.3	31.7	11.4	24.8	63.8	0	0
33	Soinidium essoi	1.53	1.53	0	0	100	0	0	0	0
34	Soinidium macmuroense	72.7	0	0	72.7	0	0	100	72.7	100
35	Soinidium paratacuiatum	13.3	0.67	12.7	0	5	95	0	0	0
36	Soinifertes crassioeilis A	11.6	9.3	2.33	0	79.9	20.1	0	0	0
37	Soinifertes crassioeilis B	2	0.67	1.33	0	33.3	66.7	0	0	0
38	Soinifertes crassioeilis C	4.47	3.27	1.2	0	73.1	26.9	0	0	0
39	Soinifertes cf. s. pterotus	0.33	0.33	0	0	100	0	0	0	0
40	Soinifertes anguatus	0.67	0.67	0	0	100	0	0	0	0
41	Soinifertes comutus A	1.33	1.33	0	0	100	0	0	0	0
42	Soinifertes comutus B	0	0	0	0	0	0	0	0	0
43	Soinifertes comutus C	0	0	0	0	100	0	0	0	0
44	Soinifertes mirapellis	0	0	0	0	0	0	0	0	0
45	Soinifertes monius	2.3	2.9	0	0	100	0	0	0	0
46	Soinifertes pseudofurcatus	11.3	11.3	0	0	100	0	0	0	0
47	Soinifertes ramosus subsp. granomemorana	4.33	4.33	0	0	100	0	0	0	0
48	Soinifertes ramosus subsp. granosus	3.97	9.97	0	0	100	0	0	0	0
49	Soinifertes ramosus subsp. membranaceus	6.67	6.67	0	0	100	0	0	0	0
50	Soinifertes ramosus subsp. multibravis	6.47	6.47	0	0	34.5	15.5	0	0	0
51	Soinifertes ramosus subsp. ramosus	72.4	15.6	43.8	13	21.5	60.5	18	0	0
52	Soinifertes ramuiferus	0.67	0.67	0	0	100	0	0	0	0
53	Soinifertes A	0	0	0	0	100	0	0	0	0
54	Soinifertes B	0	0	0	0	0	0	0	0	0
55	Soinifertes C	0	0	0	0	0	0	0	0	0
56	Systematocnora biacantha	51.7	1.67	2.33	47.7	3.23	4.52	92.3	47.7	92.3
57	Tanvosphaeridium variegatum	2	0	0	0	50	50	0	0	0
58	Tectacodinium cellitum	12.4	6.67	6.73	0	45.7	54.3	0	0	0
59	Thalassionora delicata	39.6	5	3.3	26.3	12.6	20.9	66.4	26.3	66.4
60	Thalassionora celagica	205	4.33	50	151	2.1	24.4	73.5	106	51.6
61	Trichodinium nirsuum	1.33	1.33	0	0	100	0	0	0	0
62	Trigonocoxidia gineila	0	0	0	0	100	0	0	0	0
63	Tuberculodinium vancampore	0	0	0	0	0	0	0	0	0
64	Tuberculodinium sp.	0	0	0	0	0	0	0	0	0
65	Tubidermodinium sulcatum	7.13	7.13	0	0	100	0	0	0	0
66	Turbiosphaera villosa	51.6	0.67	21.6	39.3	1.08	3.5	63.9	29.3	63.9
67	Turbiosphaera parataculata	3.33	0.67	7.67	0	3	92	0	0	0
68	Turbiosphaera rotunda	3.33	0.33	3	0	10	90	0	0	0
69	Wetzeliella cf. w. rtyshensis	1.37	1.97	0	0	100	0	0	0	0
70	Wetzeliella namodenensis	1.31	0.73	54.3	72.2	2.86	41.9	55.2	38	29.1
71	Wetzeliella unaris	466	3.33	33.5	429	1.72	7.8	92.1	406	87.1
72	Wetzeliella samancica	296	1.93	16.3	277	0.65	3.5	93.8	277	93.8
73	Wetzeliella sp. A	2.33	0	2.33	0	0	100	0	0	0
74	Wetzeliella variegatituda	12	0	1.67	10.3	0	13.9	36.1	0	0
75	Wilsonidium abuiatum	46.3	2	0	44.3	4.32	0	95.7	44.3	95.7
76	Xanikoon australis	74.6	2.13	38.1	34.3	2.86	51.1	46	0	0
77	Cycoosella elliotica	0	0	0	0	0	0	0	0	0
78	Cycoosella vieta	17.9	6.93	1	0	38.7	51.3	0	0	0
79	Micrvstridium cf. m. ragicie	4	4	0	0	100	0	0	0	0
80	Micrvstridium cf. m. variabile	27	3.67	23.3	0	13.6	36.4	0	0	0
81	Micrvstridium spp.	22.2	17.2	5	0	77.4	22.6	0	0	0

132	<i>Cymatosphaera</i> spp.	5.5	6.5	0	0	100	0	0	0	0
133	<i>Palamocetes</i> sp. A	2.67	1.33	1.33	0	50	50	0	0	0
134	<i>Palamocetes</i> sp. B	4.33	3.33	1	0	76.9	23.1	0	0	0
135	<i>Palamocetes</i> sp. C	21.0	2.33	9	10	10.9	42.2	46.9	0	0
136	<i>Pterospermopsis</i> spp.	15.2	14.2	1	0	93.4	6.58	0	0	0
137	<i>Agreodinium incompositum</i>	1	1	0	0	100	0	0	0	0
138	<i>Auria erinacea</i>	1.5	1.5	0	0	100	0	0	0	0
139	<i>Dionyesopsis buccinata</i>	4.3	3.5	1.2	0	7.5	25	0	0	0
140	<i>Dionyesopsis capitata</i>	9.8	9.8	0	0	100	0	0	0	0
141	<i>Gonyaulacysta guiseppi</i> subsp. major	1.3	2.6	10.4	0	20	30	0	0	0
142	<i>Hydrasphaera marylandense</i>	1	1	0	0	100	0	0	0	0
143	<i>Hystriochlooidium suberosa</i>	1	1	0	0	100	0	0	0	0
144	<i>Hystriochlooidium borussica</i>	2.2	2.2	0	0	100	0	0	0	0
145	<i>Lantemosphaeridium bicolor</i>	5.2	5.2	0	0	100	0	0	0	0
146	<i>Lantemosphaeridium lanosum</i>	10.1	0.2	5.8	43.1	0.2	57.3	42.5	0	0
147	<i>Lantemosphaeridium lappaceum</i>	1.2	1.2	0	0	100	0	0	0	0
148	<i>Lantemosphaeridium pilatum</i>	0.4	0.4	0	0	100	0	0	0	0
149	<i>Lantemosphaeridium radiatum</i>	1	1	0	0	100	0	0	0	0
200	<i>Coerciodinium amiculum</i>	5.8	3.5	2.2	0	52.1	37.9	0	0	0
201	<i>Coerciodinium potomacense</i>	11.4	1.3	9.5	0	15.3	84.2	0	0	0
202	<i>Scribneridium</i> (S.) australiense	24.2	1.2	23	0	4.36	95	0	0	0
203	<i>Soiniferites bulboideus</i>	0.2	0.2	0	0	100	0	0	0	0
204	<i>Soiniferites</i> cf. <i>S. crassipellis</i>	1.3	1.3	0	0	100	0	0	0	0
205	<i>Soiniferites hypercanthus</i>	1.5	1.5	0	0	100	0	0	0	0
206	<i>Soiniferites membranaceus</i>	1.6	1.6	0	0	100	0	0	0	0
207	<i>Soiniferites</i> cf. <i>S. ramosus</i> subsp. ramosus	1.3	1.3	0	0	100	0	0	0	0
208	<i>Soiniferites scaratus</i>	0.4	0.4	0	0	100	0	0	0	0
209	<i>Soiniferites supparus</i>	4.1	1	40	0	2.44	97.5	0	0	0
210	<i>Tectatodinium osilatum</i>	4.2	1.2	3	0	29.5	71.4	0	0	0
211	<i>Tubulosphaeridium pseudocurvatum</i>	1.4	1.4	0	0	100	0	0	0	0
212	<i>Deflandra asymmetrica</i>	9.2	2.3	6.4	0	30.4	69.5	0	0	0
213	<i>Deflandra dihyvensis</i>	98.7	3.4	1.6	33.7	3.44	1.62	94.9	93.7	94.9
214	<i>Deflandra macmurdoensis</i>	12	1	6.2	114	0.33	5.12	94	91	75.2
215	<i>Phthanooperidium resistente</i>	25.5	3.4	10.5	11.5	13.3	41.4	45.3	0	0
216	<i>Soinidium cilineatum</i>	239	1.2	0	237	0.5	0	99.5	21.7	91.1
217	<i>Soinidium rotundum</i>	0.4	0.4	0	0	100	0	0	0	0
218	<i>Wetzeliella</i> (Rhombodinium) <i>glabra</i>	0.2	0.2	0	0	100	0	0	0	0
219	<i>Wetzeliella</i> (W.) <i>articulata</i>	1.4	1.4	0	0	100	0	0	0	0
220	<i>Wetzeliella</i> (W.) <i>coarctata</i>	7.2	0.6	5.5	0	3.33	91.7	0	0	0
221	<i>Wetzeliella</i> (W.) <i>coleothrypta</i>	15.2	0.3	12.3	21.5	2.27	36.4	51.4	0	0
222	<i>Adnatosphaeridium vitatum</i>	4.5	3.5	1	0	78.3	21.7	0	0	0
223	<i>Canningia minor</i>	0.2	0.2	0	0	100	0	0	0	0
224	<i>Cannosphaeropsis pusuosa</i>	1.4	1.4	0	0	100	0	0	0	0
225	<i>Chlamycoconrella rauca</i>	0.2	0.2	0	0	100	0	0	0	0
226	<i>Cycloneocheilium incutum</i>	1	1	0	0	100	0	0	0	0
227	<i>Cycloneocheilium lemniscatum</i>	2.5	1.2	1.4	0	46.2	53.8	0	0	0
228	<i>Dinopteridium cladoides</i>	4	4	0	0	100	0	0	0	0
229	<i>Eisenackia scroiculata</i>	0.3	0.3	0	0	100	0	0	0	0
230	<i>Fusidium tabulatum</i>	11.7	1.2	5.2	110	1.33	4.45	94.5	92.5	79.0
231	<i>Hemicystodinium aculeatum</i>	4.6	1	3.5	0	21.7	78.3	0	0	0
232	<i>Hemicystodinium zoharyi</i>	3.5	3.5	0	0	100	0	0	0	0
233	<i>Heteraulacysta lemnamensis</i>	5.5	5.4	1.2	0	31.3	18.2	0	0	0
234	<i>Homotrydium alium</i>	1.2	1.2	0	0	100	0	0	0	0
235	<i>Impietosphaeridium transtodum</i>	5	5	0	0	100	0	0	0	0
236	<i>Litosphaeridium inversibucanum</i>	1.5	1.5	0	0	100	0	0	0	0
237	<i>Sytematocora ancylrea</i>	5.5	1.4	4.2	0	25	75	0	0	0
238	<i>Aoteocidium baculatum</i>	1.1	1.1	0	0	100	0	0	0	0
239	<i>Aquasphaeridium</i>	2	0	2	0	0	100	0	0	0
240	<i>Carcosphaeridium marylandense</i>	4	4	0	0	100	0	0	0	0
241	<i>Carcosphaeridium species A</i>	1.5	1.5	0	0	100	0	0	0	0
242	<i>Deflandra</i> cf. <i>D. magna</i>	3.5	2.5	1	0	59.4	30.5	0	0	0

243	<i>Deilanthra obscura</i>	3	5.7	2.3	0	7.0	29.3	0	0	0
244	<i>Deilanthra rhomboides</i>	0.5	0.5	0	0	1.0	0	0	0	0
245	<i>Deilanthra ruginosa</i>	3	3	0	0	6.0	0	0	0	0
246	<i>Deilanthra spiridera</i>	20.4	2.3	17.5	0	13.7	86.3	0	0	0
247	<i>Leptocinium virginianum</i>	53.0	0	20.4	39.3	4.9	32.2	62.3	0	0
248	<i>Membranoscraera labiata</i>	25.1	4.1	11.0	23.6	1.53	4.5	93.9	203	30.5
249	<i>Nematoscraecopsis trabeculata</i>	10.9	5.4	4.4	0	59.0	40.7	0	0	0
250	<i>Paraeostomacystus fragilis</i>	1.4	2	1.2	0	14.3	35.7	0	0	0
251	<i>Soinifentes cinctus subsp. reticulatus</i>	1.5	1.5	0	0	1.0	0	0	0	0
252	<i>Soinifentes seotatus</i>	10.9	9.3	1.6	0	35.0	14.7	0	0	0
253	<i>Soinifentes cf. S. seotatus</i>	10.9	9.3	1.6	0	35.0	14.7	0	0	0
254	<i>Vernacium spp.</i>	0.5	0.5	0	0	1.0	0	0	0	0
255	<i>Wetziella (W.) homomorpha subsp. zingib</i>	549	0	4.1	545	0	0.53	99.4	545	39.4
Total										
		3125	779	1802	5544					

### **Results of Chosen Phase Plane Plots**

Phase plane plots were produced for a number of species pairs from the data. Four phase plane plots, of common species, have been selected as representative (figure 14). Results of the phase plane plots are difficult to interpret. Although loops about a possible equilibrium occur, it is usually when one species is replacing another. The periods of overlap, on this scale, are short, and seldom repeat.

When replacement occurs more than once in the data set, similar loop figures occur, but are not centered around the same equilibrium point. A feature of these phase plane diagrams is a strong trend roughly perpendicular to the equilibrium axis, rather than parallel to it. This trend is expected when both species are affected by some outside factor more than by interaction with each other. On the scale of this data, the outside influence appears to parallel local environmental shifts. Although phase plane analysis does show potential for finding interaction patterns in fossil data, both environmental overprint, and data resolution on this scale limit its effectiveness.

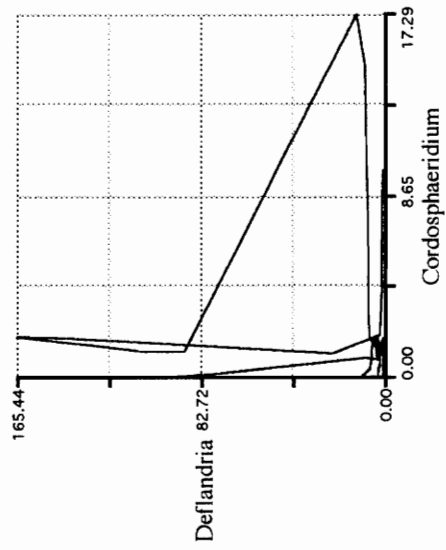
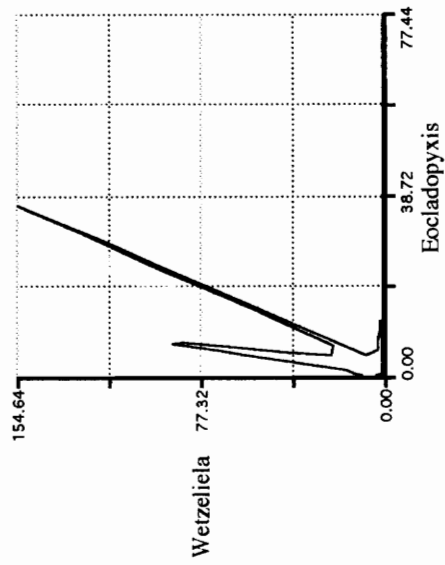
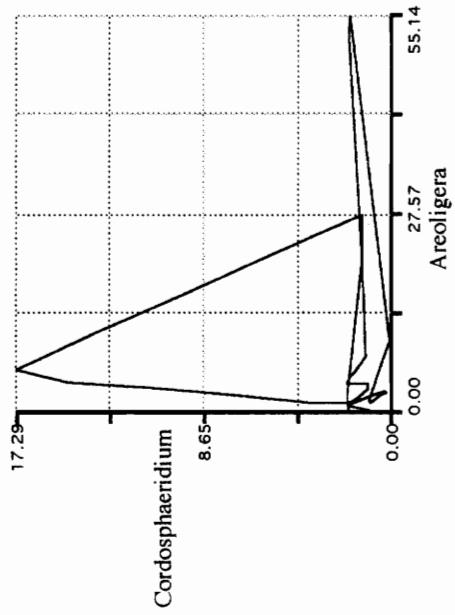
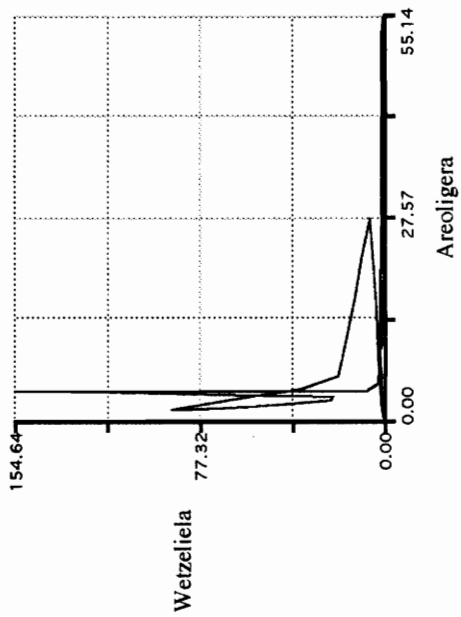


Figure 14.. Phase Plane Examples

## VII. CONCLUSIONS

The following conclusions are based on the dynamic systems modeling, and statistical analysis of fossil dinoflagellates studied during this project.

1. Dynamic systems modeling is directly applicable to both modern and fossil coastal dinoflagellate analysis. Success of modeling however depends on refinement, and on resolution of the input data. The models used in this study are preliminary.
2. Coastal dinoflagellate dynamics can be approximated using parameters of temperature, salinity, and limiting nutrient. Other parameters are necessary for model refinement.
3. The models suggest that differences in boundaries between watermasses might be discernible in dinoflagellate associations in rock or sediment. The fossil data appear consistent with this, at least within the resolution of the samples.
4. Cluster analysis of the Salisbury fossil data indicates that cluster boundaries produced are not gradient artifacts, but represent real changes in dinoflagellate associations. These parallel changes in sediment (watermass) characteristics.
5. Cluster analysis indicates 17 separate dinoflagellate associations which can be interpreted individually as occurring in estuarine, nearshore, or offshore conditions. Individual dissimilar samples between groups may represent boundary or transitional environments.
6. Samples in cluster groups are statistically similar, but not the same, and represent paleocommunity types rather than paleocommunities. Individual species reccur in similar environmental settings through time, but paleocommunity types do not repeat within the Pamunky Group data.
7. Circulation patterns of the Salisbury Embayment during the Tertiary may have resembled those of the modern Yellow Sea, with both lateral fronts, and regions of mixing. Fossil dinoflagellate data and modeling suggest that water conditions of the Salisbury Embayment may have been warmer than the modern Yellow Sea.

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**APPENDIX A:**

**TRANSFORMED DINOFLAGELLATE DATA SETS  
ANOVA ANALYSIS OF DINOFLAGELLATE DATA SETS**



PT1 "Areoligera" Association

174	174	witmer aquia ml	175	175	witmer aquia ml	176	176	witmer aquia ml
39.30	12	Areoligera sp.	37.00	12	Areoligera sp.	39.00	12	Areoligera sp.
0.20	156	Turbosphaera filosa	0.30	166	Turbosphaera filosa	0.40	166	Turbosphaera filosa
0.20	117	Paralecanella indentata	0.20	117	Paralecanella indentata	0.30	176	Xenikoon australis
0.20	176	Xenikoon australis	0.20	176	Xenikoon australis	0.20	159	Thalassoonora delicata
0.30	49	Fibradinum annetorpense	0.30	49	Fibradinum annetorpense	0.30	117	Paralecanella indentata
0.30	248	Membranosphaera tabulata	0.70	248	Membranosphaera tabulata	0.60	248	Membranosphaera tabulata
0.30	181	Micrystidium spp.	0.70	181	Micrystidium spp.	0.40	181	Micrystidium spp.
0.40	101	Microdinium ornatum	0.60	159	Thalassoonora delicata	0.40	186	Pterospermopsis spp.
0.40	159	Thalassoonora delicata	0.40	39	Impileosphaeridium rugosum	0.40	151	Spiriferites ramosus subsp. ramosus
0.30	39	Impileosphaeridium rugosum	0.40	247	Legodinium virginianum	0.30	20	Chlamydonoreia cf. C. juma
0.20	20	Chlamydonoreia cf. C. juma	0.30	192	Cymatosphaera spp.	0.30	247	Legodinium virginianum
0.20	82	Cymatosphaera spp.	0.30	246	Dellandna springera	0.30	101	Microdinium ornatum
0.20	246	Dellandna springera	0.30	101	Microdinium ornatum	0.20	178	Cyclopsiella vieta
0.20	89	Diphyesopsis buccinata	0.30	186	Pterospermopsis spp.	0.20	242	Dellandna cf. D. magnifica
0.20	110	Oercuodinium centrocarpum	0.20	20	Chlamydonoreia cf. C. juma	0.20	246	Dellandna springera
0.20	151	Spiriferites ramosus subsp. ramosus	0.20	242	Dellandna cf. D. magnifica	0.20	189	Diphyesopsis buccinata
0.10	242	Dellandna cf. D. magnifica	0.20	243	Dellandna obscura	0.20	190	Diphyesopsis capitata
0.10	243	Dellandna obscura	0.20	190	Diphyesopsis capitata	0.20	39	Impileosphaeridium rugosum
0.10	190	Diphyesopsis capitata	0.20	110	Oercuodinium centrocarpum	0.20	115	Palaeocystodinium goizowense
0.10	31	Hystriochosphaeridium rubiferum	0.20	115	Palaeocystodinium goizowense	0.20	148	Spiriferites pseudofurcatus
0.10	249	Nematosphaeropsis trabeculata	0.20	151	Spiriferites ramosus subsp. ramosus	0.20	254	Verrucium spp.
0.10	115	Palaeocystodinium goizowense	0.10	189	Diphyesopsis buccinata	0.10	243	Dellandna obscura
			0.10	31	Hystriochosphaeridium rubiferum	0.10	31	Hystriochosphaeridium rubiferum
			0.10	249	Nematosphaeropsis trabeculata	0.10	249	Nematosphaeropsis trabeculata
			0.10	110	Oercuodinium centrocarpum			

177	177	Wimmer aquia ml	178	178	Wimmer aquia ml
36.50	12	Areoligera sp.	32.10	12	Areoligera sp.
1.50	55	Turbosphaera filosa	2.00	29	Aquasphaeridium
1.50	59	Thalassiochorda cecata	1.30	166	Turbosphaera filosa
1.50	117	Paralecaniella indentata	1.50	159	Thalassiochorda cecata
1.20	248	Membranosphaera labiata	1.30	246	Deliantha spinigera
1.20	175	Xenikoon austrais	1.20	117	Paralecaniella indentata
2.30	81	Micrystidium spp.	1.10	242	Deliantha cf. D. magnifica
2.50	36	Pterospoermopsis spp.	2.30	181	Micrystidium spp.
2.50	101	Microdinium ornatum	2.30	176	Xenikoon austrais
2.40	22	Chlamydoclonella cf. C. juma	2.50	196	Pterospoermopsis spp.
2.40	242	Deliantha cf. D. magnifica	2.50	151	Spiriferites ramosus subsp. ramosus
2.40	246	Deliantha spinigera	2.40	190	Dionysososis capitata
2.30	90	Dionysososis capitata	2.40	99	Impatiensphaeridium rugosum
2.30	31	Hystichosphaeridium tubiferum	2.40	33	Kaliosphaeridium brevibarbatum
2.30	39	Impatiensphaeridium rugosum	2.40	248	Membranosphaera labiata
2.30	33	Kaliosphaeridium brevibarbatum	2.40	101	Microdinium ornatum
2.20	51	Spiriferites ramosus subsp. ramosus	2.40	151	Palaeocystodinium gorizowense
2.20	243	Deliantha obscura	2.30	201	Chlamydoclonella cf. C. juma
2.20	249	Nematosphaeropsis trabeolata	2.30	178	Cyclopsella vieta
2.20	110	Coercuodinium centrocarum	2.30	243	Deliantha obscura
2.20	115	Palaeocystodinium gorizowense	2.30	249	Nematosphaeropsis trabeolata
2.10	89	Dionysososis buccinata	2.30	250	Palaeostomacystus fragilis
2.10	146	Spiriferites pseudofurcatus	2.20	189	Dionysososis buccinata
			2.20	31	Hystichosphaeridium tubiferum
			2.20	110	Coercuodinium centrocarum
			2.10	146	Spiriferites pseudofurcatus



Corellation matrix for PT1

Correlation Matrix for Variables: X1 ... X5

	174	175	176	177	178
174	1				
175	.99996	1			
176	.99987	.99987	1		
177	.99979	.99983	.99991	1	
178	.99932	.99936	.99943	.99952	1

Anova table for PT1

One Factor ANOVA-Repeated Measures for X1 ... X5

Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	37545.0858	147.8153	3494.61349	.0001
Within subjects	1020	43.144	.0423		
treatments	4	.00282	.00071	.01662	.9995
residual	1016	43.14118	.04246		
Total	1274	37588.2298			

Reliability Estimates for- All treatments: .99971    Single Treatment: .99857

PT2 "Areoligera/Eocladopyxis peniculatum " Association

0.00	180	0.00	181
49.50	12: Areoligera sp.	47.50	12: Areoligera sp.
20.50	47: Eocladopyxis peniculatum	21.50	47: Eocladopyxis peniculatum
6.50	96: Deliantha carmona	7.50	96: Lantermosphaeridium anosum
4.50	3: Adnatosphaeridium robustum	7.30	3: Adnatosphaeridium robustum
4.50	29: Corcosphaeridium inodes	4.10	29: Corcosphaeridium inodes
4.40	96: Lantermosphaeridium anosum	2.50	247: Leptodinium virginianum
3.70	37: Deliantha phosonortica	1.30	36: Deliantha carmona
3.30	33: Kaiiosphaeridium breviarbatum	1.30	33: Kaiiosphaeridium breviarbatum
3.30	247: Leptodinium virginianum	1.70	37: Deliantha phosonortica
3.30	248: Membranosphaera tabulata	0.90	253: Spiniferites cf. S. septatus
3.40	81: Micrystidium spp.	0.90	252: Spiniferites septatus
3.30	243: Deliantha obscura	0.30	248: Membranosphaera tabulata
3.30	190: Diphyesopsis capitata	0.30	81: Micrystidium spp.
3.30	250: Palaeostomacystus fragilis	0.30	249: Nematosphaeropsis trabeculata
3.30	117: Paracaniella indentata	0.20	20: Chlamydomonella cf. C. juma
3.20	228: Aptrodinium baculatum	0.20	241: Corcosphaeridium species A
3.20	39: Impletosphaeridium rugosum	0.20	241: Corcosphaeridium species A
3.20	115: Palaeocystodinium goizowense	0.20	242: Deliantha cf. D. magnifica
3.20	253: Spiniferites cf. S. septatus	0.20	190: Diphyesopsis capitata
3.20	141: Spiniferites cornutus A	0.20	38: Linguodinium macnaerophorum
3.20	136: Spiniferites crassioeilis A	0.20	248: Membranosphaera tabulata
3.20	252: Spiniferites septatus	0.10	250: Palaeostomacystus fragilis
3.10	20: Chlamydomonella cf. C. juma	0.10	238: Aptrodinium baculatum
3.10	241: Corcosphaeridium species A	0.10	178: Cyclosetella vieta
3.10	242: Deliantha cf. D. magnifica	0.10	39: Impletosphaeridium rugosum
3.10	110: Operculodinium centrocarpum	0.10	38: Linguodinium macnaerophorum
3.10	251: Spiniferites angulatus subsp. reticulatus	0.10	110: Operculodinium centrocarpum
3.10	106: Spiniferites cornutus A	0.10	115: Palaeocystodinium goizowense
3.10	136: Spiniferites crassioeilis A	0.10	141: Spiniferites cornutus A
3.10	254: Verrucium spp.	0.10	136: Spiniferites crassioeilis A
		0.10	254: Verrucium spp.



Corellation Matrix for PT2

Correlation Matrix for Variables: X<sub>1</sub> ... X<sub>4</sub>

	179	180	181	182
179	1			
180	.99113	1		
181	.88925	.92817	1	
182	.70533	.75445	.90491	1

ANOVA Table for PT2

One Factor ANOVA-Repeated Measures for X<sub>1</sub> ... X<sub>4</sub>

Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	8488.78035	33.4204	24.50405	.0001
Within subjects	765	1043.3625	1.36387		
treatments	3	.04097	.01366	.00997	.9986
residual	762	1043.32153	1.36919		
Total	1019	9532.14285			

Reliability Estimates for- All treatments: .95919    Single Treatment: .85457

PT3 "Wetzeliiella (W.) homomorpha/ Membranosphaera tabulata/

Eocladopyxis peniculatum" Association

:801	:841
01:301 255 Wetzeliiella (W.) homomorpha subsp. quinq.	49:801 255 Wetzeliiella (W.) homomorpha subsp. quinq.
22:401 47 Eocladopyxis peniculatum	14:601 47 Eocladopyxis peniculatum
17:301 247 Laetodinium virginianum	10:501 247 Laetodinium virginianum
10:501 12 Arcoigera sp.	10:301 12 Arcoigera sp.
5:101 196 Lantimosphaeridium lanosum	2:501 196 Lantimosphaeridium lanosum
3:701 28 Corcosphaeridium inodes	1:701 33 Kalkosphaeridium brevisarbatum
1:301 93 Kalkosphaeridium brevisarbatum	1:501 28 Corcosphaeridium inodes
1:301 31 Aonatosphaeridium roostum	1:201 31 Aonatosphaeridium roostum
0:801 37 Dellandina onosporitica	1:101 249 Nematosphaeroopsis trabeculata
1:701 253 Spiniferites cf. S. septatus	0:901 190 Diphyesopsis capitata
0:701 252 Spiniferites septatus	0:701 311 Hystichosphaeridium tubiferum
0:501 249 Nematosphaeroopsis trabeculata	0:501 181 Microtyndium spp.
0:401 190 Diphyesopsis capitata	0:401 371 Dellandina onosporitica
0:301 241 Corcosphaeridium species A	0:401 101 Microdinium ornatum
0:301 311 Hystichosphaeridium tubiferum	0:401 115 Palaecocystodinium goizowense
0:301 151 Spiniferites ramosus subsp. ramosus	0:301 242 Dellandina cf. D. magnifica
0:201 201 Chlamydomonadea cf. C. urna	0:301 243 Dellandina obscura
0:201 243 Dellandina obscura	0:301 246 Dellandina springera
0:201 246 Dellandina springera	0:301 248 Membranosphaera tabulata
0:201 248 Membranosphaera tabulata	0:301 151 Spiniferites ramosus subsp. ramosus
0:201 110 Oercciodinium constrictum	0:201 89 Impietosphaeridium rugosum
0:201 250 Palaecostomacystis family	0:201 98 Linguodinium macrosporum
0:201 141 Spiniferites comutus A	0:201 253 Spiniferites cf. S. septatus
0:101 178 Cyclopsiella vieta	0:201 251 Spiniferites cingulatus subsp. reticulatus
0:101 242 Dellandina cf. D. magnifica	0:201 252 Spiniferites septatus
0:101 245 Dellandina ruginosa	0:101 201 Chlamydomonadea cf. C. urna
0:101 194 Hystichosphaeroopsis borussica	0:101 240 Corcosphaeridium marylandense
0:101 391 Impietosphaeridium rugosum	0:101 241 Corcosphaeridium species A
0:101 38 Linguodinium macrosporum	0:101 179 Cyclopsiella vieta
0:101 115 Palaecocystodinium goizowense	0:101 245 Dellandina ruginosa
0:101 186 Perospermopsis spp.	0:101 194 Hystichosphaeroopsis borussica
0:101 251 Spiniferites cingulatus subsp. reticulatus	0:101 110 Oercciodinium constrictum
0:101 136 Spiniferites crassipellis A	0:101 136 Spiniferites crassipellis A
0:101 254 Verhucium spp.	0:101 148 Spiniferites ramosus subsp. granosus
0:101 169 Wetzeliiella cf. W. rhyssensis	



195:	196:	187:
54.30: 255: <i>Weitzelia</i> (W.) <i>homomorpha</i> subsp. <i>quind</i>	58.00: 255: <i>Weitzelia</i> (W.) <i>homomorpha</i> subsp. <i>quind</i>	25.40: 255: <i>Weitzelia</i> (W.) <i>homomorpha</i> subsp. <i>quind</i>
2.00: 247: <i>Leptodinium</i> <i>virginianum</i>	3.70: 248: <i>Membranosphaera</i> <i>tabulata</i>	25.80: 248: <i>Membranosphaera</i> <i>tabulata</i>
3.10: 2: <i>Areoligera</i> sp.	7.30: 247: <i>Leptodinium</i> <i>virginianum</i>	16.10: 20: <i>Chlamydomonada</i> cf. <i>C. jma</i>
5.30: 31: <i>Hyalinosphaeridium</i> <i>tubiferum</i>	5.80: 12: <i>Areoligera</i> sp.	5.80: 12: <i>Areoligera</i> sp.
2.30: 196: <i>Lammosphaeridium</i> <i>lanosum</i>	5.20: 20: <i>Chlamydomonada</i> cf. <i>C. jma</i>	2.40: 196: <i>Lammosphaeridium</i> <i>lanosum</i>
1.60: 253: <i>Spiriferites</i> cf. <i>S. septatus</i>	2.00: 196: <i>Lammosphaeridium</i> <i>lanosum</i>	2.20: 248: <i>Deilandra</i> <i>spinigera</i>
1.60: 252: <i>Spiriferites</i> <i>septatus</i>	1.70: 248: <i>Deilandra</i> <i>spinigera</i>	1.70: 247: <i>Leptodinium</i> <i>virginianum</i>
1.50: 33: <i>Kaliosphaeridium</i> <i>brevibarbatum</i>	1.20: 115: <i>Palaeocystodinium</i> <i>goizowense</i>	1.40: 115: <i>Palaeocystodinium</i> <i>goizowense</i>
1.40: 248: <i>Membranosphaera</i> <i>tabulata</i>	1.20: 250: <i>Palaeostomacystus</i> <i>fragilis</i>	1.40: 250: <i>Palaeostomacystus</i> <i>fragilis</i>
1.20: 30: <i>Chlamydomonada</i> cf. <i>C. jma</i>	1.10: 249: <i>Nematosphaerosis</i> <i>irabeculata</i>	1.00: 33: <i>Kaliosphaeridium</i> <i>brevibarbatum</i>
1.20: 248: <i>Deilandra</i> <i>spinigera</i>	1.00: 33: <i>Kaliosphaeridium</i> <i>brevibarbatum</i>	0.90: 31: <i>Hyalinosphaeridium</i> <i>tubiferum</i>
1.20: 249: <i>Nematosphaerosis</i> <i>irabeculata</i>	0.90: 31: <i>Hyalinosphaeridium</i> <i>tubiferum</i>	0.80: 253: <i>Spiriferites</i> cf. <i>S. septatus</i>
1.10: 250: <i>Palaeostomacystus</i> <i>fragilis</i>	0.90: 253: <i>Spiriferites</i> cf. <i>S. septatus</i>	0.80: 252: <i>Spiriferites</i> <i>septatus</i>
0.90: 190: <i>Diovesopsis</i> <i>capitata</i>	0.90: 252: <i>Spiriferites</i> <i>septatus</i>	0.50: 249: <i>Nematosphaerosis</i> <i>irabeculata</i>
0.90: 115: <i>Palaeocystodinium</i> <i>goizowense</i>	0.40: 110: <i>Operculodinium</i> <i>centrocarpum</i>	0.40: 251: <i>Corosphaeridium</i> <i>librospinosum</i>
0.80: 47: <i>Eocladopyxis</i> <i>penicillatum</i>	0.40: 151: <i>Spiriferites</i> <i>ramosus</i> subsp. <i>ramosus</i>	0.40: 190: <i>Diovesopsis</i> <i>capitata</i>
0.70: 28: <i>Corosphaeridium</i> <i>inodes</i>	0.30: 28: <i>Corosphaeridium</i> <i>inodes</i>	0.30: 47: <i>Eocladopyxis</i> <i>penicillatum</i>
0.60: 38: <i>Linguodinium</i> <i>macnaerophorum</i>	0.20: 241: <i>Corosphaeridium</i> <i>species</i> A	0.30: 181: <i>Micrystidium</i> spp.
0.40: 243: <i>Deilandra</i> <i>obscura</i>	0.20: 243: <i>Deilandra</i> <i>obscura</i>	0.30: 101: <i>Microdinium</i> <i>ornatum</i>
0.40: 245: <i>Deilandra</i> <i>ruginosa</i>	0.20: 190: <i>Diovesopsis</i> <i>capitata</i>	0.20: 29: <i>Corosphaeridium</i> <i>inodes</i>
0.40: 110: <i>Operculodinium</i> <i>centrocarpum</i>	0.20: 47: <i>Eocladopyxis</i> <i>penicillatum</i>	0.20: 182: <i>Cymatosphaera</i> spp.
0.30: 181: <i>Micrystidium</i> spp.	0.20: 38: <i>Linguodinium</i> <i>macnaerophorum</i>	0.20: 242: <i>Deilandra</i> cf. <i>D. magnifica</i>
0.20: 242: <i>Deilandra</i> cf. <i>D. magnifica</i>	0.20: 101: <i>Microdinium</i> <i>ornatum</i>	0.20: 245: <i>Deilandra</i> <i>ruginosa</i>
0.20: 194: <i>Hyalinosphaerosis</i> <i>borussica</i>	0.20: 186: <i>Pterospermopsis</i> spp.	0.20: 39: <i>Impatiensphaeridium</i> <i>rugosum</i>
0.20: 186: <i>Pterospermopsis</i> spp.	0.10: 240: <i>Corosphaeridium</i> <i>marylandense</i>	0.20: 10: <i>Operculodinium</i> <i>centrocarpum</i>
0.20: 136: <i>Spiriferites</i> <i>crassipellis</i> A	0.10: 242: <i>Deilandra</i> cf. <i>D. magnifica</i>	0.20: 186: <i>Pterospermopsis</i> spp.
0.20: 151: <i>Spiriferites</i> <i>ramosus</i> subsp. <i>ramosus</i>	0.10: 37: <i>Deilandra</i> <i>onosphortica</i>	0.20: 169: <i>Weitzelia</i> cf. <i>w. rhyssensis</i>
0.10: 240: <i>Corosphaeridium</i> <i>marylandense</i>	0.10: 245: <i>Deilandra</i> <i>ruginosa</i>	0.10: 240: <i>Corosphaeridium</i> <i>marylandense</i>
0.10: 37: <i>Deilandra</i> <i>onosphortica</i>	0.10: 39: <i>Impatiensphaeridium</i> <i>rugosum</i>	0.10: 78: <i>Cycloosella</i> <i>vieta</i>
0.10: 244: <i>Deilandra</i> <i>omboneora</i>	0.10: 181: <i>Micrystidium</i> spp.	0.10: 243: <i>Deilandra</i> <i>obscura</i>
0.10: 18: <i>Hyalinoclopidium</i> <i>unisanum</i>	0.10: 251: <i>Spiriferites</i> <i>cingulatus</i> subsp. <i>reticulatus</i>	0.10: 37: <i>Deilandra</i> <i>onosphortica</i>
0.10: 39: <i>Impatiensphaeridium</i> <i>rugosum</i>	0.10: 136: <i>Spiriferites</i> <i>crassipellis</i> A	0.10: 76: <i>Hyalinoclopidium</i> <i>unisanum</i>
0.10: 101: <i>Microdinium</i> <i>ornatum</i>	0.10: 148: <i>Spiriferites</i> <i>ramosus</i> subsp. <i>granosus</i>	0.10: 38: <i>Linguodinium</i> <i>macnaerophorum</i>
0.10: 251: <i>Spiriferites</i> <i>cingulatus</i> subsp. <i>reticulatus</i>	0.10: 169: <i>Weitzelia</i> cf. <i>w. rhyssensis</i>	0.10: 251: <i>Spiriferites</i> <i>cingulatus</i> subsp. <i>reticulatus</i>
0.10: 148: <i>Spiriferites</i> <i>ramosus</i> subsp. <i>granosus</i>		0.10: 136: <i>Spiriferites</i> <i>crassipellis</i> A
0.10: 136: <i>Spiriferites</i> <i>crassipellis</i> A		0.10: 48: <i>Spiriferites</i> <i>ramosus</i> subsp. <i>granosus</i>
		0.10: 151: <i>Spiriferites</i> <i>ramosus</i> subsp. <i>ramosus</i>

188	189	190
48.20: 255 Weitzelia (W.) nomomorona subsp. quinq.	33.40: 255 Weitzelia (W.) nomomorona subsp. quinq.	30.80: 255 Weitzelia (W.) nomomorona subsp. quinq.
2.50: 248 Membranospaera tabuiata	26.00: 248 Membranospaera tabuiata	27.50: 248 Membranospaera tabuiata
3.60: 196 Lantemosphaeridium lanosum	7.00: 20 Chlamydomonella cf. C. urna	6.70: 20 Chlamydomonella cf. C. urna
7.40: 20 Chlamydomonella cf. C. urna	6.10: 12 Aeoligera sp.	5.90: 12 Aeoligera sp.
5.20: 12 Aeoligera sp.	3.40: 196 Lantemosphaeridium lanosum	5.60: 196 Lantemosphaeridium lanosum
3.40: 246 Deltandna spinigera	1.50: 246 Deltandna spinigera	1.30: 246 Deltandna spinigera
1.10: 247 Lactodinium virginianum	1.20: 115 Palaecystodinium goizowense	1.50: 81 Hystichosphaeridium tubiferum
1.10: 115 Palaecystodinium goizowense	1.20: 250 Palaecostomacystus fragilis	1.20: 77 Hystichokolpoma tumescens
1.10: 250 Palaecostomacystus fragilis	1.10: 77 Hystichokolpoma tumescens	1.10: 115 Palaecystodinium goizowense
1.00: 249 Nematosphaerosus trabeculata	3.00: 28 Cordosphaeridium inodes	1.10: 250 Palaecostomacystus fragilis
0.90: 25 Cordosphaeridium fibrosinosum	3.00: 81 Hystichosphaeridium tubiferum	0.80: 29 Cordosphaeridium inodes
0.90: 253 Spiniferes cf. S. septatus	0.90: 253 Spiniferes cf. S. septatus	0.80: 249 Nematosphaerosus trabeculata
0.90: 252 Spiniferes septatus	0.90: 252 Spiniferes septatus	0.70: 253 Spiniferes cf. S. septatus
0.80: 81 Hystichosphaeridium tubiferum	0.80: 25 Cordosphaeridium fibrosinosum	0.70: 252 Spiniferes septatus
0.70: 240 Cordosphaeridium marylandense	0.80: 190 Dihyvesopsis capitata	0.60: 25 Cordosphaeridium fibrosinosum
0.70: 190 Dihyvesopsis capitata	0.40: 240 Cordosphaeridium marylandense	0.40: 240 Cordosphaeridium marylandense
0.60: 28 Cordosphaeridium inodes	0.40: 93 Kaiiosphaeridium breviarbatum	0.30: 243 Deltandna obscura
0.60: 93 Kaiiosphaeridium breviarbatum	0.30: 195 Lantemosphaeridium bioiare	0.30: 245 Deltandna ruginosa
0.60: 110 Operculodinium centrocarpum	0.30: 247 Lactodinium virginianum	0.30: 93 Kaiiosphaeridium breviarbatum
0.40: 136 Spiniferes crassipellis A	0.30: 249 Nematosphaerosus trabeculata	0.30: 10 Operculodinium centrocarpum
0.30: 243 Deltandna obscura	0.30: 148 Spiniferes ramosus subsp. granosus	0.20: 178 Cyclopsella veta
0.30: 77 Hystichokolpoma tumescens	0.20: 243 Deltandna obscura	0.20: 195 Lantemosphaeridium bioiare
0.30: 146 Spiniferes pseudofurcatus	0.20: 78 Hystichokolpoma unisonum	0.20: 38 Linguodinium machaerophorum
0.20: 241 Cordosphaeridium species A	0.20: 98 Linguodinium machaerophorum	0.20: 186 Pterodermopsis spp.
0.20: 244 Deltandna mombocera	0.20: 181 Microstadium spp.	0.20: 136 Spiniferes crassipellis A
0.20: 47 Sociadopyxis penicularum	0.20: 101 Microstadium ornatum	0.20: 146 Spiniferes pseudofurcatus
0.20: 194 Hystichosphaerosus borussica	0.20: 110 Operculodinium centrocarpum	0.20: 148 Spiniferes ramosus subsp. granosus
0.20: 195 Lantemosphaeridium bioiare	0.20: 136 Spiniferes crassipellis A	0.10: 244 Deltandna mombocera
0.20: 251 Spiniferes angulatus subsp. reticulatus	0.20: 146 Spiniferes pseudofurcatus	0.10: 190 Dihyvesopsis capitata
0.20: 69 Weitzelia cf. w. rhysschenis	0.20: 151 Spiniferes ramosus subsp. ramosus	0.10: 78 Hystichokolpoma unisonum
0.10: 242 Deltandna cf. D. magnifica	0.10: 241 Cordosphaeridium species A	0.10: 194 Hystichosphaerosus borussica
0.10: 245 Deltandna ruginosa	0.10: 179 Cyclopsella veta	0.10: 39 Impletosphaeridium rugosum
0.10: 78 Hystichokolpoma unisonum	0.10: 245 Deltandna ruginosa	0.10: 247 Lactodinium virginianum
0.10: 39 Impletosphaeridium rugosum	0.10: 76 Hystichokolpoma nquace	0.10: 181 Microstadium spp.
0.10: 38 Linguodinium machaerophorum	0.10: 194 Hystichosphaerosus borussica	0.10: 151 Spiniferes ramosus subsp. ramosus
0.10: 146 Spiniferes ramosus subsp. granosus	0.10: 39 Impletosphaeridium rugosum	
0.10: 151 Spiniferes ramosus subsp. ramosus	0.10: 186 Pterodermopsis spp.	



94	95	96
34.00  355 Weizaelella (W.) homomorpha subsp. quinque	40.10  255 Weizaelella (W.) homomorpha subsp. quinque	30.40  255 Weizaelella (W.) homomorpha subsp. quinque
32.50  248 Membranosphaera tabulata	29.20  248 Membranosphaera tabulata	30.00  248 Membranosphaera tabulata
6.30  20 Chlamydomonella cf. C. urna	6.00  20 Chlamydomonella cf. C. urna	15.50  20 Chlamydomonella cf. C. urna
3.20  245 Dellandina spingera	2.20  245 Dellandina spingera	7.20  196 Lanternosphaeridium lanosum
2.20  196 Lanternosphaeridium lanosum	2.10  115 Palaecocystodinium goizowense	2.20  93 Kaiosphaeridium brevisarbatum
1.40  28 Cordosphaeridium nodos	2.10  250 Palaecostomacystus fragilis	1.50  115 Palaecocystodinium goizowense
1.30  33 Kaiosphaeridium brevisarbatum	1.90  196 Lanternosphaeridium lanosum	1.50  250 Palaecostomacystus fragilis
1.20  12 Aeroigera sp.	1.20  243 Dellandina obscura	1.20  12 Aeroigera sp.
1.10  243 Dellandina obscura	1.10  12 Aeroigera sp.	1.20  246 Dellandina spingera
1.50  25 Cordosphaeridium fibrosinosum	0.60  01 Microdinium ornatum	0.70  240 Cordosphaeridium mariandense
1.50  77 Hystichosphaeridium tumescens	0.40  81 Microdinium sp.	0.70  311 Hystichosphaeridium tubiferum
1.50  250 Palaecostomacystus fragilis	0.30  86 Perospermopsis sp.	0.50  251 Cordosphaeridium fibrosinosum
0.50  81 Microdinium sp.	0.20  238 Aptodinium baculatum	0.50  243 Dellandina obscura
0.50  115 Palaecocystodinium goizowense	0.20  251 Cordosphaeridium fibrosinosum	0.50  77 Hystichosphaeridium tumescens
0.40  240 Cordosphaeridium mariandense	0.20  281 Cordosphaeridium nodos	0.50  381 Lingulodinium machaerophorum
0.40  249 Nematosphaeropsis trabeoculata	0.20  244 Dellandina rhomboides	0.50  249 Nematosphaeropsis trabeoculata
0.40  253 Spiriferes cf. S. septatus	0.20  245 Dellandina ruginosa	0.50  136 Spiriferes crassipellis A
0.40  252 Spiriferes septatus	0.20  391 Impietosphaeridium rugosum	0.50  151 Spiriferes ramosus subsp. ramosus
0.30  17 Dellandina onosphontica	0.20  33 Kaiosphaeridium brevisarbatum	0.40  281 Cordosphaeridium nodos
0.30  245 Dellandina ruginosa	0.20  247 Laptodinium virginianum	0.40  195 Lanternosphaeridium bipolare
0.30  391 Impietosphaeridium rugosum	0.20  249 Nematosphaeropsis trabeoculata	0.30  245 Dellandina ruginosa
0.30  195 Lanternosphaeridium bipolare	0.20  136 Spiriferes crassipellis A	0.30  190 Diphyesopsis capitata
0.30  247 Laptodinium virginianum	0.20  511 Spiriferes ramosus subsp. ramosus	0.30  391 Impietosphaeridium rugosum
0.30  136 Spiriferes crassipellis A	0.20  169 Weizaelella cf. w. ruschensis	0.30  247 Laptodinium virginianum
0.20  190 Diphyesopsis capitata	0.10  240 Cordosphaeridium mariandense	0.30  01 Microdinium ornatum
0.20  311 Hystichosphaeridium tubiferum	0.10  190 Diphyesopsis capitata	0.30  253 Spiriferes cf. S. septatus
0.20  381 Lingulodinium machaerophorum	0.10  77 Hystichosphaeridium tumescens	0.30  148 Spiriferes ramosus subsp. granosus
0.20  01 Microdinium ornatum	0.10  781 Hystichosphaeridium unispinum	0.30  252 Spiriferes septatus
0.20  01 Coercuodinium centrocarpum	0.10  311 Hystichosphaeridium tubiferum	0.20  238 Aptodinium baculatum
0.20  196 Perospermopsis sp.	0.10  195 Lanternosphaeridium bipolare	0.20  82 Cymatosphaera sp.
0.20  511 Spiriferes ramosus subsp. ramosus	0.10  381 Lingulodinium machaerophorum	0.20  781 Hystichosphaeridium unispinum
0.10  238 Aptodinium baculatum	0.10  01 Coercuodinium centrocarpum	0.20  194 Hystichosphaeropsis borussica
0.10  75 Hystichosphaeridium guade	0.10  253 Spiriferes cf. S. septatus	0.20  110 Coercuodinium centrocarpum
0.10  781 Hystichosphaeridium unispinum	0.10  146 Spiriferes pseudoturcatus	0.20  146 Spiriferes pseudoturcatus
0.10  146 Spiriferes pseudoturcatus	0.10  148 Spiriferes ramosus subsp. granosus	0.10  781 Cyclosetella vieta
0.10  148 Spiriferes ramosus subsp. granosus	0.10  252 Spiriferes septatus	0.10  86 Perospermopsis sp.
0.10  169 Weizaelella cf. w. ruschensis		0.10  169 Weizaelella cf. w. ruschensis

	197	
41	40	255: Wetzelia (W.) homomorpha subsp. quinque
31	50	248: Membranosphaera tabulata
14	20	20: Chlamydomonella cf. C. urna
2	40	196: Lanternosphaeridium lanosum
1	40	2: Aroloigera sp.
0	80	93: Kalliosphaeridium brevispatulum
0	50	25: Corcosphaeridium fibropinosum
0	50	240: Corcosphaeridium marylandense
0	50	245: Delandona rugulosa
0	40	243: Delandona obscura
0	40	246: Delandona spingera
0	40	98: Linguodinium macraeophorum
0	40	181: Micrystidium sp.
0	40	250: Palaeostomacystus fragilis
0	40	151: Spiniferites ramosus subsp. ramosus
0	30	28: Corcosphaeridium inopes
0	30	77: Hystichosphaeridium tumescens
0	30	89: Impatiensphaeridium rugosum
0	20	238: Aptedinium baculatum
0	20	190: Dipnemosopsis capitata
0	20	78: Hystichosphaeridium unispinum
0	20	81: Hystichosphaeridium tubiferum
0	20	195: Lanternosphaeridium bipolare
0	20	247: Leptodinium virginianum
0	20	101: Microdinium ornatum
0	20	249: Nematospaeropsis trabeulata
0	20	110: Operculodinium centrocarpum
0	20	115: Palaeocystodinium gozowense
0	20	186: Pterospemopsis sp.
0	20	253: Spiniferites cf. S. setatus
0	20	251: Spiniferites cingulatus subsp. reticulatus
0	20	136: Spiniferites crassipellis A
0	20	252: Spiniferites setatus
0	20	169: Wetzelia cf. W. rhysschensis
0	10	178: Cyclopsella vieta
0	10	194: Hystichosphaeropsis borussica
0	10	146: Spiniferites pseudofurcatus
0	10	148: Spiniferites ramosus subsp. granosus

Correlation Matrix for PT3

Correlation Matrix for Variables: X1 ... X13

	183	184	185	186	187	188	189	190
183	1							
184	.92104	1						
185	.80824	.95885	1					
186	.76673	.94216	.98162	1				
187	.58132	.72576	.76791	.85738	1			
188	.71181	.88932	.92917	.9734	.91282	1		
189	.55201	.69553	.73805	.8328	.99757	.89957	1	
190	.52405	.65604	.69874	.79761	.99103	.87937	.99642	1
191	.62173	.77427	.81607	.89201	.97384	.95895	.97351	.96739
194	.49197	.6332	.67804	.78548	.98541	.8586	.98724	.98814
195	.54982	.71283	.75957	.85607	.99233	.91239	.98856	.98128
196	.49126	.62143	.66643	.77213	.97907	.86298	.98365	.9911
197	.55276	.71423	.759	.8546	.98904	.91245	.9856	.97982

Correlation Matrix for Variables: X1 ... X13

	191	194	195	196	197
191	1				
194	.94293	1			
195	.96549	.98977	1		
196	.95451	.99224	.98083	1	
197	.96219	.99004	.99654	.98187	1

## ANOVA Table for PT3

One Factor ANOVA-Repeated Measures for X <sub>1</sub> ... X <sub>13</sub>					
Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	27994.76687	110.21562	69.39686	.0001
Within subjects	3060	4859.87077	1.58819		
treatments	12	.00524	.00044	.00027	1
residual	3048	4859.86553	1.59444		
Total	3314	32854.63763			

Reliability Estimates for- All treatments: .98559    Single Treatment: .34029

One Factor ANOVA-Repeated Measures for X <sub>1</sub> ... X <sub>13</sub>				
Group:	Count:	Mean:	Std. Dev.:	Std. Error:
183	255	.39412	2.72401	.17058
184	255	.39176	3.3715	.21113
185	255	.39647	3.56685	.22336
186	255	.39373	3.73425	.23385
187	255	.39373	2.93882	.18404

One Factor ANOVA-Repeated Measures for X <sub>1</sub> ... X <sub>13</sub>				
Group:	Count:	Mean:	Std. Dev.:	Std. Error:
188	255	.39529	3.23527	.2026
189	255	.3949	2.87663	.18014
190	255	.39451	2.82223	.17674
191	255	.39529	2.93586	.18385
194	255	.39451	3.12191	.1955

PT4 "Wetzeliella (W.) homomorpha/Deflandria dilwynensis/Muratodinium fimbriatum" Association

199	200	201
42.10: 255:Wetzeliella (W.) homomorpha subsp. quinct	40.70: 255:Wetzeliella (W.) homomorpha subsp. quinct	36.50: 255:Wetzeliella (W.) homomorpha subsp. quinct
30.00: 213:Deflandria dilwynensis	31.20: 213:Deflandria dilwynensis	32.50: 213:Deflandria dilwynensis
25.20: 103:Muratodinium fimbriatum	25.30: 103:Muratodinium fimbriatum	29.50: 103:Muratodinium fimbriatum
1.00: 186:Pterospermopsis spp.	0.20: 186:Pterospermopsis spp.	0.50: 186:Pterospermopsis spp.
0.50: 181:Michystidium spp.	0.40: 181:Michystidium spp.	0.40: 181:Michystidium spp.
0.20: 20:Chlamydomonada cf. Clona	0.40: 151:Spiniferites ramosus subsp. ramosus	0.20: 243:Deflandria obscura
0.20: 249:Nematosphaeropsis trabeculata	0.20: 249:Nematosphaeropsis trabeculata	0.20: 249:Nematosphaeropsis trabeculata
0.10: 182:Cymatosphaera spp.	0.10: 182:Cymatosphaera spp.	0.10: 182:Cymatosphaera spp.
0.10: 243:Deflandria obscura		0.10: 151:Spiniferites ramosus subsp. ramosus
0.10: 246:Deflandria spinigera		
0.10: 195:Lantimosphaeridium bipolare		
0.10: 151:Spiniferites ramosus subsp. ramosus		



Correlation Matrix for Variables: X<sub>1</sub> ... X<sub>3</sub>

	199	200	201
199	1		
200	.99938	1	
201	.99133	.99489	1

One Factor ANOVA-Repeated Measures for X<sub>1</sub> ... X<sub>3</sub>

Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	9716.17783	38.25267	624.75003	.0001
Within subjects	510	31.22667	.06123		
treatments	2	.00003	.00001	.00021	.9998
residual	508	31.22664	.06147		
Total	764	9747.4045			

Reliability Estimates for- All treatments: .9984 Single Treatment: .99521

PT5 "Glaphyrocysta exuberans complex/Fibradinium annetorpense"

Association

3031	winter aqua percentages	3041	winter aqua percentages
28.97	331 Glaphyrocysta exuberans spix	26.00	531 Glaphyrocysta exuberans spix
14.00	491 Fibradinium annetorpense	20.00	491 Fibradinium annetorpense
7.32	461 Electrocysta obscurotabulata	3.00	751 Xenkoon australe
9.33	751 Xenkoon australe	7.50	461 Electrocysta obscurotabulata
9.30	261 Corrososphaeridium giganteum	4.33	116 Palaeosphaeridium synporum
5.87	116 Palaeosphaeridium synporum	4.67	161 Dallanona garrmana
3.33	311 Hystriochosphaeridium lupiferum	3.33	1801 Micrystidium cf. m. ranae
3.20	1801 Micrystidium cf. m. ranae	3.00	281 Corrososphaeridium nodosum
2.67	281 Corrososphaeridium nodosum	2.33	261 Corrososphaeridium giganteum
1.67	161 Dallanona garrmana	2.33	551 Floeremina tergy
3.33	37 Corrososphaeridium gracile	2.33	311 Hystriochosphaeridium lupiferum
1.67	34 Danaea sp.	2.00	117 Paraeacanthella indentata
1.33	171 Calogadinium amiculum	1.67	171 Calogadinium amiculum
1.33	1831 Palaeoagras sp. A	1.33	271 Corrososphaeridium gracile
1.00	201 Chlamydothorea cf. C. urna	2.67	341 Danaea sp.
1.00	1101 Coercuodinium centrocarpum	2.67	451 Electrocysta densopaculata
1.00	511 Soriniferites ramosus suoso ramosus	2.67	101 Coercuodinium centrocarpum
1.00	1341 Palaeoagras sp. B	2.67	511 Soriniferites ramosus suoso ramosus
2.67	311 Cladoxanthidium saeculum	2.67	591 Thalassiothora delicata
2.67	331 Conniximura limbrata	2.67	931 Palaeoagras sp. A
2.67	451 Electrocysta densopaculata	2.33	51 Andalusella rombolocera
2.67	541 Fibrocysta sp.	2.33	201 Chlamydothorea cf. C. urna
2.67	351 Floeremina terox	2.33	211 Cladoxanthidium saeculum
2.67	581 Harnissphaera septata	2.33	231 Conniximura limbrata
2.67	151 Palaeocystodinium gozowense	2.33	251 Corrososphaeridium fibrosinosum
2.67	117 Paraeacanthella indentata	2.33	331 Corrososphaeridium caliosum
2.67	591 Thalassiothora delicata	2.33	411 Dionyes colligerum
2.33	251 Corrososphaeridium fibrosinosum	2.33	431 Dionyes colligerum sensu Cookson 1965
2.33	331 Corrososphaeridium caliosum	2.33	481 Exochosphaeridium striatum
2.33	411 Dionyes colligerum	2.33	541 Fibrocysta sp.
2.33	431 Dionyes colligerum sensu Cookson 1965	2.33	561 Forma A
2.33	481 Exochosphaeridium striatum	2.33	511 Fromea fragilis
2.33	561 Forma A	2.33	581 Harnissphaera septata
2.33	511 Fromea fragilis	2.33	731 Horologipella punctata
2.33	781 Hystriochosphaeridium unispinum	2.33	891 Impatiensphaeridium rugosum
2.33	321 Hystriochosphaeridium lupiferum brevispin	2.33	101 Microdinium ornatum
2.33	391 Impatiensphaeridium rugosum	2.33	051 Nematosphaeropsis pertusa
2.33	311 Inversidinium eximium	2.33	081 Oligosphaeridium complex
2.33	1011 Microdinium ornatum	2.33	151 Palaeocystodinium gozowense
2.33	051 Nematosphaeropsis pertusa	2.33	201 Phaeodinium magnificum
2.33	081 Oligosphaeridium complex	2.33	1321 Senegalinum? diluviana
2.33	101 Phaeodinium magnificum	2.33	161 Soriniferites crassipellis A
2.33	1251 Renedinium membraniferum	2.33	461 Soriniferites pseudofurcatus
2.33	1321 Senegalinum? diluviana	2.33	1471 Soriniferites ramosus suoso granomemoran
2.33	1361 Soriniferites crassipellis A	2.33	1491 Soriniferites ramosus suoso membranaceo
2.33	1461 Soriniferites pseudofurcatus	2.33	501 Soriniferites ramosus suoso multibrevis
2.33	1471 Soriniferites ramosus suoso granomemoran	2.33	571 Tanvossphaeridium varacatum
2.33	1481 Soriniferites ramosus suoso granosus	2.33	1791 Micrystidium cf. m. fragile
2.33	1501 Soriniferites ramosus suoso multibrevis	2.33	1811 Micrystidium sp.
2.33	1571 Tanvossphaeridium varacatum	2.33	821 Cymatosphaera sp.
2.33	1801 Thalassiothora pelagica	2.33	861 Pterosphaeropsis sp.
2.33	1821 Trigonopyxidia gineia		
2.33	1791 Micrystidium cf. m. fragile		
2.33	1821 Cymatosphaera sp.		
2.33	1861 Pterosphaeropsis sp.		

3091	wimer aquia percentages	3101	wimer aquia percentages	3111	wimer aquia percentages
34.33	53: Glaphrocysta exuberans cplx	33.00	49: Foradinium annetorpense	50.00	53: Glaphrocysta exuberans cplx
31.33	49: Foradinium annetorpense	21.33	53: Glaphrocysta exuberans cplx	23.00	49: Foradinium annetorpense
9.67	75: Xenkoon australe	4.67	76: Xenkoon australe	4.67	46: Electrocysta obscurotubulata
4.00	46: Electrocysta obscurotubulata	3.67	46: Electrocysta obscurotubulata	3.00	55: Glaphrocysta sp A
3.33	35: Glaphrocysta sp A	3.00	26: Cordosphaeridium giganteum	2.33	36: Deltanona darmonia
1.00	7: Caigodinium amicum	2.57	35: Glaphrocysta sp A	1.33	37: Cordosphaeridium gracilis
2.57	25: Cordosphaeridium giganteum	2.33	78: Cyclosetella vieta	1.00	61: Microvistridium sop
2.00	80: Microvistridium cf. m. vanabone	2.00	27: Cordosphaeridium gracile	0.57	28: Cordosphaeridium nodos
1.33	29: Cordosphaeridium nodos	2.00	66: Turbosphaera hlosa	0.67	45: Electrocysta densocuculata
1.33	15: Palaeocystodinium pyrohorum	1.57	36: Deltanona darmonia	0.67	37: Lentinia spongiera
0.57	5: Batacasphaera sp	1.57	17: Parazacanelia indentata	0.57	66: Turbosphaera hlosa
0.57	34: Danae sp	1.33	80: Microvistridium cf. m. vanabone	0.57	76: Xenkoon australe
0.67	36: Deltanona darmonia	1.00	97: Lentinia spongiera	0.33	13: Ascotomocystus hydra
0.67	45: Electrocysta densocuculata	1.00	15: Palaeocystodinium gozowense	0.33	15: Batacasphaera sp
0.57	31: Fromea fragilis	1.00	51: Sonfientes ramosus subsp. ramosus	0.33	7: Caigodinium amicum
0.57	58: Halmisphaera septata	0.57	28: Cordosphaeridium nodos	0.33	20: Chlamydocornella cf. C. uma
0.57	17: Parazacanelia indentata	0.57	58: Halmisphaera septata	0.33	25: Cordosphaeridium fibrososolum
0.57	84: Palaeopages sp. B	0.57	05: Nematospaeropsis pertusa	0.33	26: Cordosphaeridium giganteum
0.33	31: Ascotomocystus hydra	0.57	120: Pheodinium magnificum	0.33	47: Diphyes colligerum
0.33	20: Chlamydocornella cf. C. uma	0.57	48: Sonfientes ramosus subsp. granosus	0.33	42: Diphyes colligerum
0.33	21: Cladocystidium saetorum	0.33	7: Caigodinium amicum	0.33	56: Forma A
0.33	33: Conneximura remorata	0.33	20: Chlamydocornella cf. C. uma	0.33	51: Fromea fragilis
0.33	27: Cordosphaeridium gracile	0.33	21: Cladocystidium saetorum	0.33	58: Halmisphaera septata
0.33	37: Cordosphaeridium calosum	0.33	33: Cordosphaeridium calosum	0.33	73: Horologinella apiculata
0.33	46: Exochosphaeridium bifidum	0.33	42: Diphyes colligerum	0.33	31: Hystrichosphaeridium tubiferum
0.33	55: Halmisphaera ferox	0.33	43: Diphyes colligerum (sensu Cookson 1985)	0.33	89: Impletosphaeridium rugosum
0.33	73: Horologinella apiculata	0.33	45: Electrocysta densocuculata	0.33	31: Inversidinium eximurum
0.33	81: Hystrichosphaeridium tubiferum	0.33	56: Forma A	0.33	01: Microdinium ornatum
0.33	01: Microdinium ornatum	0.33	51: Fromea fragilis	0.33	05: Nematospaeropsis pertusa
0.33	05: Nematospaeropsis pertusa	0.33	52: Fromea? laevigata	0.33	15: Palaeocystodinium gozowense
0.33	08: Oligosphaeridium complex	0.33	75: Hystrichoboloma mamatum	0.33	17: Parazacanelia indentata
0.33	10: Oerccuodinium centrocarpum	0.33	30: Hystrichosphaeridium sop	0.33	120: Pheodinium magnificum
0.33	20: Pheodinium magnificum	0.33	81: Hystrichosphaeridium tubiferum	0.33	25: Rendingium membranarium
0.33	27: Rottneistra borussica	0.33	07: Nematospaeropsis rabeculata	0.33	29: Samandia reticulifera
0.33	29: Samandia reticulifera	0.33	10: Oerccuodinium centrocarpum	0.33	33: Sponidium essei
0.33	32: Senegalinum? dityrrense	0.33	127: Rottneistra borussica	0.33	51: Sonfientes ramosus subsp. ramosus
0.33	33: Sponidium essei	0.33	29: Samandia reticulifera	0.33	59: Thalassiohara delicata
0.33	39: Sonfientes cf. s. terotus	0.33	30: Senegalinum? obscurum	0.33	62: Trigonopyxidia gineia
0.33	51: Sonfientes ramosus subsp. ramosus	0.33	30: Senegalinum? obscurum	0.33	79: Cyclosetella vieta
0.33	56: Systematohara placacantha	0.33	47: Sonfientes ramosus subsp. granomembrans	0.33	79: Microvistridium cf. m. fragile
0.33	59: Thalassiohara delicata	0.33	49: Sonfientes ramosus subsp. membranaceus	0.33	84: Palaeopages sp. B
0.33	79: Microvistridium cf. m. fragile	0.33	52: Sonfientes ramosifera	0.33	86: Pterocermopsis sop
0.33	82: Cymatospaera sop	0.33	59: Thalassiohara delicata		
0.33	86: Pterocermopsis sop	0.33	62: Trigonopyxidia gineia		

112'	wtmer aqua percentages
58.67'	331 Glaphrocysta exuberans cplx
7.33'	*661 Turosohaera hiosa
4.57'	361 Delandria carmens
3.33'	*751 Xenophon australis
3.00'	551 Glaphrocysta sp. A
3.33'	271 Cordosphaeridium gracilis
*.57'	*781 Cyclospora vieta
*.33'	461 Electrocyta obscuratoluita
*.00'	291 Cordosphaeridium nodos
3.57'	*.71 Calogonium amiculum
2.57'	451 Electrocyta densobaculata
3.57'	311 Fromea fragilis
2.57'	371 Lentia spongosa
2.57'	*461 Sommerites pseudoturcatus
3.57'	*61 Micromystridium spp.
3.57'	*841 Paalimages sp. B
3.33'	41 Alcocysta cf. a. margarita
3.33'	131 Ascotomocystus hydria
3.33'	*51 Baracasonaera sp.
3.33'	201 Chlamydomonada cf. C. urna
3.33'	261 Cordosphaeridium giganteum
3.33'	271 Delandria phosphorica
3.33'	491 Faradrium annetense
3.33'	581 Halmasphaera sectata
3.33'	301 Hysterochaetium spp.
3.33'	311 Hysterochaetium tubiferum
3.33'	*941 Nematosphaerosis cf. n. obtusa
3.33'	*851 Nematosphaerosis pertusa
3.33'	*871 Nematosphaerosis vapidulata
3.33'	*101 Oertruodinium centrocarpum
3.33'	*151 Pataocystodinium gozowense
3.33'	*171 Paracanieta indanata
3.33'	*251 Renidium membraniferum
3.33'	*321 Seneganium? demense
3.33'	*511 Sommerites ramosus subsp. ramosus
3.33'	*591 Thalesionora delicata
3.33'	*801 Micromystridium cf. m. variable
3.33'	*821 Cymatosphaera spp.

Correlation Matrix for PT 5

Correlation Matrix for Variables:		X <sub>1</sub> ... X <sub>5</sub>				
	304	309	310	311	312	
304	1					
309	.93835	1				
310	.90098	.85814	1			
311	.89215	.96045	.75493	1		
312	.74139	.87105	.52544	.92808	1	

ANOVA Table for PT 5

One Factor ANOVA-Repeated Measures for		X <sub>1</sub> ... X <sub>5</sub>			
Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	14311.97736	56.34637	18.42889	.0001
Within subjects	1020	3118.65252	3.0575		
treatments	4	5.43712	1.35928	.4436	.7771
residual	1016	3113.2154	3.06419		
Total	1274	17430.62988			

Reliability Estimates for- All treatments: .94574      Single Treatment: .77707

PT6 "Glaphyrocysta exuberans complex/ Cordosphaeridium giganteum/  
 Fibradinium annetorpense" Association

1051	water aqua percentages
22.67	49 Fibradinium annetorpense
16.33	26 Cordosphaeridium giganteum
7.33	46 Electrocysta ooscuroradiata
6.67	36 Dallanona darmorata
4.67	63 Glaphyrocysta exuberans complex
4.00	34 Danea sp.
3.33	80 Microthidium cf. m. variabile
2.67	01 Microthidium ornatum
2.00	27 Cordosphaeridium gracile
2.00	45 Electrocysta gensoacuiata
1.67	20 Chaetomyxonorella cf. C. urna
1.67	28 Cordosphaeridium inodes
1.67	55 Florentina ferax
1.67	105 Nematospiraerosa pertusa
1.67	117 Parasaccharia inerrata
1.67	129 Santandia ruscifera
1.33	56 Forma A
1.33	81 Hystichosphaeridium tubiferum
1.33	176 Xenicon australe
1.00	5 Baltica-sphaera sp.
1.00	151 Soinferes ramosus suoso ramosus
0.67	5 Andalusella rhomboida
0.67	7 Calogonidium amiculum
0.67	68 Halimosaera senata
0.67	110 Opertuodinium centrocarpum
0.67	116 Palaeoendinium cyroponorum
0.67	136 Soinferes crassipellis A
0.67	82 Trigonopyxidia gneiss
0.33	13 Ascotomocystus myrina
0.33	21 Craoopyxidium sagittum
0.33	23 Cohensitura limonata
0.33	25 Cordosphaeridium hirsutiniosum
0.33	42 Dermis colligerum
0.33	48 Exochosphaeridium bifidum
0.33	61 Fromea fragilis
0.33	82 Hystichosphaeridium tubiferum brevipes
0.33	85 Limogonidium sp. A
0.33	89 Limnetosphaeridium rugosum
0.33	91 Inversorium eximium
0.33	97 Lentia spongosa
0.33	115 Palaeocystodinium goizowense
0.33	120 Phaeodinium magnificum
0.33	125 Rendrium membraniferum
0.33	127 Rodnesia borussica
0.33	133 Soinferium essei
0.33	141 Soinferes cornutus A
0.33	146 Soinferes pseudoturcicus
0.33	147 Soinferes ramosus suoso granomembrans
0.33	148 Soinferes ramosus suoso granosus
0.33	149 Soinferes ramosus suoso membranaceus
0.33	179 Microthidium cf. m. fragile
0.33	182 Cymatosphaera sp.
0.33	183 Palamosges sp. A

206	wilmer aqua percentages	207	wilmer aqua percentages	208	wilmer aqua percentages
200	33 Glaphyrocysta exuberans coxi	19 57	26 Cordosphaeridium giganteum	24 33	49 Fibracium annelorperse
207	25 Cordosphaeridium giganteum	6 67	46 Electrocyta obscuratobuata	9 33	46 Electrocyta obscuratobuata
209	46 Electrocyta obscuratobuata	11 00	53 Glaphyrocysta exuberans coxi	12 00	26 Cordosphaeridium giganteum
3 33	7 Paracaneia indentata	4 33	7 Caligodinium amicum	6 33	53 Glaphyrocysta exuberans coxi
5 37	49 Fibracium annelorperse	5 00	49 Fibracium annelorperse	5 00	176 Xenkoon austrais
6 33	3 Hystichosphaeridium tuiferum	2 67	16 Dallanona darmona	3 67	28 Cordosphaeridium nodes
2 67	80 Micrystidium cf. m. vanabae	3 33	180 Micrystidium cf. m. vanabae	3 00	34 Danae sp.
3 33	55 Florentina terox	3 33	28 Cordosphaeridium nodes	3 00	80 Micrystidium cf. m. vanabae
2 00	28 Cordosphaeridium nodes	3 00	27 Cordosphaeridium gracilis	2 67	17 Caligodinium amicum
1 00	14 Danae sp.	3 00	55 Glaphyrocysta sp. A	2 00	01 Microdinum ornatum
2 00	58 Halmisphaera seotata	3 00	58 Halmisphaera seotata	1 67	20 Chlamydomonade cf. C. urna
1 67	17 Caligodinium amicum	3 00	116 Palaeodinium pyrodonum	1 67	36 Dallanona darmona
1 67	27 Cordosphaeridium gracilis	2 57	55 Florentina terox	1 67	81 Hystichosphaeridium tuiferum
1 67	78 Xenkoon austrais	2 33	81 Hystichosphaeridium tuiferum	1 67	29 Samandia reticulifera
1 33	16 Dallanona darmona	2 00	17 Paracaneia indentata	1 33	117 Paracaneia indentata
1 33	45 Electrocyta densobaculata	1 67	101 Microdinum ornatum	1 00	45 Electrocyta densobaculata
1 33	151 Sorinertes ramosus subsp. ramosus	1 57	129 Samandia reticulifera	1 00	91 Inversotium astrimurum
1 00	31 Fromea fragilis	1 33	20 Chlamydomonade cf. C. urna	1 00	116 Palaeodinium pyrodonum
1 00	10 Oocrocodium centrocarpum	1 33	34 Danae sp.	3 67	15 Batacasphaera sp.
1 00	16 Palaeodinium pyrodonum	1 00	10 Oocrocodium centrocarpum	0 67	27 Cordosphaeridium gracilis
1 00	57 Tanosphaeridium vanecialium	1 00	51 Sorinertes ramosus subsp. ramosus	0 67	35 Florentina terox
2 57	15 Batacasphaera sp.	1 00	178 Xenkoon austrais	2 67	58 Forma A
2 67	20 Chlamydomonade cf. C. urna	2 67	23 Connimura limonata	2 67	110 Oocrocodium centrocarpum
2 67	3 Cladopyxidum saeptum	2 67	45 Electrocyta densobaculata	0 67	151 Palaeocystodinium gozowense
2 67	35 Cordosphaeridium fibrosinuosum	2 57	48 Exochosphaeridium bifidum	2 67	50 Sorinertes ramosus subsp. multibrevis
2 67	55 Glaphyrocysta sp. A	2 67	58 Forma A	2 67	51 Sorinertes ramosus subsp. ramosus
2 67	101 Microdinum ornatum	2 67	105 Nematosphaeropsis pertusa	2 67	81 Micrystidium spp.
2 67	05 Nematosphaeropsis pertusa	2 67	159 Thalassiothra delicata	0 33	23 Connimura limonata
2 67	36 Sorinertes crassipellis A	2 67	81 Micrystidium spp.	0 33	33 Cordosphaeridium calosum
2 33	5 Ancaryssea rhomboides	0 33	3 Ascotomocystus hydria	0 33	48 Exochosphaeridium bifidum
2 33	1 Ascotomocystus hydria	2 33	5 Batacasphaera sp.	0 33	51 Fromea fragilis
2 33	23 Connimura imbricata	0 33	30 Cordosphaeridium multibrevis	0 33	55 Glaphyrocysta sp. A
2 33	33 Cordosphaeridium calosum	0 33	33 Cordosphaeridium calosum	0 33	88 Halmisphaera seotata
2 33	41 Dionyes colligerum	0 33	41 Dionyes colligerum	0 33	30 Hystichosphaeridium spp.
2 33	48 Exochosphaeridium bifidum	0 33	43 Dionyes colligerum (sensu cookson 1985)	0 33	35 Impagidinium sp. A
2 33	58 Forma A	0 33	80 Hystichosphaeridium spp.	0 33	89 Impatiensphaeridium rugosum
2 33	57 Halmisphaera cf. h. seotata	0 33	39 Impatiensphaeridium rugosum	0 33	32 Isabellidium cooksoniae
2 33	37 Lentina spongiera	0 33	15 Palaeocystodinium gozowense	0 33	105 Nematosphaeropsis pertusa
2 33	08 Oligosphaeridium complex	0 33	27 Rottnesta ocellata	0 33	48 Sorinertes ramosus subsp. granosus
2 33	15 Palaeocystodinium gozowense	0 33	32 Senegalinum? dhvynense	0 33	57 Tanosphaeridium vanecialium
2 33	29 Samandia reticulifera	0 33	36 Sorinertes crassipellis A	0 33	59 Thalassiothra delicata
2 33	32 Senegalinum? dhvynense	0 33	47 Sorinertes ramosus subsp. granosus	0 33	79 Micrystidium cf. m. fragile
2 33	33 Sorinertes crassipellis A	0 33	48 Sorinertes ramosus subsp. granosus	0 33	186 Pterospemopsis spp.
2 33	48 Sorinertes ramosus subsp. granosus	0 33	149 Sorinertes ramosus subsp. membranaceus		
2 33	49 Sorinertes ramosus subsp. membranaceus	0 33	150 Sorinertes ramosus subsp. multibrevis		
2 33	52 Sorinertes ramuliferus	0 33	56 Systematophora placacantha		
2 33	58 Systematophora placacantha	0 33	62 Ngonopyxidia gneiss		
2 33	59 Thalassiothra delicata	0 33	79 Micrystidium cf. m. fragile		
2 33	60 Thalassiothra delicata	0 33	86 Pterospemopsis spp.		
2 33	61 Microdinum nitidum				
2 33	79 Micrystidium cf. m. fragile				
2 33	83 Palumbages sp. A				
2 33	86 Pterospemopsis spp.				

Correlation Matrix for PT6

Correlation Matrix for Variables: X1 ... X4

	305	306	307	308
305	1			
306	.59792	1		
307	.73364	.80327	1	
308	.92007	.62847	.75536	1

ANOVA Table for PT6

One Factor ANOVA-Repeated Measures for X1 ... X4

Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	3160.85165	12.4443	12.36574	.0001
Within subjects	765	769.86015	1.00635		
treatments	3	.08939	.0298	.0295	.9932
residual	762	769.77076	1.0102		
Total	1019	3930.7118			

Reliability Estimates for- All treatments: .91913    Single Treatment: .73968



Sample 313

31.31	water adula percentages
19.331	*881Turbinosphaera fibosa
9.671	551Glaucocystis sp A
9.001	381Deltandna darmona
6.871	*781Xenomon australe
5.331	*171Paralecanella indentata
4.671	631Glaucocystis exuberans cpx
4.001	*591Thalassiosira decata
3.331	611Fomes fragilis
2.331	371Lemna spingera
2.001	*71Calquodinium unicum
*.571	41Aiscocystis cf. A. margaria
*.331	131Ascotomocystis nydra
*.331	201Chlamydomonada cf. C. urna
*.331	461Electrocystis obscuroradiata
*.331	*781Cyclopsella veta
*.001	581Harmosphaera septata
*.001	*511Sordieria ramosa subsp. ramosa
3.671	1101Obolodinium centrocarum
3.671	*201Phaeodinium magnificum
3.671	1481Sordieria ramosa subsp. ramosa
3.671	1801Microcystidium cf. M. vanaone
3.671	*841Palamogon sp. B
3.331	211Cladopyxium saetum
0.331	271Cordosphaeridium gracile
0.331	281Cordosphaeridium nodos
0.331	371Deltandna phosonothica
0.331	451Electrocystis densobaculata
0.331	621Fomes? laevigata
0.331	731Horodogonea apiculata
0.331	901Hystichosphaeridium sp.
0.331	811Hystichosphaeridium tubiferum
0.331	891Impatiensphaeridium rugosum
0.331	*011Microdium ornatum
0.331	*051Nematosphaeropsis dentata
0.331	*071Nematosphaeropsis radeculata
0.331	*151Palaecocystidium gozowense
0.331	*361Sordieria crassipes A
0.331	*461Sordieria pseudofurcata
0.331	*821Cymatosphaera sp.
0.331	*861Parasphaera sp.

Statistical Information for Sample 313

X1: 304					
Mean:	Std. Dev.:	Std. Error:	Variance:	Coef. Var.:	Count:
.41008	2.22636	.13942	4.95669	542.91125	255
Minimum:	Maximum:	Range:	Sum:	Sum of Sqr.:	# Missing:
0	26	26	104.57	1301.8805	0

X2: 313					
Mean:	Std. Dev.:	Std. Error:	Variance:	Coef. Var.:	Count:
.41416	2.68898	.16839	7.2306	649.26547	255
Minimum:	Maximum:	Range:	Sum:	Sum of Sqr.:	# Missing:
0	39.33	39.33	105.61	1880.3115	0

PT7 "Glaphytocysta species A/ Eocladopyxis peniculatum" Association

wimer aqua percentages		wimer aqua percentages		wimer aqua percentages	
314	55 Glaphytocysta sp A	315	55 Glaphytocysta sp A	316	55 Glaphytocysta sp A
35 00	59 Thalassiochorda delicata	41 57	47 Eocladopyxis peniculatum	38 00	47 Eocladopyxis peniculatum
26 33	17 Paralecanella indentata	14 00	34 Impaginium cf. speciosum	21 33	31 Adriatosphaeridium robustum
5 00	66 Dellandria carmoria	3 33	36 Lentinia ruginosa	6 33	37 Dellandria oshonhorica
3 33	36 Dellandria carmoria	4 67	36 Dellandria carmoria	3 33	28 Coridosphaeridium inodes
1 33	58 Halimnospaera setolata	2 57	32 Senegalinum? dhymense	2 00	67 Turosophaera parastabulata
1 33	9 Inversidinium exilimum	1 57	3 Adriatosphaeridium robustum	1 67	27 Coridosphaeridium gracilis
1 00	46 Electrocysta obscuratubulata	1 47	28 Coridosphaeridium inodes	1 57	46 Electrocysta obscuratubulata
1 00	53 Glaphytocysta exuberans cox	1 33	15 Palaeocystodinium gozdownense	1 57	34 Impaginium cf. speciosum
1 00	51 Sponferites ramosus subsp. ramosus	1 33	120 Pheodinium magnificum	1 33	58 Halimnospaera setolata
1 00	81 Microstridium spp.	1 33	30 Senegalinum? obscurum	1 00	24 Coridosphaeridium biomatum
0 57	41 Alisocysta cf. a. margarita	1 00	31 Andauserella rhomboides	1 00	50 Fibrocysta bipolaris
0 57	28 Coridosphaeridium inodes	1 00	13 Ascolomocystus hydra	1 00	53 Glaphytocysta exuberans cox
0 57	41 Dionives colligerum	1 00	46 Electrocysta obscuratubulata	1 00	37 Impaginium speciosum
0 57	99 Impetosphaeridium rugosum	1 00	37 Impaginium speciosum	1 00	89 Impetosphaeridium rugosum
0 57	101 Microdinium ornatum	1 00	37 Lentinia springera	1 00	31 Kairosphaeridium brevibrachium
0 57	107 Nematospaeropsis rabeculata	1 00	67 Turosophaera parastabulata	1 00	96 Lentinia ruginosa
0 57	110 Operculodinium centrocarpum	0 57	18 Cassidium paleocenium	1 00	110 Operculodinium centrocarpum
0 57	76 Xanxon australe	0 57	24 Coridosphaeridium biomatum	1 00	126 Sponferites crassipellis A
0 57	80 Microstridium cf. m. variable	0 57	35 Coridosphaeridium fibrososum	1 00	68 Turosophaera rotunda
0 33	3 Adriatosphaeridium robustum	0 57	53 Glaphytocysta exuberans cox	0 57	8 Cassidium paleocenium
0 33	7 Caigoonidium amicum	0 57	58 Halimnospaera setolata	0 57	25 Coridosphaeridium fibrososum
0 33	8 Cassidium paleocenium	0 57	31 Hyalinosphaeridium tubiferum	0 57	45 Electrocysta densobaculata
0 33	25 Coridosphaeridium fibrososum	0 57	98 Lingulodinium machaerophorum	0 57	51 Fronia fragilis
0 33	26 Coridosphaeridium giganteum	0 57	110 Operculodinium centrocarpum	0 57	51 Fronia fragilis
0 33	27 Coridosphaeridium gracilis	0 57	117 Paralecanella indentata	0 57	51 Hyalinosphaeridium tubiferum
0 33	33 Coridosphaeridium calosum	0 57	25 Pseudonidium membraniferum	0 57	107 Nematospaeropsis rabeculata
0 33	43 Dionives colligerum (sensu Cookson 1985)	0 57	136 Sponferites crassipellis A	0 57	151 Sponferites ramosus subsp. granosus
0 33	45 Electrocysta densobaculata	0 57	146 Sponferites pseudofurcatus	0 57	151 Sponferites ramosus subsp. ramosus
0 33	51 Fronia fragilis	0 57	178 Cycloosella vieta	0 33	80 Microstridium cf. m. variable
0 33	50 Hyalinosphaeridium spp.	0 33	7 Caigoonidium amicum	0 33	51 Apectodinium Homomorphum complex
0 33	51 Hyalinosphaeridium tubiferum	0 33	20 Chlamydomonada cf. C. urna	0 33	11 Apectodinium robustum
0 33	37 Lentinia springera	0 33	21 Cladopyxidium saeptum	0 33	7 Caigoonidium amicum
0 33	20 Pheodinium magnificum	0 33	27 Coridosphaeridium gracilis	0 33	21 Cladopyxidium saeptum
0 33	25 Pseudonidium membraniferum	0 33	37 Dellandria oshonhorica	0 33	33 Coridosphaeridium calosum
0 33	27 Rottnechia borussica	0 33	41 Dionives colligerum	0 33	41 Dionives colligerum
0 33	46 Sponferites pseudofurcatus	0 33	45 Electrocysta densobaculata	0 33	77 Hyalinosphaeridium tumescens
0 33	52 Trigonopyxidia girella	0 33	51 Fronia fragilis	0 33	80 Hyalinosphaeridium spp.
0 33	79 Microstridium cf. m. fragile	0 33	76 Hyalinosphaeridium rugosum	0 33	115 Palaeocystodinium gozdownense
0 33	84 Psalimoides sp. B	0 33	77 Hyalinosphaeridium tumescens	0 33	125 Pseudonidium membraniferum
0 33	86 Pteroppermopsis spp.	0 33	99 Impetosphaeridium rugosum	0 33	27 Rottnechia borussica
		0 33	33 Kairosphaeridium brevibrachium	0 33	32 Senegalinum? dhymense
		0 33	91 Microdinium ornatum	0 33	37 Sponferites crassipellis B
		0 33	105 Nematospaeropsis pertusa	0 33	49 Sponferites ramosus subsp. membranaceo
		0 33	107 Nematospaeropsis rabeculata	0 33	79 Cycloosella vieta
		0 33	127 Rottnechia borussica	0 33	79 Microstridium cf. m. fragile
		0 33	137 Sponferites crassipellis B	0 33	81 Microstridium spp.
		0 33	148 Sponferites ramosus subsp. granosus		
		0 33	149 Sponferites ramosus subsp. membranaceo		
		0 33	68 Turosophaera rotunda		
		0 33	79 Microstridium cf. m. fragile		
		0 33	80 Microstridium cf. m. variable		
		0 33	81 Microstridium spp.		

	wtmer aqua percentages
25.331	551Glaucocystis sp. A
18.671	31Acanthosphaeridium robustum
13.001	461Electrocysta obscuritabulata
5.671	371Imoagidium speciosum
5.331	371Diatonia pinozonica
4.001	471Ectacoopyxis pectinatum
3.671	451Electrocysta pectinatum
2.671	1321Senecogonium? dihyminense
2.331	181Cassidium paleocenium
1.671	51Anaxissella monoonidra
1.671	931Kairosphaeridium brevisaratum
1.671	1011Microdinium ornatum
1.671	1301Senecogonium? obscurum
1.001	81Aeetodinium Homomorphum complex
1.001	1251Rensidium membraniferum
1.001	1681Turboisphaera rotunda
0.671	111Aetodinium retortum
0.671	201Chlamydomonella cf. C. jma
0.671	241Cordosphaeridium biarmatum
0.671	271Cordosphaeridium gracile
0.671	281Cordosphaeridium inodes
0.671	341Imoagidium cf. l. speciosum
0.671	311Inversidium extimurum
0.671	361Lentrea ruginosa
0.671	1151Paleocystodinium golzowense
0.671	1431Solenites cornutus C
0.671	1671Turboisphaera paritabulata
0.671	1811Microdinium spp.
0.331	211Cyclopyxidium saeptum
0.331	431Domes collicerum (sensu cookson 1985)
0.331	511Forma fragile
0.331	831Glaucocystis esuberans colx
0.331	731Heteroginella apiculata
0.331	771Heteroginella tumescens
0.331	1071Nematospaeropsis tabaculata
0.331	1201Pheodinium magnificum
0.331	1271Rothestia borussica
0.331	1481Solenites ramosus suoso. granosus
0.331	1511Solenites ramosus suoso. ramosus
0.331	1621Tropocoida gineia
0.331	1781Cycloisella vieta
0.331	1791Microdinium cf. m. fragile
0.331	1801Microdinium cf. m. variae
0.331	1821Cymatosphaera spp.
0.331	1861Pterospermoisa spp.

Correlation Matrix for PT7

Correlation Matrix for Variables: X<sub>1</sub> ... X<sub>4</sub>

	314	315	316	317
314	1			
315	.69485	1		
316	.64003	.95364	1	
317	.53587	.72663	.76434	1

ANOVA Table for PT7

One Factor ANOVA-Repeated Measures for X<sub>1</sub> ... X<sub>4</sub>

Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	5756.58743	22.66373	10.96607	.0001
Within subjects	765	1581.0357	2.06671		
treatments	3	.02361	.00787	.00379	.9997
residual	762	1581.01209	2.07482		
Total	1019	7337.62313			

Reliability Estimates for- All treatments: .90881    Single Treatment: .71359

Sample number 318

1 91		wilmeri sp. percentages
31 20	3	<i>Apectoedinium</i> monomorphum complex
39 20	46	<i>Electrocysta</i> obscuroradiata
4 57	97	<i>Turboedonera</i> parvifurcata
4 20	45	<i>Electrocysta</i> densoculata
3 57	3	<i>Asciotricocystus</i> nuda
3 20	33	<i>Kaliosphaeridium</i> brevicaudatum
1 57	81	<i>Froesia</i> fragilis
1 33	36	<i>Lentina</i> rugosa
1 33	15	<i>Paraeocystodinium</i> gottohwense
1 33	137	<i>Soriferites</i> crassipellis B
1 00	23	<i>Cordosphaeridium</i> nodes
1 00	55	<i>Glyptrocysta</i> sp. A
1 00	37	<i>Impatioidinium</i> speciosum
1 00	97	<i>Lentina</i> svingeri
1 00	98	<i>Unquiodinium</i> machaerophorum
1 20	01	<i>Microdinium</i> ornatum
1 20	17	<i>Paralecanella</i> indentata
1 20	68	<i>Turboedonera</i> rotunda
2 67	11	<i>Apctoedinium</i> rotatum
2 57	20	<i>Chlamydoxonella</i> cf. <i>C. urna</i>
2 67	47	<i>Ectocystus</i> pumiliatum
2 67	53	<i>Fibrocysta</i> radiata
2 67	30	<i>Hysteroedonidium</i> spp.
2 67	31	<i>Hysteroedonidium</i> tubiferum
2 67	107	<i>Nematodesma</i> radiatum
2 67	10	<i>Coercuodinium</i> centrocarum
2 67	30	<i>Senecidium</i> ? obscurum
2 67	51	<i>Soriferites</i> ramosus subsp. ramosus
2 67	69	<i>Wetzelaella</i> cf. <i>w. mythenis</i>
2 67	80	<i>Micromystrium</i> cf. <i>m. vanaae</i>
2 67	81	<i>Micromystrium</i> spp.
2 67	84	<i>Palaeodinium</i> sp. B
2 33	3	<i>Aquilosphaeridium</i> robustum
2 33	5	<i>Anaulusella</i> rhomboides
2 33	18	<i>Cassidulum</i> paleoboreum
2 33	24	<i>Cordosphaeridium</i> palmatum
2 33	25	<i>Cordosphaeridium</i> fibrosinuosum
2 33	30	<i>Cordosphaeridium</i> mutuosinuosum
2 33	33	<i>Cordosphaeridium</i> calosum
2 33	17	<i>Dalmanella</i> oliviformis
2 33	42	<i>Diphyes</i> colligatum
2 33	50	<i>Fibrocysta</i> ovalare
2 33	57	<i>Halmosphaera</i> cf. <i>h. seotata</i>
2 33	58	<i>Halmosphaera</i> seotata
2 33	76	<i>Hysteroedonidium</i> rugosum
2 33	77	<i>Hysteroedonidium</i> tumescens
2 33	78	<i>Hysteroedonidium</i> uncinatum
2 33	39	<i>Impatioidinium</i> rugosum
2 33	31	<i>Inversidinium</i> aculeatum
2 33	32	<i>Senecidium</i> ? oliveense
2 33	136	<i>Soriferites</i> crassipellis A
2 33	43	<i>Soriferites</i> cornutus C
2 33	46	<i>Soriferites</i> pseudohurcatus
2 33	78	<i>Cyclopsella</i> vieta

PT8 "Adnatosphaeridium multispinosum/ Apectodinium homomorphum"  
Association

324	324 witmer, Nant. zone 6	325	325 witmer, Nant. zone 6	329	witmer nanemov percentages
29.20	2 Adnatosphaeridium multispinosum	28.33	2 Adnatosphaeridium multispinosum	20.00	2 Adnatosphaeridium multispinosum
23.33	5 Apectodinium homomorphum complex	22.00	5 Apectodinium homomorphum complex	3.33	1 Anomigera sp.
4.67	12 Senegalinium? diiwyense	15.33	132 Senegalinium? diiwyense	2.33	156 Systematopora placacantha
4.67	12 Coerculodinium israelianum	4.33	103 Muratodinium limbratum	2.00	51 Spiniferites ramosus subsp. ramosus
3.33	103 Muratodinium limbratum	4.00	151 Spiniferites ramosus subsp. ramosus	1.57	172 Wertheimella samianica
3.00	281 Cordosphaeridium inodes	3.33	281 Cordosphaeridium inodes	1.33	37 Dellandria phosporatica
2.67	391 Dinopterygium cladioides	3.33	112 Coerculodinium israelianum	1.33	47 Ectoocopyxis penicillatum
2.33	107 Nematosphaerosopsis irabeculata	2.57	107 Nematosphaerosopsis irabeculata	1.00	5 Apectodinium Homomorphum complex
2.33	151 Spiniferites ramosus subsp. ramosus	1.67	130 Senegalinium? obscurum	1.00	103 Muratodinium limbratum
1.00	58 Halmisphaera septata	1.33	39 Dinopterygium cladioides	2.57	27 Cordosphaeridium gracillius
1.00	32 Millouodinium guiseppi major	1.33	58 Halmisphaera septata	2.57	41 Dionyes colligerum
1.00	91 Micrystidium spp.	1.33	39 Impetosphaeridium rugosum	2.67	45 Ectocysta densobaculata
0.67	37 Dellandria phosporatica	1.30	27 Cordosphaeridium gracillius	2.67	58 Forma C
0.67	130 Senegalinium? obscurum	1.00	150 Spiniferites ramosus subsp. multibrevis	2.57	46 Glaphrocysta sp B
0.67	160 Thalassiphora pelagica	0.67	33 Cordosphaeridium caliosum	0.67	37 Lentima springera
0.33	27 Cordosphaeridium gracillius	0.67	51 Fibrocysta lappacea	0.67	102 Millouodinium guiseppi major
0.33	291 Cordosphaeridium inodes robustum	0.67	110 Coerculodinium centrocarpum	0.67	107 Nematosphaerosopsis irabeculata
0.33	301 Cordosphaeridium multispinosum	0.67	181 Micrystidium spp.	0.67	149 Spiniferites ramosus subsp. membranaceo
0.33	33 Cordosphaeridium caliosum	0.33	30 Cordosphaeridium multispinosum	0.67	170 Wertheimella rampdenensis
0.33	41 Dionyes colligerum	0.33	37 Dellandria phosporatica	0.33	281 Cordosphaeridium inodes
0.33	42 Dionyes colligerum	0.33	42 Dionyes colligerum	0.33	10 Cordosphaeridium multispinosum
0.33	50 Fibrocysta bipolare	0.33	47 Ectoocopyxis penicillatum	0.33	31 Cordosphaeridium soaster
0.33	51 Fibrocysta lappacea	0.33	50 Fibrocysta bipolare	0.33	39 Dinopterygium cladioides
0.33	53 Glaphrocysta exuberans cox	0.33	55 Glaphrocysta sp A	0.33	43 Dionyes colligerum (sensu Cookson 1985)
0.33	30 Hysterochloa spp.	0.33	78 Hysterochloa unispinum	0.33	46 Ectocysta obscurotubulata
0.33	89 Impetosphaeridium rugosum	0.33	80 Hysterochloa spp.	0.33	33 Fibrocysta radiata
0.33	33 Kalosphaeridium brevibarbatum	0.33	98 Linguodinium macnaerophorum	0.33	57 Forma B
0.33	37 Lentima springera	0.33	102 Millouodinium guiseppi major	0.33	53 Glaphrocysta exuberans cox
0.33	98 Linguodinium macnaerophorum	0.33	125 Rensidium membraniferum	0.33	55 Glaphrocysta sp A
0.33	10 Coerculodinium centrocarpum	0.33	127 Rottmestria borussica	0.33	58 Halmisphaera septata
0.33	25 Rensidium membraniferum	0.33	136 Spiniferites crassipellis A	0.33	76 Hysterochloa rugosa
0.33	46 Spiniferites pseudofurcatus	0.33	145 Spiniferites monilis	0.33	77 Hysterochloa tumescens
0.33	147 Spiniferites ramosus subsp. granomembrans	0.33	146 Spiniferites pseudofurcatus	0.33	81 Hysterochloa tubiferum
0.33	148 Spiniferites ramosus subsp. granosus	0.33	147 Spiniferites ramosus subsp. granomembrans	0.33	33 Kalosphaeridium brevibarbatum
0.33	150 Spiniferites ramosus subsp. multibrevis	0.33	148 Spiniferites ramosus subsp. granosus	0.33	36 Lentima rugosa
0.33	161 Trichodinium hirsutum	0.33	160 Thalassiphora pelagica	0.33	98 Linguodinium macnaerophorum
0.33	182 Cymatosphaera spp.	0.33	161 Trichodinium hirsutum	0.33	391 Meliosphaeridium pseudocurvatum
0.33	186 Pterospemopsis spp.	0.33	182 Cymatosphaera spp.	0.33	112 Coerculodinium israelianum
		0.33	186 Pterospemopsis spp.	0.33	151 Palaeocystodinium gozowense
				0.33	171 Paralecanella inornata
				0.33	124 Polysphaeridium zonari
				0.33	125 Rensidium membraniferum
				0.33	127 Rottmestria borussica
				0.33	136 Spiniferites crassipellis A
				0.33	145 Spiniferites monilis
				0.33	146 Spiniferites pseudofurcatus
				0.33	158 Tectadodinium psillum
				0.33	160 Thalassiphora pelagica
				0.33	175 Wilsonidium tabulatum
				0.33	191 Micrystidium spp.
				0.33	186 Pterospemopsis spp.

126:	326	wilmer, Nani, zone 7
4.33:	*75:	Wilsonidium tabulatum
3.57:	37:	Deilanona gnosphornica
3.33:	*02:	Milioidodinium guiseppi maior
2.57:	*03:	Muratodinium limbratum
2.33:	2:	Adnatospaeridium mutispinosum
2.00:	*32:	Senegalinum? dilwynense
1.57:	33:	Kaliosphaeridium brevicastratum
1.57:	97:	Lentina spinigera
1.33:	46:	Electrocysta obscuratolobata
*.00:	*10:	Operculodinium centrocarpum
1.00:	*51:	Spiniferes ramosus subsp. ramosus
*.00:	*81:	Micrystidium spp.
3.57:	50:	Fibrocysta bipolare
3.57:	*07:	Nematospaeropsis iraeoculata
3.57:	56:	Systematophora placacantha
3.57:	86:	Pterospermopsis spp.
3.33:	20:	Chlamydomonadea cf. C. uma
3.33:	25:	Cordosphaeridium fibrosinuosum
3.33:	27:	Cordosphaeridium gracilis
3.33:	28:	Cordosphaeridium inodes
3.33:	39:	Dinodieridium cladodes
3.33:	41:	Dionyes colligerum
3.33:	42:	Dionyes colligerum
3.33:	43:	Dionyes colligerum (sensu Cookson 1985)
3.33:	53:	Glaucocysta exuberans cxu
3.33:	58:	Halmisphaera septata
3.33:	76:	Hysterochloa riguada
3.33:	81:	Hysterochloa tubiferum
3.33:	39:	Impatiensphaeridium rugosum
3.33:	31:	Inversidium exilimurum
3.33:	36:	Lentina ruginosa
3.33:	38:	Linguodinium machaerophorum
3.33:	39:	Melissosphaeridium pseudocurvatum
3.33:	*01:	Microdinium ornatum
3.33:	*17:	Paralecanella indentata
3.33:	*25:	Rendinium membraniferum
3.33:	*30:	Senegalinum? obscurum
3.33:	*60:	Thalassiphora delagrea
3.33:	*82:	Cymatosphaera spp.

327	327	wilmer, Nani, zone 7
4.57:	*56:	Systematophora placacantha
5.57:	57:	Forma B
4.57:	*51:	Spiniferes ramosus subsp. ramosus
4.00:	31:	Hysterochloa tubiferum
3.33:	*17:	Paralecanella indentata
3.33:	*32:	Senegalinum? dilwynense
2.57:	2:	Adnatospaeridium mutispinosum
2.33:	41:	Dionyes colligerum
2.33:	*07:	Nematospaeropsis iraeoculata
2.00:	37:	Deilanona gnosphornica
1.57:	38:	Linguodinium machaerophorum
1.57:	*02:	Milioidodinium guiseppi maior
1.33:	97:	Lentina spinigera
1.33:	*36:	Spiniferes crassipellis A
1.00:	45:	Electrocysta densoculata
1.00:	50:	Fibrocysta bipolare
1.00:	*10:	Operculodinium centrocarpum
0.57:	28:	Cordosphaeridium inodes
0.57:	42:	Dionyes colligerum
0.57:	46:	Electrocysta obscuratolobata
0.57:	47:	Eocadopyxis paniculatum
0.57:	103:	Muratodinium limbratum
0.57:	*50:	Spiniferes ramosus subsp. multibrevis
0.57:	*81:	Micrystidium spp.
0.33:	5:	Apectodinium Homomorpnium complex
0.33:	20:	Chlamydomonadea cf. C. uma
0.33:	25:	Cordosphaeridium fibrosinuosum
0.33:	27:	Cordosphaeridium gracilis
0.33:	33:	Cordosphaeridium calosum
0.33:	78:	Hysterochloa riguada
0.33:	93:	Kaliosphaeridium brevicastratum
0.33:	96:	Lentina ruginosa
0.33:	99:	Melissosphaeridium pseudocurvatum
0.33:	*12:	Operculodinium israelianum
0.33:	*15:	Palaeocystodinium gozowense
0.33:	*25:	Rendinium membraniferum
0.33:	*46:	Spiniferes pseudofurcatus
0.33:	147:	Spiniferes ramosus subsp. granomemoran
0.33:	*48:	Spiniferes ramosus subsp. granosus
0.33:	*49:	Spiniferes ramosus subsp. membranaceol
0.33:	*58:	Tectatodinium peltitum
0.33:	*75:	Wilsonidium tabulatum
0.33:	*78:	Cyclopsella vieta
0.33:	*82:	Cymatosphaera spp.
0.33:	*86:	Pterospermopsis spp.



Correlation Matrix for PT8

Correlation Matrix for Variables: X<sub>1</sub> ... X<sub>3</sub>

	324	325	328
324	1		
325	.99585	1	
328	.68368	.68571	1

ANOVA Table for PT8

One Factor ANOVA-Repeated Measures for X<sub>1</sub> ... X<sub>3</sub>

Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	3683.1426	14.50056	12.34664	.0001
Within subjects	510	598.9718	1.17445		
treatments	2	2.91387	1.45694	1.2417	.2898
residual	508	596.05793	1.17334		
Total	764	4282.1144			

Reliability Estimates for- All treatments: .91901    Single Treatment: .79089

PT9 "Wetzeliella hampdenensis/ Adnatosphaeridium multispinosum/  
Thalassiphora pelagica" Association

329	329	wimer, Nani, zone 7
0.30	70	Wetzeliella hampdenensis
17.00	2	Adnatosphaeridium multispinosum
9.67	160	Thalassiphora pelagica
9.00	28	Corcosphaeridium inodes
5.97	71	Homotryblum tasmanense
5.33	56	Glaphyrocysta sp. B
5.00	58	Forma C
4.33	45	Electrocysta densobaculata
4.33	47	Eociadopyxis oeniculatum
4.33	72	Wetzeliella samlandica
2.97	151	Spiriferites ramosus subsp. ramosus
2.33	98	Linguodinium machaerophorum
1.67	3	Adnatosphaeridium robustum
1.67	12	Arcoligera sp.
1.67	27	Corcosphaeridium gracilis
1.67	41	Dionyes colligerum
1.67	57	Forma B
1.67	55	Glaphyrocysta sp. A
1.33	42	Dionyes colligerum
1.33	37	Lentina spongera
1.00	58	Halmisphaera septata
1.00	06	Nematosphaeropsis pousuosa
1.00	07	Nematosphaeropsis irabeculata
1.00	10	Oercuodinium centrocarum
1.00	27	Rothnesia borussica
1.00	81	Micrystidium spp.
0.67	25	Corcosphaeridium librosinosum
0.67	30	Corcosphaeridium multispinosum
0.67	39	Dinoerygium claidodes
0.67	46	Electrocysta obscuratoulata
0.67	31	Hysterochaetidium tubiferum
0.67	36	Lentina ruginosa
0.67	39	Mentosphaeridium pseudocurvatum
0.67	02	Millooudinium guiseppi maior
0.67	126	Rendinium sp. A
0.67	145	Spiriferites montis
0.67	46	Spiriferites pseudofurcatus
0.67	47	Spiriferites ramosus subsp. granomembrana
0.67	48	Spiriferites ramosus subsp. granosus
0.67	49	Spiriferites ramosus subsp. membranaceus
0.33	158	Tectadodinium pelitum
0.33	1	Achnella bifurcata n. gen. n. comb.
0.33	51	Oercuodinium homomorphum complex
0.33	29	Corcosphaeridium inodes robustum
0.33	33	Corcosphaeridium caliosum
0.33	43	Dionyes colligerum (sensu cookson '985)
0.33	48	Erochrosphaeridium bifidum
0.33	50	Fibrocysta bicolare
0.33	51	Fibrocysta iapopaea
0.33	52	Fibrocysta coarctosinosa
0.33	53	Fibrocysta radiata
0.33	76	Hysterochaetopoma rugade
0.33	77	Hysterochaetopoma tumescens
0.33	80	Hysterochaetidium spp.
0.33	39	Imetosphaeridium rugosum
0.33	33	Kaliosphaeridium brevisaratum
0.33	03	Muratodinium limonatum
0.33	11	Oercuodinium israelianum
0.33	115	Palaeocystodinium goizowense
0.33	17	Paralecanella indentata
0.33	23	Polysphaeridium cf. p. zonaryi
0.33	24	Polysphaeridium zonaryi
0.33	30	Senegalinum? obscurum
0.33	36	Spiriferites crassioellus A
0.33	50	Spiriferites ramosus subsp. multibrevis
0.33	53	Spiriferites A
0.33	61	Trochodinium firmum
0.33	65	Tuodermodinium sulcatum
0.33	75	Wilsonodinium tabulatum

331	331	wimer, Nani, zone 8
9.00	2	Adnatosphaeridium multispinosum
12.67	47	Eociadopyxis oeniculatum
12.67	58	Forma C
10.33	174	Wetzeliella vancouveritica
9.00	3	Adnatosphaeridium robustum
4.67	117	Paralecanella indentata
3.33	98	Linguodinium machaerophorum
3.00	45	Electrocysta densobaculata
2.67	103	Muratodinium limonatum
2.67	72	Wetzeliella samlandica
2.00	70	Wetzeliella hampdenensis
1.67	124	Polysphaeridium zonaryi
1.67	160	Thalassiphora pelagica
1.00	12	Arcoligera sp.
1.00	37	Dellandra prosaeritica
1.00	46	Electrocysta obscuratoulata
0.67	18	Cassidium paleocenicum
0.67	53	Fibrocysta radiata
0.67	59	Heteroculacysta campanula
0.67	75	Hysterochaetopoma rugade
0.67	80	Hysterochaetidium spp.
0.67	102	Millooudinium guiseppi maior
0.67	107	Nematosphaeropsis irabeculata
0.67	109	Oercuodinium brevisinosum
0.67	110	Oercuodinium centrocarum
0.67	22	Phenacodinium echinatum
0.67	151	Spiriferites ramosus subsp. ramosus
0.67	158	Tectadodinium pelitum
0.67	179	Cyclosetella vista
0.33	20	Chlamydocoelata cf. C. urna
0.33	27	Corcosphaeridium gracilis
0.33	28	Corcosphaeridium inodes
0.33	29	Corcosphaeridium inodes robustum
0.33	33	Corcosphaeridium caliosum
0.33	39	Dinoerygium claidodes
0.33	41	Dionyes colligerum
0.33	43	Dionyes colligerum (sensu cookson '985)
0.33	48	Erochrosphaeridium bifidum
0.33	68	Halmisphaera septata
0.33	71	Homotryblum tasmanense
0.33	77	Hysterochaetopoma tumescens
0.33	31	Hysterochaetidium tubiferum
0.33	33	Kaliosphaeridium brevisaratum
0.33	36	Lentina ruginosa
0.33	100	Membranifilmacia leptoderma
0.33	111	Oercuodinium cf. d. brevisinosum
0.33	112	Oercuodinium israelianum
0.33	115	Palaeocystodinium goizowense
0.33	123	Polysphaeridium cf. p. zonaryi
0.33	126	Rendinium sp. A
0.33	136	Spiriferites crassioellus A
0.33	145	Spiriferites montis
0.33	146	Spiriferites pseudofurcatus
0.33	148	Spiriferites ramosus subsp. granosus
0.33	149	Spiriferites ramosus subsp. membranaceus
0.33	153	Spiriferites A
0.33	65	Tuodermodinium sulcatum
0.33	81	Micrystidium spp.
0.33	185	Palambages sp. C

Correlation Matrix for PT9

Correlation Matrix for Variables: X<sub>1</sub> ... X<sub>2</sub>

	329	331
329	1	
331	.4135	1

ANOVA Table for PT9

One Factor ANOVA-Repeated Measures for X<sub>1</sub> ... X<sub>2</sub>

Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	1926.28088	7.58378	2.23858	.0001
Within subjects	255	863.88175	3.38777		
treatments	1	2.4948	2.4948	.73565	.3919
residual	254	861.38695	3.39129		
Total	509	2790.16263			

Reliability Estimates for- All treatments: .55329    Single Treatment: .38244

PT10 "Wetziella samlandica" Association

3301	330	witmer, Nani, zone δ
32.301	72	Wetziella samlandica
6.57	57	Forma B
6.57	45	Electrocysta densobaculata
6.57	60	Thaassiphora pelagica
6.57	58	Forma C
6.57	170	Wetziella ramodensis
6.57	174	Wetziella vaneiongtuda
1.331	46	Electrocysta obscurotabulata
1.331	47	Ectocarpus penicillatus
1.331	2	Adnatosphaeridium multispinosum
1.331	51	Spiniferites ramosus subsp. ramosus
3.571	21	Adnatosphaeridium robustum
3.571	27	Corcosphaeridium gracilis
3.571	28	Corcosphaeridium nodos
3.571	37	Deltanona pnosphontica
3.571	96	Lentina ruginosa
3.571	97	Lentina spinigera
3.571	98	Linguodinium machaerophorum
3.571	101	Microdinium ornatum
3.571	107	Nematospaeropsis irabeculata
3.571	115	Palaeocystodinium goizowense
3.571	122	Phthanopendinium echinatum
3.571	165	Tubodermoidium sulcatum
3.571	181	Micrystidium sp.
3.571	186	Pterospemopsis sp.
3.331	1	Achnelia biformoides n. gen., n. comb.
3.331	51	Apectodinium homomorphy complex
3.331	12	Aneoligera sp.
3.331	25	Corcosphaeridium fibrosinosum
3.331	43	Olenes colligerum (sensu cooxson 1985)
3.331	52	Fibrocysta coarctospinosa
3.331	71	Homotrypium tasmanense
3.331	78	Hysterochloa rigida
3.331	80	Hysterochloa sp.
3.331	91	Inversidinium exilimurum
3.331	93	Kallosphaeridium brevispinatum
3.331	100	Membranillamaca leptodema
3.331	102	Miloudocodium guseepi major
3.331	103	Muratodinium limbatum
3.331	110	Coercuodinium centrocarum
3.331	112	Oercuodinium israelianum
3.331	117	Paraeacanthia indentata
3.331	124	Polysphaeridium zonari
3.331	125	Rendinium membraniferum
3.331	125	Rendinium membraniferum
3.331	127	Rottnesia berussica
3.331	132	Seneganium? dilwynense
3.331	146	Spiniferites pseudoturcatus
3.331	149	Spiniferites ramosus subsp. membranaceo
3.331	153	Spiniferites A
3.331	162	Trochocoxys gineia
3.331	175	Wilsonidium tabulatum
3.331	179	Cyclopsella vieta
3.331	182	Cymatosphaera sp.
3.331	185	Palambages sp. C

3321	332	witmer, Nani, zone δ
51.301	172	Wetziella samlandica
6.671	181	Cassidium paleocanicum
3.331	21	Adnatosphaeridium multispinosum
5.331	31	Apectodinium australense
3.001	11	Achnelia biformoides n. gen., n. comb.
1.331	12	Aneoligera sp.
1.331	47	Ectocarpus penicillatus
1.331	100	Membranillamaca leptodema
1.331	151	Spiniferites ramosus subsp. ramosus
1.331	170	Wetziella ramodensis
1.001	40	Dinoperygum lennartense
1.001	76	Hysterochloa rigida
1.001	103	Muratodinium limbatum
3.671	28	Corcosphaeridium nodos
0.671	53	Fibrocysta radiata
3.571	98	Linguodinium machaerophorum
0.671	181	Micrystidium sp.
0.671	186	Pterospemopsis sp.
0.331	41	Apectodinium homomorphy complex
3.331	27	Corcosphaeridium gracilis
3.331	37	Deltanona pnosphontica
3.331	43	Olenes colligerum (sensu cooxson 1985)
0.331	45	Electrocysta densobaculata
0.331	46	Electrocysta obscurotabulata
0.331	50	Fibrocysta bipolare
0.331	58	Forma C
0.331	71	Homotrypium tasmanense
0.331	80	Hysterochloa sp.
0.331	89	Imetosphaeridium rugosum
0.331	91	Inversidinium exilimurum
0.331	93	Kallosphaeridium brevispinatum
0.331	97	Lentina spinigera
0.331	101	Microdinium ornatum
0.331	102	Miloudocodium guseepi major
0.331	107	Nematospaeropsis irabeculata
0.331	109	Oercuodinium brevispinosum
0.331	110	Oercuodinium centrocarum
0.331	115	Palaeocystodinium goizowense
0.331	117	Paraeacanthia indentata
0.331	122	Phthanopendinium echinatum
0.331	123	Polysphaeridium cf. o. zonari
0.331	124	Polysphaeridium zonari
0.331	125	Rendinium membraniferum
0.331	132	Seneganium? dilwynense
0.331	146	Spiniferites pseudoturcatus
0.331	150	Spiniferites ramosus subsp. multibrevis
0.331	160	Thaassiphora pelagica
0.331	165	Tubodermoidium sulcatum
0.331	182	Cymatosphaera sp.
0.331	185	Palambages sp. C

Correlation Matrix for PT10

Correlation Matrix for Variables: X<sub>1</sub> ... X<sub>2</sub>

	330	332
330	1	
332	.71924	1

ANOVA Table for PT10

One Factor ANOVA-Repeated Measures for X<sub>1</sub> ... X<sub>2</sub>

Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	3954.2721	15.568	5.5413	.0001
Within subjects	255	716.4093	2.80945		
treatments	1	.00311	.00311	.0011	.9735
residual	254	716.40619	2.8205		
Total	509	4670.6814			

Reliability Estimates for- All treatments: .81954    Single Treatment: .69425

PT11 "Spinidinium macmurdoense / Forma C" Association

333	333 wimer, Nanj. zone 3	334	334 wimer, Nanj. zone 3
40.57	34 Spinidinium macmurdoense	32.00	134 Spinidinium macmurdoense
19.33	58 Forma C	21.33	58 Forma C
7.57	85 Palaeobages sp. C	10.00	185 Palaeobages sp. C
2.57	47 Eociadopyxis peniculatum	3.00	160 Thalassionora pelagica
2.00	37 Dallanona phosporatica	2.57	37 Dallanona phosporatica
1.57	30 Senegalinum? obscurum	2.33	170 Wetzeiella namodenensis
1.33	2 Aroeligera sp.	2.00	12 Aroeligera sp.
1.00	2 Adnatospaendium mutispinosum	1.57	158 Tectadodinium pelitum
1.00	34 Kisselovia coelothrypta	1.33	151 Spinifertes ramosus subsp. ramosus
1.00	38 Linguodinium machaerophorum	1.00	72 Homotrybium pallidum
1.00	13 Coercuodinium mutispinosum	1.00	33 Kalkosphaendium brevispinosum
1.00	7 Paralecanella indentata	1.00	98 Linguodinium machaerophorum
1.00	22 Phthanopendinium echinatum	1.00	99 Obercuodinium brevispinosum
1.00	60 Thalassionora pelagica	1.00	117 Paralecanella indentata
0.57	1 Achnella bifurcoides n. gen., n. comb.	1.00	130 Senegalinum? obscurum
0.57	38 Dallanona wardnerensis	1.00	132 Senegalinum? diuynense
0.57	40 Dinotterygium lehmannense	1.00	171 Wetzeiella unans
0.57	59 Heteraulacacysta campanula	0.57	1 Achnella bifurcoides n. gen., n. comb.
0.57	71 Homotrybium tasmanense	0.57	22 Cliestosphaendium diversispinosum
0.57	72 Homotrybium pallidum	0.57	29 Cordosphaendium inodes
0.57	33 Kalkosphaendium brevispinosum	0.57	38 Dallanona wardnerensis
0.57	09 Coercuodinium brevispinosum	0.57	41 Diphyes colligerum
0.57	31 Senegalinum? asymmetricum	0.57	46 Electrocyta obscurotabulata
0.57	32 Senegalinum? diuynense	0.57	47 Eociadopyxis peniculatum
0.57	51 Spinifertes ramosus subsp. ramosus	0.57	52 Fibrocysta coarctosponosa
0.57	73 Wetzeiella namodenensis	0.57	59 Forma D
0.57	72 Wetzeiella samianica	0.57	71 Homotrybium tasmanense
0.57	78 Cycloosella vieta	0.57	74 Hystriochlopora esenackii
0.33	0 Apieodinium labyrinthum	0.57	34 Kisselovia coelothrypta
0.33	20 Chlamydochorella cf. C. urna	0.57	13 Obercuodinium mutispinosum
0.33	22 Cliestosphaendium diversispinosum	0.57	18 Paucisphaendium inversibuccinum
0.33	25 Cordosphaendium fibrososum	0.57	128 Samianella chlamydochora
0.33	27 Cordosphaendium gracilis	0.57	131 Senegalinum? asymmetricum
0.33	29 Cordosphaendium inodes	0.57	165 Tubidermodinium sucatum
0.33	33 Cordosphaendium calosum	0.33	3 Asteodinium cribosum
0.33	39 Dinotterygium claddoes	0.33	25 Cordosphaendium fibrososum
0.33	41 Diphyes colligerum	0.33	27 Cordosphaendium gracilis
0.33	42 Diphyes colligerum	0.33	39 Dinotterygium claddoes
0.33	43 Diphyes colligerum (sensu cookson 1985)	0.33	40 Dinotterygium lehmannense
0.33	46 Electrocyta obscurotabulata	0.33	42 Diphyes colligerum
0.33	70 Homotrybium calcicum	0.33	53 Fibrocysta radiata
0.33	74 Hystriochlopora esenackii	0.33	59 Heteraulacacysta campanula
0.33	3 Inversidinium axillurum	0.33	76 Hystriochlopora riguada
0.33	0 Membranilamacia leptodema	0.33	30 Hystriochlopora endium sop.
0.33	02 Milliodonum guiseppi major	0.33	93 Hystriochlopora ovum
0.33	10 Obercuodinium centrocarpum	0.33	91 Inversidinium axillurum
0.33	12 Obercuodinium israelianum	0.33	37 Lentinia springera
0.33	15 Paucisphaendium gozowense	0.33	10 Obercuodinium centrocarpum
0.33	18 Paucisphaendium inversibuccinum	0.33	22 Phthanopendinium echinatum
0.33	23 Polysphaendium cf. p. zoharyi	0.33	23 Polysphaendium cf. p. zoharyi
0.33	28 Samianella chlamydochora	0.33	24 Polysphaendium zoharyi
0.33	36 Spinifertes crassipellis A	0.33	127 Rotmestia porusica
0.33	45 Spinifertes monilus	0.33	136 Spinifertes crassipellis A
0.33	46 Spinifertes pseudofurcatus	0.33	146 Spinifertes pseudofurcatus
0.33	48 Spinifertes ramosus subsp. granosus	0.33	149 Spinifertes ramosus subsp. membranaceus
0.33	49 Spinifertes ramosus subsp. membranaceus	0.33	172 Wetzeiella samianica
0.33	50 Spinifertes ramosus subsp. multibravis	0.33	175 Wilsonidium tabuiatum
0.33	65 Tubidermodinium sucatum	0.33	178 Cycloosella vieta
0.33	71 Wetzeiella unans		

Correlation Matrix for PT11

Correlation Matrix for Variables: X<sub>1</sub> ... X<sub>2</sub>

	333	334
333	1	
334	.97969	1

ANOVA Table for PT11

One Factor ANOVA-Repeated Measures for X<sub>1</sub> ... X<sub>2</sub>

Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	3609.80249	14.21182	67.7084	.0001
Within subjects	255	53.52385	.2099		
treatments	1	.0056	.0056	.02658	.8706
residual	254	53.51825	.2107		
Total	509	3663.32634			

Reliability Estimates for- All treatments: .98523    Single Treatment: .97089

PT12 "Deflandria phosphoritical Areoligera" Association

0.361	336	witmer, Nani, zone 10	0.331	335	witmer, Nani, zone 10
57.001	371	Deflandria phosphoritica	00.331	371	Deflandria phosphoritica
9.331	170	Wetzeliella hamdenensis	20.001	121	Areoligera sp.
0.331	121	Areoligera sp.	19.001	221	Cleistosphaeridium diversispinosum
2.331	180	Thalassionora pelagica	1.671	160	Thalassionora pelagica
2.301	135	Spondinium paratabulatum	2.671	158	Tectatodinium pelitum
1.001	117	Paralecaniella indentata	2.331	117	Paralecaniella indentata
1.001	132	Senegalinum? diwynense	2.331	173	Wetzeliella sp. A
0.671	221	Cleistosphaeridium diversispinosum	2.001	721	Homotryblum calcium
0.671	591	Forma D	1.671	1131	Ooeruciodinium multispinosum
0.671	721	Homotryblum calcium	1.671	1701	Wetzeliella hamdenensis
0.671	741	Hysterochokoloma eisenacki	1.671	1781	Cyclopsella vieta
0.671	171	Wetzeliella lunans	1.331	641	Glaucocystis ordinata
0.671	185	Palambages sp. C	1.331	185	Palambages sp. C
0.331	11	Achniella biformoides n. gen., n. comb.	1.001	1351	Spondinium paratabulatum
0.331	21	Adriatosphaeridium multispinosum	0.671	21	Adriatosphaeridium multispinosum
0.331	61	Apectodinium Homomorphum complex	0.671	931	Kaliosphaeridium brevispinosum
0.331	471	Eocladopyxis penicillatum	0.671	381	Linguodinium machaerophorum
0.331	501	Fibrocysta bipolare	0.671	1861	Phaeosporopsis spp.
0.331	521	Fibrocysta caalispinosa	0.331	11	Achniella biformoides n. gen., n. comb.
0.331	531	Fibrocysta radiata	2.331	381	Deflandria wardenensis
0.331	641	Glaucocystis ordinata	0.331	501	Fibrocysta bipolare
0.331	761	Hysterochokoloma nguade	0.331	531	Fibrocysta radiata
0.331	931	Kaliosphaeridium brevispinosum	0.331	591	Forma D
0.331	941	Kisselovia coleothrypta	0.331	691	Heteraulacocystis carmoana
0.331	981	Linguodinium machaerophorum	0.331	701	Homotryblum calcium
0.331	1001	Membranillamacia leptoderma	0.331	711	Homotryblum tasmanense
0.331	1021	Melliodinium guseppi major	0.331	821	Hysterochokoloma tubiferum brevispin.
0.331	1071	Nematosphaeropsis irabeculata	0.331	941	Kisselovia coleothrypta
0.331	1131	Ooeruciodinium multispinosum	0.331	1221	Phthalopendium acinatum
0.331	1151	Palaeocystodinium galzowense	0.331	1241	Polysphaeridium zonari
0.331	1281	Samtancia chlamydozona	0.331	1281	Samtancia chlamydozona
0.331	1301	Senegalinum? obscurum	0.331	1301	Senegalinum? obscurum
0.331	1461	Spiniferites pseudofurcatus	0.331	1321	Senegalinum? diwynense
0.331	1511	Spiniferites ramosus subsp. ramosus	0.331	1511	Spiniferites ramosus subsp. ramosus
0.331	1581	Tectatodinium pelitum	0.331	1651	Tubidermodinium sulcatum
0.331	1721	Wetzeliella samandica	0.331	1711	Wetzeliella lunans
0.331	1781	Cyclopsella vieta	0.331	1751	Walsbyidium tabulatum
0.331	1821	Cymatosphaera spp.	0.331	1811	Microstadium spp.



Correlation Matrix for PT12

Correlation Matrix for Variables: X1 ... X2

	335	336
335	1	
336	.76476	1

ANOVA Table for PT12

One Factor ANOVA-Repeated Measures for X1 ... X2

Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	4296.06572	16.91364	6.28092	.0001
Within subjects	255	686.6793	2.69286		
treatments	1	.44235	.44235	.16373	.6861
residual	254	686.23695	2.70172		
Total	509	4982.74502			

Reliability Estimates for- All treatments: .84079    Single Treatment: .72531



Statistical Information for Sample 337

X<sub>1</sub>: 337

Mean:	Std. Dev.:	Std. Error:	Variance:	Coef. Var.:	Count:
.50173	2.85452	.17876	8.1483	568.94104	255
Minimum:	Maximum:	Range:	Sum:	Sum of Sqr.:	# Missing:
0	28.33	28.33	127.94	2133.8582	0

PT13 "Glaphyrocysta ordinata/ Homotriblium palladium" Association

339	339	witmer, Nani, zone 11
31.67	54	Glaphyrocysta ordinata
31.66	72	Homotriblium palladium
3.33	22	Cleistosphaeridium diversispinosum
5.33	171	Wetzelieella unans
3.33	38	Lingulodinium machaerophorum
1.57	78	Cyclopsella vieta
1.57	22	Phthanopendrium echinatum
1.33	33	Corosphaeridium callosum
1.33	151	Spiniferites ramosus subsp. ramosus
1.00	00	Membranilamaca leioderma
1.00	17	Paracaneila inornata
0.67	28	Corosphaeridium inodes
0.67	47	Eocladopyxis peniculatum
0.67	35	Spinidium paratabulatum
0.67	40	Spiniferites cingulatus
0.67	49	Spiniferites ramosus subsp. membranaceus
0.33	1	Achnella biformoides n. gen., n. comb.
0.33	13	Ascotomocystus hydra
0.33	39	Dinopterygium cacoides
0.33	41	Diphyes colligerum
0.33	43	Diphyes colligerum (sensu cookson 1985)
0.33	70	Homotriblium calcicum
0.33	74	Hystriochokorboma eisenackii
0.33	75	Hystriochokorboma nguade
0.33	80	Hystriochosphaeridium spp.
0.33	83	?Hystriochosphaeropsis ovum
0.33	88	Impietosphaeridium kroemmelbairni
0.33	99	Impietosphaeridium rugosum
0.33	33	Kaliosphaeridium breviarbatum
0.33	06	Nematospaeropsis pusillosa
0.33	07	Nematospaeropsis irabeculata
0.33	10	Oerculodinium centrocarpum
0.33	13	Oerculodinium multispinosum
0.33	32	Seneganium? davyanense
0.33	38	Spiniferites crassioleis C
0.33	58	Tectatodinium pelitum
0.33	81	Micrystidium spp.
0.33	82	Cymatosphaera spp.
0.33	85	Palmabages sp. C
0.33	86	Perosermoopsis spp.

338	338	witmer, Nani, zone 11
27.33	72	Homotriblium palladium
19.67	54	Glaphyrocysta ordinata
9.33	1	Achnella biformoides n. gen., n. comb.
6.67	71	Wetzelieella unans
6.33	98	Lingulodinium machaerophorum
5.00	40	Dinopterygium namamense
5.67	47	Eocladopyxis peniculatum
4.00	17	Paracaneila inornata
2.67	70	Wetzelieella namponensis
1.57	78	Cyclopsella vieta
1.00	13	Ascotomocystus hydra
1.00	35	Spinidium paratabulatum
1.00	51	Spiniferites ramosus subsp. ramosus
0.67	2	Adnatospaeridium multispinosum
0.67	22	Cleistosphaeridium diversispinosum
0.67	33	Corosphaeridium callosum
0.67	38	Delianona waltonensis
0.67	100	Membranilamaca leioderma
0.67	107	Nematospaeropsis irabeculata
0.67	22	Phthanopendrium echinatum
0.67	32	Seneganium? davyanense
0.67	60	Thalassiphora pelagica
0.33	9	Apteodinium cribosum
0.33	42	Diphyes colligerum
0.33	59	Forma D
0.33	70	Homotriblium calcicum
0.33	82	Hystriochosphaeridium tuofferum brevispin
0.33	83	?Hystriochosphaeropsis ovum
0.33	88	Impietosphaeridium kroemmelbairni
0.33	89	Impietosphaeridium rugosum
0.33	13	Oerculodinium multispinosum
0.33	124	Polyosphaeridium zonary
0.33	127	Rottneusia oorusica
0.33	138	Spiniferites crassioleis C
0.33	149	Spiniferites ramosus subsp. membranaceus
0.33	158	Tectatodinium pelitum

Correlation Matrix for PT13

Correlation Matrix for Variables: X1 ... X2

	338	339
338	1	
339	.79852	1

ANOVA Table for PT13

One Factor ANOVA-Repeated Measures for X1 ... X2

Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	2875.11607	11.31935	8.47392	.0001
Within subjects	255	340.6259	1.33579		
treatments	1	.00738	.00738	.0055	.9409
residual	254	340.61852	1.34102		
Total	509	3215.74197			

Reliability Estimates for- All treatments: .88199      Single Treatment: .78889

PT14 "Wetzeliella (W.) samlandica Fusidinium tabulatum" Association

239	239	goodman, Nani woodstock zone 16	240	240	goodman, Nani woodstock zone 16	241	241	goodman, Nani woodstock zone 16			
41	60	72	Wetzeliella (W.) samlandica	53	80	172	Wetzeliella (W.) samlandica	72	40	172	Wetzeliella (W.) samlandica
39	60	230	Fusidinium tabulatum n. gen., n. sp.	16	80	38	Linguodinium machaerophorum	17	80	230	Fusidinium tabulatum n. gen., n. sp.
7	60	202	Scnrodinium (S.) australiense	5	00	209	Spiniferes supparus	5	30	209	Spiniferes supparus
4	20	209	Spiniferes supparus	4	80	21	Adnatosphaeridium multisporosum	2	60	38	Linguodinium machaerophorum
1	90	38	Linguodinium machaerophorum	4	20	202	Scnrodinium (S.) australiense	2	20	200	Operculodinium amiculum n. sp.
1	00	101	Microdinium ornatum	3	40	230	Fusidinium tabulatum n. gen., n. sp.	2	20	202	Scnrodinium (S.) australiense
0	60	03	Muratodinium limbnatum	2	80	103	Muratodinium limbnatum	2	20	12	Araoiigera sp.
0	30	110	Operculodinium centrocarpum	1	40	227	Cycloneoneium lemiscatum	1	80	21	Adnatosphaeridium multisporosum
0	60	213	Deflandra dilwynensis	1	00	100	Memoranillamacia leotoderma	0	60	100	Memoranillamacia leotoderma
0	60	21	Araoiigera sp.	2	80	215	Phthanoperidinium resistente	0	60	222	Adnatosphaeridium vittatum
0	60	233	Heteraulacysta lehmannensis	0	80	222	Adnatosphaeridium vittatum	0	40	76	Hystichokolpoma nguade
0	40	41	Diphyes colligerum	0	80	101	Microdinium ornatum	0	40	195	Lantemosphaeridium bipolare
0	40	76	Hystichokolpoma nguade	0	60	200	Operculodinium amiculum n. sp.	0	40	103	Muratodinium limbnatum
0	40	93	Kaliosphaeridium breviarbatum	0	40	11	Achiella biformoides n. gen., n. comb.	0	40	215	Phthanoperidinium resistente
0	40	195	Lantemosphaeridium bipolare	0	40	148	Spiniferes ramosus subsp. granosus	0	40	228	Dinopterygium cladoides
0	40	100	Memoranillamacia leotoderma	0	40	47	Eociadopyxis peniculatum	0	40	39	Impietosphaeridium rugosum
0	40	200	Operculodinium amiculum n. sp.	0	40	233	Heteraulacysta lehmannensis	0	40	235	Impietosphaeridium transodum
0	40	207	Spiniferes cf. S. ramosus subsp. ramosus	0	40	89	Impietosphaeridium rugosum	0	20	11	Achiella biformoides n. gen., n. comb.
0	40	115	Paiaocystodinium gozowense	0	40	235	Impietosphaeridium transodum	0	20	33	Cordosphaeridium caliosum
0	40	21	Adnatosphaeridium multisporosum	0	20	33	Cordosphaeridium caliosum	0	20	27	Cordosphaeridium gracilliss
0	40	222	Adnatosphaeridium vittatum	0	20	27	Cordosphaeridium gracilliss	0	20	28	Corosphaeridium nodos
0	40	226	Cycloneoneium incutum	0	20	41	Diphyes colligerum	0	20	41	Diphyes colligerum
0	40	235	Impietosphaeridium transodum	0	20	191	Gonyaulacysta guiseppi subsp. major	0	20	194	Hystichosphaeropsis borussica
0	20	11	Achiella biformoides n. gen., n. comb.	0	20	76	Hystichokolpoma nguade	0	20	199	Lantemosphaeridium radiatum
0	20	33	Cordosphaeridium caliosum	0	20	194	Hystichosphaeropsis borussica	0	20	201	Operculodinium potomacense n. sp.
0	20	27	Cordosphaeridium gracilliss	0	20	93	Kaliosphaeridium breviarbatum	0	20	146	Spiniferes pseudofurcatus
0	20	191	Gonyaulacysta guiseppi subsp. major	0	20	195	Lantemosphaeridium bipolare	0	20	50	Spiniferes ramosus subsp. multibrevis
0	20	194	Hystichosphaeropsis borussica	0	20	199	Lantemosphaeridium radiatum	0	20	151	Spiniferes ramosus subsp. ramosus
0	20	147	Spiniferes ramosus subsp. granosus	0	20	110	Operculodinium centrocarpum	0	20	208	Spiniferes scabratus
0	20	50	Thalassiosira pelagica	0	20	204	Spiniferes cf. S. crassioleis	0	20	313	Deflandra dilwynensis
0	20	211	Tubiosphaeridium pseudocurvatum n. gen.	0	20	208	Spiniferes membranaceus	0	20	38	Deflandra wardenensis
0	20	38	Deflandra wardenensis	0	20	145	Spiniferes monius	0	20	151	Paiaocystodinium gozowense
0	20	215	Phthanoperidinium resistente	0	20	150	Spiniferes ramosus subsp. multibrevis	0	20	219	Wetzeliella (W.) articulata
0	20	227	Cycloneoneium lemiscatum	0	20	151	Spiniferes ramosus subsp. ramosus	0	20	224	Cannosphaeropsis pusilliosa
0	20	228	Dinopterygium cladoides	0	20	158	Tectatodinium sellitum	0	20	226	Cycloneoneium incutum
0	20	39	Impietosphaeridium rugosum	0	20	160	Thalassiosira pelagica	0	20	227	Cycloneoneium lemiscatum
0	20	165	Tubidermodinium sulcatum	0	20	211	Tubiosphaeridium pseudocurvatum n. gen.	0	20	59	Heteraulacysta campanula
				0	20	213	Deflandra dilwynensis	0	20	233	Heteraulacysta lehmannensis
				0	20	38	Deflandra wardenensis	0	20	101	Microdinium ornatum
				0	20	219	Wetzeliella (W.) articulata	0	20	165	Tubidermodinium sulcatum
				0	20	12	Araoiigera sp.				
				0	20	228	Dinopterygium cladoides				
				0	20	69	Heteraulacysta campanula				
				0	20	236	Litosphaeridium inensibucinum				
				0	20	165	Tubidermodinium sulcatum				



Correlation Matrix for PT14

Correlation Matrix for Variables: X<sub>1</sub> ... X<sub>3</sub>

	240	241	242
240	1		
241	.94496	1	
242	.48557	.64424	1

ANOVA Table for PT14

One Factor ANOVA-Repeated Measures for X<sub>1</sub> ... X<sub>3</sub>

Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	9826.52528	38.68711	7.68774	.0001
Within subjects	510	2566.48	5.03231		
treatments	2	.28936	.14468	.02864	.9718
residual	508	2566.19064	5.05156		
Total	764	12393.00528			

Reliability Estimates for- All treatments: .86992    Single Treatment: .69033



Sample 243

243:	243	goodman, Nani woodstock zone 18
55.80:	101	Microdinium ornatum
28.40:	21	Achnatosphaeridium multispinosum
3.40:	103	Muratoridium limbatum
3.00:	172	Weißeella (W.) samandica
2.20:	98	Lingulodinium machaerophorum
1.20:	209	Spiniferites supparus
1.00:	215	Phthanoperidinium resistente
0.80:	41	Diphyes colligerum
0.80:	235	Impletosphaeridium transodum
0.60:	29	Cordosphaeridium inodes
0.60:	100	Membranilamaca leptoderma
0.60:	110	Operculodinium centrocarpum
0.60:	151	Spiniferites ramosus subsp. ramosus
0.60:	222	Achnatosphaeridium vitatum
0.60:	230	Fusulinium tabusatum n. gen., n. sp.
0.40:	147	Spiniferites ramosus subsp. granomemoratum
0.40:	38	Delandria wardenensis
0.40:	89	Heteraulacysta carnosula
0.40:	233	Heteraulacysta lehmannensis
0.40:	99	Impletosphaeridium rugosum
0.40:	65	Tubidermodinium succatum
0.20:	11	Achnella bifurcata n. gen., n. comb.
0.20:	33	Cordosphaeridium callosum
0.20:	25	Cordosphaeridium fibrospinatum
0.20:	27	Cordosphaeridium gracile
0.20:	189	Diphyesopsis buccinata n. gen., n. sp.
0.20:	90	Diphyesopsis capsata n. gen., n. sp.
0.20:	48	Exochosphaeridium bilidum
0.20:	191	Gonyaulacysta guiseppi subsp. major
0.20:	92	Hyalosphaera marylandensis n. gen., n. sp.
0.20:	193	Hysterochokeopoma tubosum
0.20:	76	Hysterochokeopoma rugatum
0.20:	79	Hysterochokeopoma tubosum
0.20:	194	Hysterochokeopoma bonasica
0.20:	93	Kaliosphaeridium ornitibaratum
0.20:	195	Laternosphaeridium bicolare
0.20:	199	Laternosphaeridium radiatum
0.20:	200	Operculodinium anticum n. sp.
0.20:	202	Scniodinium (S.) austrakense
0.20:	203	Spiniferites bulbosus
0.20:	198	Spiniferites crassipertis
0.20:	204	Spiniferites cf. S. crassipertis
0.20:	205	Spiniferites hyperandrus
0.20:	208	Spiniferites membranaceus
0.20:	146	Spiniferites pseudofurcatus
0.20:	148	Spiniferites ramosus subsp. granosus
0.20:	150	Spiniferites ramosus subsp. multibrevis
0.20:	207	Spiniferites cf. S. ramosus subsp. ramosus
0.20:	208	Spiniferites scabratus
0.20:	158	Tectadocinium bellum
0.20:	211	Tubulosphaeridium pseudocurvatum n. gen.
0.20:	37	Delandria onosphortica
0.20:	115	Palaocystodinium gotowense
0.20:	219	Weißeella (W.) articulata
0.20:	12	Areligera sp.
0.20:	225	Chlamydotheca ruca n. sp.
0.20:	22	Cleistosphaeridium diversispinosum
0.20:	226	Cyclosetonium incurtum
0.20:	227	Cyclosetonium lemmacatum
0.20:	228	Dinodorygium cladose
0.20:	231	Hemicyclodinium aculeatum n. sp.

Statistical information for Sample 243

X<sub>1</sub>: 243

Mean:	Std. Dev.:	Std. Error:	Variance:	Coef. Var.:	Count:
.42588	3.86698	.24216	14.9535	907.99152	255
Minimum:	Maximum:	Range:	Sum:	Sum of Sq.:	# Missing:
0	55.8	55.8	108.6	3844.44	0

PT15 "Deflandria macmurdoensis" Association

244	244 goodman, Nanj woodstock zone '6	245	245 goodman, Nanj woodstock zone '6	246	246 goodman, Nanj woodstock zone '6
22.90	2141Deflandria macmurdoensis	39.20	2141Deflandria macmurdoensis	51.80	2141Deflandria macmurdoensis
12.00	1011Microdinium ornatum	9.80	2211Wetziella (W.) coleothrypa	11.00	2211Wetziella (W.) coleothrypa
10.60	2211Wetziella (W.) coleothrypa	7.20	271Corcosphaendum gracilis	3.20	221Cleistosphaendum diversispinosum
3.40	121Arocligera sp.	7.00	221Cleistosphaendum diversispinosum	6.20	2151Phthanoendinium resistente
5.80	981Linguodinium machaerophorum	6.20	121Arocligera sp.	0.60	2311Hemicyclostodinium aculeatum n. sp.
5.40	371Deflandria phosponitica	3.20	2371Systematophora ancyrea	3.00	2121Deflandria asymmetrica
4.80	411Diphyes colligerum	2.50	981Linguodinium machaerophorum	2.90	411Diphyes colligerum
3.60	271Corcosphaendum gracilis	2.50	1101Oercuodinium centrocarpum	2.50	2011Coercuodinium ootomacense n. sp.
3.20	1721Wetziella (W.) samlandica	2.00	2011Coercuodinium ootomacense n. sp.	2.20	591Heterauacysta campanula
2.80	2011Coercuodinium ootomacense n. sp.	2.00	2151Phthanoendinium resistente	1.90	271Corcosphaendum gracilis
1.80	2101Tectatodinium psilatum	1.90	411Diphyes colligerum	1.60	1701Wetziella (W.) hamdenensis
1.30	1601Thalassionora pelagica	1.30	371Deflandria phosponitica	1.60	1011Microdinium ornatum
1.20	2301Fusidinium tabulatum n. gen., n. sp.	1.50	2131Deflandria dihyvensis	1.40	381Linguodinium machaerophorum
1.40	1031Muratodinium limbatum	1.40	1721Wetziella (W.) samlandica	1.20	1511Spiniferites ramosus subsp. ramosus
1.00	1511Spiniferites ramosus subsp. ramosus	1.20	1001Membranilamacia leptodema	1.20	2331Heterauacysta lehmannensis
1.00	2091Spiniferites supparus	1.00	511Spiniferites ramosus subsp. ramosus	0.80	121Arocligera sp.
1.00	2221Adnatosphaendum vitatum	0.80	331Kalliosphaendum brevibarbatum	0.60	1901Diphyesopsis capitata n. gen., n. sp.
1.00	221Cleistosphaendum diversispinosum	0.80	1581Spiniferites hypercanthus	0.60	1461Spiniferites pseudolurcatus
0.80	2151Phthanoendinium resistente	0.80	2121Deflandria asymmetrica	0.60	2071Spiniferites cf. S. ramosus subsp. ramosus
0.80	21Adnatosphaendum multispinosum	0.80	1711Wetziella (W.) lunans	0.60	2091Spiniferites supparus
0.80	2321Hemicyclostodinium zonaryi	0.80	2321Hemicyclostodinium zonaryi	0.60	381Deflandria wardenensis
0.80	391Impietosphaendum rugosum	0.60	1881Aurina ennaea n. gen., n. comb.	0.60	471Eocladopyxis peniculatum
0.60	2001Oercuodinium amiculum n. sp.	0.60	761Hystriochloa riguae	0.60	2321Hemicyclostodinium zonaryi
0.60	1381Spiniferites crassipellis	0.60	1951Lantemosphaendum bipolare	0.60	391Impietosphaendum rugosum
0.60	381Deflandria wardenensis	0.60	2101Tectatodinium psilatum	0.40	1881Aurina ennaea n. gen., n. comb.
0.40	251Corcosphaendum fibrosipinosum	0.60	2281Dinopterygium cladoides	0.40	1001Membranilamacia leptodema
0.40	761Hystriochloa riguae	0.40	691Heterauacysta campanula	0.40	1101Coercuodinium centrocarpum
0.40	331Kalliosphaendum brevibarbatum	0.60	2351Impietosphaendum transtodum	0.40	2131Deflandria dihyvensis
0.40	1101Coercuodinium centrocarpum	0.40	301Corcosphaendum multispinosum	0.40	371Deflandria phosponitica
0.40	1121Coercuodinium israelianum	0.40	1901Diphyesopsis capitata n. gen., n. sp.	0.40	1711Wetziella (W.) lunans
0.40	2121Deflandria asymmetrica	0.40	1381Spiniferites crassipellis	0.40	2281Dinopterygium cladoides
0.40	471Eocladopyxis peniculatum	0.40	2051Spiniferites hypercanthus	0.20	11Achiella biformoides n. gen., n. comb.
0.40	2311Hemicyclostodinium aculeatum n. sp.	0.40	1701Wetziella (W.) hamdenensis	0.20	331?Corcosphaendum callosum
0.40	2351Impietosphaendum transtodum	0.20	891Impietosphaendum rugosum	0.20	251Corcosphaendum fibrosipinosum
0.20	11Achiella biformoides n. gen., n. comb.	0.20	11Achiella biformoides n. gen., n. comb.	0.20	1891Diphyesopsis buccinata n. gen., n. sp.
0.20	1881Aurina ennaea n. gen., n. comb.	0.20	331?Corcosphaendum callosum	0.20	1921Hydrasphaera marylandense n. gen., n. sp.
0.20	331?Corcosphaendum callosum	0.20	281Corcosphaendum inodes	0.20	1931Hystriochloa tubosa
0.20	281Corcosphaendum inodes	0.20	481Exochosphaendum bifidum	0.20	761Hystriochloa riguae
0.20	1891Diphyesopsis buccinata n. gen., n. sp.	0.20	1921Hydrasphaera marylandense n. gen., n. sp.	0.20	331Kalliosphaendum brevibarbatum
0.20	1901Diphyesopsis capitata n. gen., n. sp.	0.20	1951Lantemosphaendum bipolare	0.20	1991Lantemosphaendum radiatum
0.20	481Exochosphaendum bifidum	0.20	1931Hystriochloa tubosa	0.20	1001Membranilamacia leptodema
0.20	1921Hydrasphaera marylandense n. gen., n. sp.	0.20	791Hystriochloa tuiferum	0.20	2021Scrinodinium (S.) austraiense
0.20	1931Hystriochloa tubosa	0.20	971Lantemosphaendum lappaceum	0.20	141Spiniferites comutus
0.20	1941Hystriochloa borussica	0.20	1031Muratodinium limbatum	0.20	2061Spiniferites membranaceus
0.20	1951Lantemosphaendum bipolare	0.20	041Nematosphaeropsis calcomiana	0.20	1471Spiniferites ramosus subsp. granomembran.
0.20	1991Lantemosphaendum radiatum	0.20	2001Coercuodinium amiculum n. sp.	0.20	1581Tectatodinium psilatum
0.20	1001Membranilamacia leptodema	0.20	141Spiniferites comutus	0.20	1481Spiniferites ramosus subsp. granosus
0.20	2021Scrinodinium (S.) austraiense	0.20	2041Spiniferites cf. S. crassipellis	0.20	1601Thalassionora pelagica
0.20	2041Spiniferites cf. S. crassipellis	0.20	2061Spiniferites membranaceus	0.20	2111Tubulosphaendum pseudocurvatum n. gen.
0.20	2051Spiniferites hypercanthus	0.20	1451Spiniferites monius	0.20	21Adnatosphaendum multispinosum
0.20	2061Spiniferites membranaceus	0.20	1471Spiniferites pseudolurcatus	0.20	381Deflandria wardenensis
0.20	1461Spiniferites pseudolurcatus	0.20	1471Spiniferites ramosus subsp. granomembran.	0.20	221Adnatosphaendum vitatum
0.20	1471Spiniferites ramosus subsp. granomembran.	0.20	1581Tectatodinium psilatum	0.20	21Adnatosphaendum multispinosum
0.20	1481Spiniferites ramosus subsp. granosus	0.20	1601Thalassionora pelagica	0.20	2221Adnatosphaendum vitatum
0.20	1581Tectatodinium psilatum	0.20	2111Tubulosphaendum pseudocurvatum n. gen.	0.20	221Cannosphaeropsis pusillosa
0.20	1601Thalassionora pelagica	0.20	2171Spinidinium rotundum	0.20	2261Cyclonephium incutum
0.20	2111Tubulosphaendum pseudocurvatum n. gen.	0.20	1721Wetziella (W.) samlandica	0.20	2271Cyclonephium lemiscatum
0.20	2171Spinidinium rotundum	0.20	21Adnatosphaendum multispinosum	0.20	2301Fusidinium tabulatum n. gen., n. sp.
0.20	1721Wetziella (W.) samlandica	0.20	221Adnatosphaendum vitatum	0.20	2311Hemicyclostodinium aculeatum n. sp.
0.20	21Adnatosphaendum multispinosum	0.20	2221Adnatosphaendum vitatum	0.20	2331Heterauacysta lehmannensis
0.20	221Adnatosphaendum vitatum	0.20	221Cannosphaeropsis pusillosa	0.20	1011Microdinium ornatum
0.20	2221Adnatosphaendum vitatum	0.20	2261Cyclonephium incutum	0.20	1651Tubidermodinium sulcatum
0.20	221Cannosphaeropsis pusillosa	0.20	2271Cyclonephium lemiscatum		
0.20	2261Cyclonephium incutum	0.20	2301Fusidinium tabulatum n. gen., n. sp.		
0.20	2271Cyclonephium lemiscatum	0.20	2311Hemicyclostodinium aculeatum n. sp.		
0.20	2301Fusidinium tabulatum n. gen., n. sp.	0.20	2331Heterauacysta lehmannensis		
0.20	2311Hemicyclostodinium aculeatum n. sp.	0.20	1011Microdinium ornatum		
0.20	2331Heterauacysta lehmannensis	0.20	1651Tubidermodinium sulcatum		
0.20	1011Microdinium ornatum				
0.20	1651Tubidermodinium sulcatum				

Correlation Matrix for PT15

Correlation Matrix for Variables: X1 ... X3

	244	245	246
244	1		
245	.84606	1	
246	.80409	.96751	1

ANOVA Table for PT15

One Factor ANOVA-Repeated Measures for X1 ... X3

Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	4981.02975	19.61035	14.80823	.0001
Within subjects	510	675.38667	1.32429		
treatments	2	.23414	.11707	.08809	.9157
residual	508	675.15252	1.32904		
Total	764	5656.41642			

Reliability Estimates for- All treatments: .93247 Single Treatment: .82152



Statistical information for Sample 247

X1: 247

Mean:	Std. Dev.:	Std. Error:	Variance:	Coef. Var.:	Count:
.43216	2.64987	.16594	7.0218	613.17255	255
Minimum:	Maximum:	Range:	Sum:	Sum of Sq.:	# Missing:
0	35.6	35.6	110.2	1831.16	0



267	SPECIES	268	SPECIES	270	SPECIES
17.50	171 Weizaelella (W.) lunans	38.00	160 Thaisaenora pelagica	11.50	171 Weizaelella (W.) lunans
33.20	150 Thaisaenora pelagica	37.50	171 Weizaelella (W.) lunans	20.00	216 Spindinium bilineatum n. sp.
7.20	170 Weizaelella (W.) hamboenensis	9.40	121 Arealigera sp.	3.40	170 Weizaelella (W.) hamboenensis
5.50	121 Arealigera sp.	7.40	21 Adnatospaendium multispinosum	5.20	150 Thaisaenora pelagica
2.50	21 Adnatospaendium multispinosum	5.50	170 Weizaelella (W.) hamboenensis	0.40	151 Spinfertes ramosus subsp. ramosus
2.40	71 Homotrybium tasmanense	2.90	221 Cleistosphaendium diversispinosum	0.40	381 Delfandria wardenensis
1.20	1891 Diphyesopsis buccinata n. gen., n. sp.	0.80	151 Spinfertes ramosus subsp. ramosus	0.20	1381 Spinfertes crassipeilis
1.20	151 Spinfertes ramosus subsp. ramosus	0.80	216 Spindinium bilineatum n. sp.	0.20	721 Homotrybium pallidum
1.20	2091 Spinfertes suoparus	0.50	11 Achiella biformodes n. gen., n. comb.		
1.20	221 Cleistosphaendium diversispinosum	0.50	1871 Aptodinium incompositum n. sp.		
0.50	41 Diphyes colligerum	0.60	2331 Heteraulacysta tehamaensis		
0.50	381 Delfandria wardenensis	0.40	1891 Diphyesopsis buccinata n. gen., n. sp.		
0.40	1 Achiella biformodes n. gen., n. comb.	0.40	981 Linguodinium macnaerophorum		
0.40	191 Gonyaulacysta guiseppi subsp. major	0.20	31 Aptodinium cribosum		
0.40	201 Operculodinium potomacense n. sp.	0.20	271 Cordosphaendium gracilis		
0.40	1381 Spinfertes crassipeilis	0.20	41 Diphyes colligerum		
0.40	216 Spindinium bilineatum n. sp.	0.20	1901 Diphyesopsis capitata n. gen., n. sp.		
0.20	271 Cordosphaendium gracilis	0.20	1911 Gonyaulacysta guiseppi subsp. major		
0.20	1971 Lanternosphaendium lappaceum	0.20	761 Hystriochloa rigida		
0.20	381 Linguodinium macnaerophorum	0.20	1971 Lanternosphaendium lappaceum		
0.20	2001 Operculodinium amiculum n. sp.	0.20	1101 Operculodinium carmocarpum		
0.20	2221 Adnatospaendium vittatum	0.20	2011 Operculodinium potomacense n. sp.		
0.20	471 Eociadopyxis penicillatum	0.20	1461 Spinfertes pseudofurcatus		
0.20	591 Heteraulacysta campanula	0.20	381 Delfandria wardenensis		
0.20	2341 Homotrybium altum n. sp.	0.20	1151 Palaeocystodinium gotzowense		
0.20	891 Impletosphaendium rugosum	0.20	2221 Adnatospaendium vittatum		
0.20	2351 Impletosphaendium transfocum	0.20	711 Homotrybium tasmanense		



Correlation Matrix for PT16

Correlation Matrix for Variables: X<sub>1</sub> ... X<sub>5</sub>

	250	251	267	268	270
250	1				
251	.7864	1			
267	.62844	.80887	1		
268	.53083	.68455	.97611	1	
270	.74225	.95447	.82454	.71318	1

ANOVA Table for PT16

One Factor ANOVA-Repeated Measures for X<sub>1</sub> ... X<sub>5</sub>

Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	20121.07464	79.21683	14.9525	.0001
Within subjects	1020	5403.856	5.2979		
treatments	4	.20342	.05085	.00956	.9998
residual	1016	5403.65258	5.31856		
Total	1274	25524.93064			

Reliability Estimates for- All treatments: .93312    Single Treatment: .73618

PT17 "Spinidium bilineatum" Association

254	SPECIES	265	SPECIES	266	SPECIES
51.40	216 Spinidium bilineatum n. sp.	71.20	216 Spinidium bilineatum n. sp.	35.20	171 Wetzelia (W.) lunans
1.30	22 Cleistosphaeridium diversispinosum	7.30	191 Gonyaulacysta guseepei subsp. major	25.30	216 Spinidium bilineatum n. sp.
5.60	220 Wetzelia (W.) coalita n. sp.	5.00	171 Wetzelia (W.) lunans	5.20	170 Wetzelia (W.) hamdenensis
4.30	209 Spinifertes supparus	4.20	209 Spinifertes supparus	5.40	12 Araoligera sp.
1.40	171 Wetzelia (W.) lunans	2.00	22 Cleistosphaeridium diversispinosum	5.20	3 Aptecodinium cnbosum
3.40	11 Homotrybium tasmaniense	2.00	71 Homotrybium tasmaniense	5.20	160 Thalassonora pelagica
2.60	191 Gonyaulacysta guseepei subsp. major	1.80	170 Wetzelia (W.) hamdenensis	4.20	22 Cleistosphaeridium diversispinosum
2.20	170 Wetzelia (W.) hamdenensis	1.40	41 Diphyes colligerum	3.40	71 Homotrybium tasmaniense
1.80	160 Thalassonora pelagica	1.20	11 Achiella biformoides n. gen., n. comb.	2.60	209 Spinifertes supparus
1.80	12 Araoligera sp.	1.20	160 Thalassonora pelagica	2.00	151 Spinifertes ramosus subsp. ramosus
1.20	201 Operculodinium potomacense n. sp.	1.00	38 Linguodinium macnaeeronorum	1.40	215 Phthanopendinium resistente
1.00	11 Achiella biformoides n. gen., n. comb.	1.00	201 Operculodinium potomacense n. sp.	1.40	2 Agnatosphaeridium multispinosum
0.80	47 Eocladopyxis peniculatum	0.80	31 Aptecodinium cnbosum	1.20	221 Wetzelia (W.) coleothrypta
0.60	33 Cordosphaeridium calosum	0.80	235 Impietosphaeridium transtodum	1.00	11 Achiella biformoides n. gen., n. comb.
0.80	41 Diphyes colligerum	0.60	33 Cordosphaeridium calosum	1.00	158 Tectatodinium pelitum
0.60	204 Spinifertes cf. S. crassipalis	0.60	28 Cordosphaeridium inodes	1.00	72 Homotrybium pallidum
0.60	215 Phthanopendinium resistente	0.60	151 Spinifertes ramosus subsp. ramosus	3.50	41 Diphyes colligerum
0.60	89 Impietosphaeridium rugosum	0.60	220 Wetzelia (W.) coalita n. sp.	3.60	191 Gonyaulacysta guseepei subsp. major
0.60	165 Tubidermodinium sucatum	0.60	47 Eocladopyxis peniculatum	0.60	38 Linguodinium macnaeeronorum
0.40	28 Cordosphaeridium inodes	0.40	189 Diphyesopsis buccinata n. gen., n. sp.	0.60	47 Eocladopyxis peniculatum
0.40	38 Linguodinium macnaeeronorum	0.40	204 Spinifertes cf. S. crassipalis	0.60	165 Tubidermodinium sucatum
0.40	100 Membranillamacia leptoderma	0.40	69 Heteraulacysta campanula	0.40	115 Palaeocystodinium gozowense
0.40	224 Carnosphaeropsis ovisuosa	0.40	72 Homotrybium pallidum	0.40	39 Impietosphaeridium rugosum
0.40	229 Eisenackia scrobiculata	0.20	27 Cordosphaeridium gracilis	0.40	236 Litosphaeridium inversibuccinum
0.40	233 Heteraulacysta tenmamensis	0.20	190 Diphyesopsis capitata n. gen., n. sp.	0.20	27 Cordosphaeridium gracilis
0.40	72 Homotrybium pallidum	0.20	76 Hysterochloa nguade	0.20	28 Cordosphaeridium inodes
0.20	31 Aptecodinium cnbosum	0.20	79 Hysterosphaeridium tuoferum	0.20	189 Diphyesopsis buccinata n. gen., n. sp.
0.20	25 Cordosphaeridium fibrosipinosum	0.20	93 Kalliosphaeridium breviarbatum	0.20	195 Lantemosphaeridium bipolare
0.20	27 Cordosphaeridium gracilis	0.20	197 Lantemosphaeridium laopaceum	0.20	197 Lantemosphaeridium laopaceum
0.20	89 Diphyesopsis buccinata n. gen., n. sp.	0.20	100 Membranillamacia leptoderma	0.20	198 Lantemosphaeridium dilatatum n. sp.
0.20	190 Diphyesopsis capitata n. gen., n. sp.	0.20	103 Muratodinium limonatum	0.20	100 Membranillamacia leptoderma
0.20	48 Exochosphaeridium bifidum	0.20	200 Operculodinium amiculum n. sp.	0.20	103 Muratodinium limonatum
0.20	192 Hyndosphaera marylandense n. gen., n. sp.	0.20	110 Operculodinium centrocarpum	0.20	200 Operculodinium amiculum n. sp.
0.20	76 Hysterochloa nguade	0.20	158 Tectatodinium pelitum	0.20	110 Operculodinium centrocarpum
0.20	195 Lantemosphaeridium bipolare	0.20	210 Tectatodinium psilatum	0.20	201 Operculodinium potomacense n. sp.
0.20	197 Lantemosphaeridium laopaceum	0.20	38 Dellandria wardenensis	0.20	181 Spinifertes crassipalis
0.20	198 Lantemosphaeridium dilatatum n. sp.	0.20	115 Palaeocystodinium gozowense	0.20	213 Dellandria dilwynensis
0.20	200 Operculodinium amiculum n. sp.	0.20	215 Phthanopendinium resistente	0.20	38 Dellandria wardenensis
0.20	110 Operculodinium centrocarpum	0.20	221 Wetzelia (W.) coleothrypta	0.20	228 Dinoterygium cladoides
0.20	202 Scnrodinium (S.) australiense	0.20	12 Araoligera sp.	0.20	229 Eisenackia scrobiculata
0.20	205 Spinifertes hypercanthus	0.20	229 Eisenackia scrobiculata	0.20	233 Heteraulacysta tenmamensis
0.20	151 Spinifertes ramosus subsp. ramosus	0.20	233 Heteraulacysta tenmamensis	0.20	234 Homotrybium alisum n. sp.
0.20	207 Spinifertes cf. S. ramosus subsp. ramosus	0.20	89 Impietosphaeridium rugosum	0.20	235 Impietosphaeridium transtodum
0.20	158 Tectatodinium pelitum	0.20	236 Litosphaeridium inversibuccinum		
0.20	210 Tectatodinium psilatum	0.20	165 Tubidermodinium sucatum		
0.20	213 Dellandria dilwynensis				
0.20	38 Dellandria wardenensis				
0.20	115 Palaeocystodinium gozowense				
0.20	69 Heteraulacysta campanula				
0.20	34 Homotrybium alisum n. sp.				
0.20	235 Impietosphaeridium transtodum				
0.20	236 Litosphaeridium inversibuccinum				



Correlation Matrix for PT17

Correlation Matrix for Variables: X<sub>1</sub> ... X<sub>5</sub>

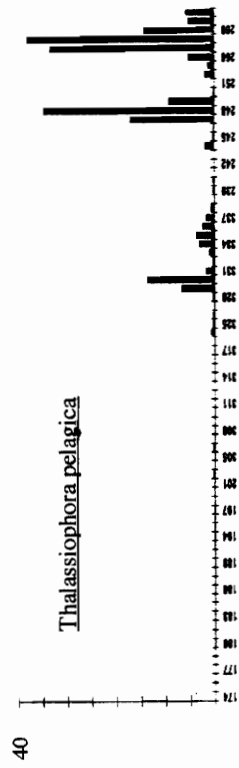
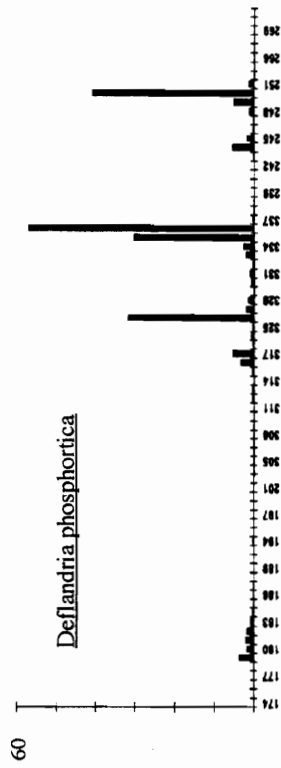
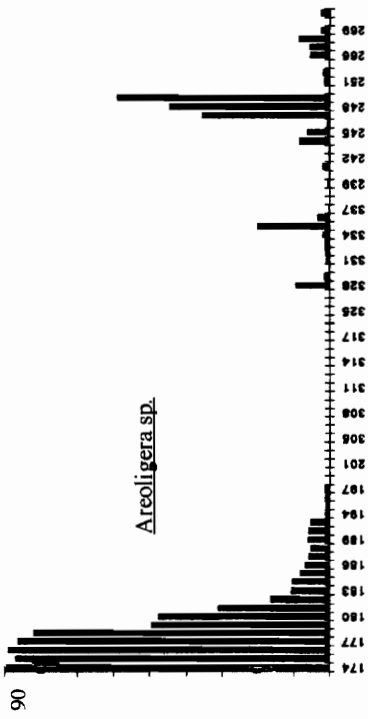
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264	1				
265	.97045	1			
266	.62956	.6265	1		
269	.58464	.59889	.96461	1	
271	.81201	.83367	.90005	.90565	1

ANOVA Table for PT17

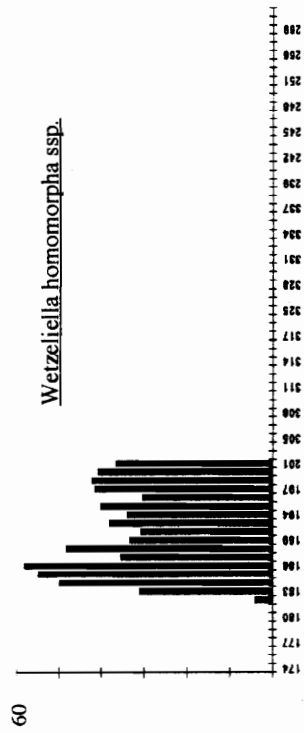
One Factor ANOVA-Repeated Measures for X<sub>1</sub> ... X<sub>5</sub>

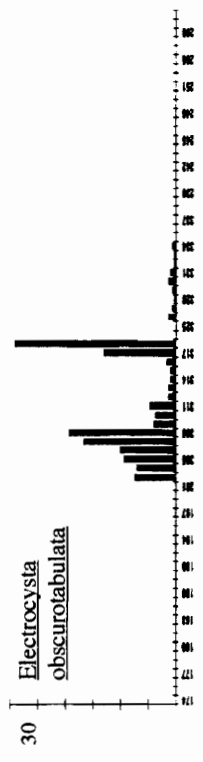
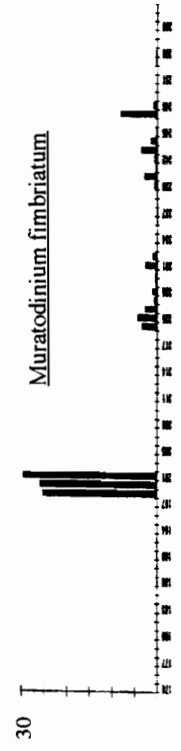
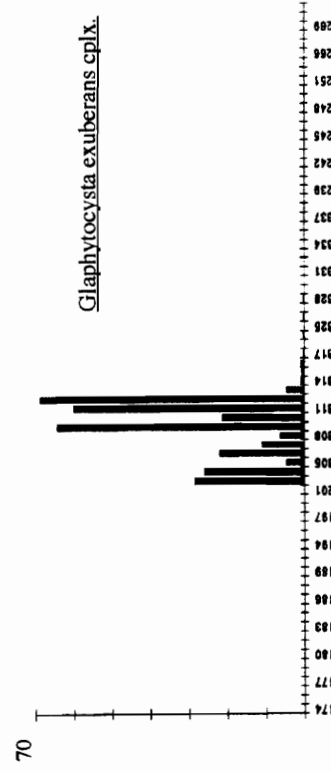
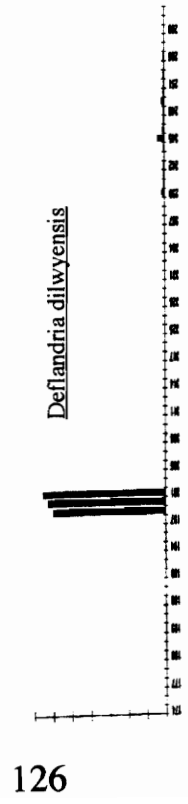
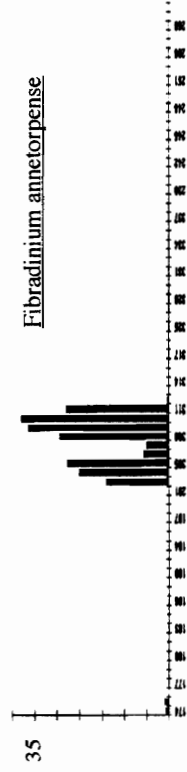
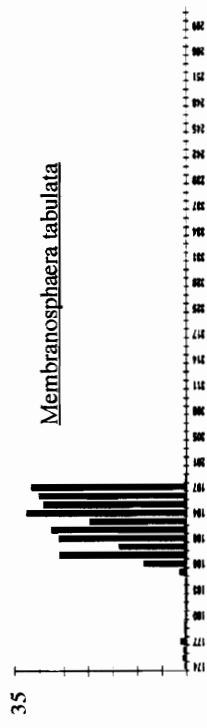
Source:	df:	Sum of Squares:	Mean Square:	F-test:	P value:
Between subjects	254	12255.06867	48.2483	16.25591	.0001
Within subjects	1020	3027.408	2.96805		
treatments	4	.15354	.03838	.01288	.9997
residual	1016	3027.25446	2.97958		
Total	1274	15282.47667			

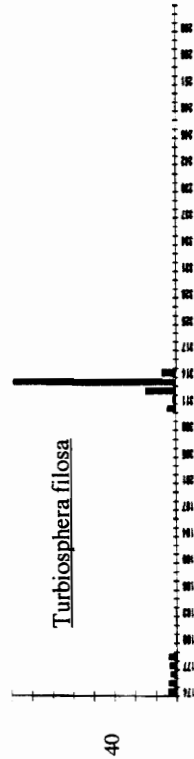
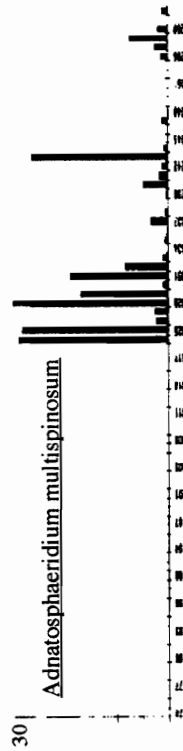
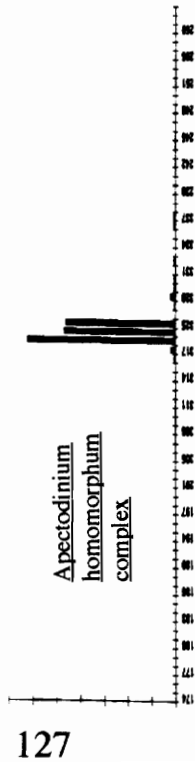
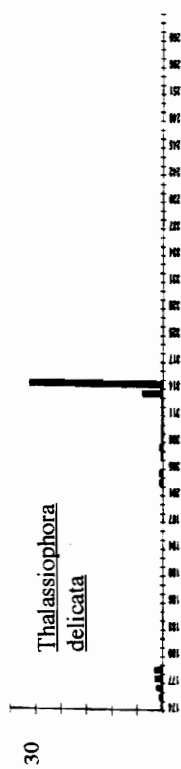
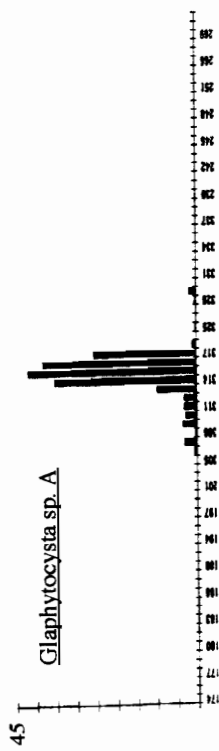
Reliability Estimates for- All treatments: .93848    Single Treatment: .75316



Graphical representation of very common (> 25%) dinoflagellate species.

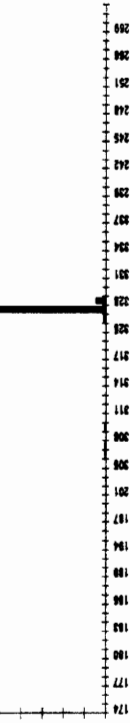






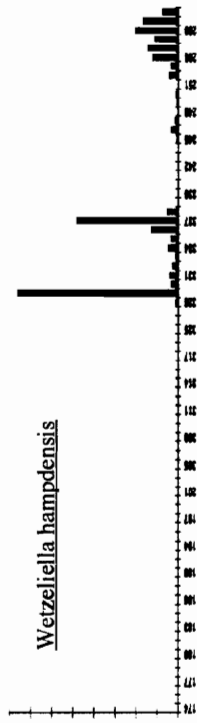
50

Systematophora placacantha



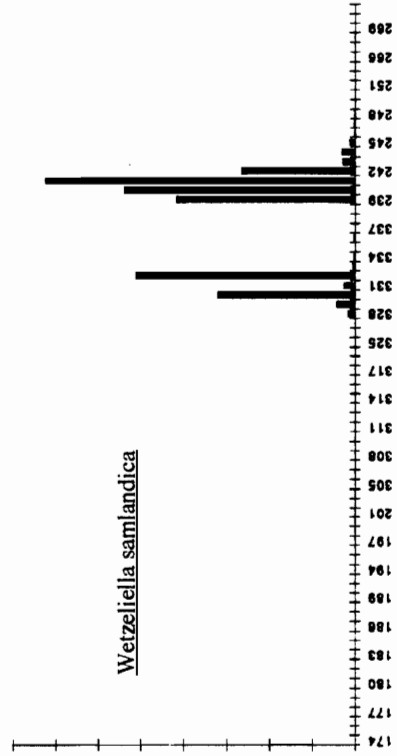
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Wetzeliella hampdensis



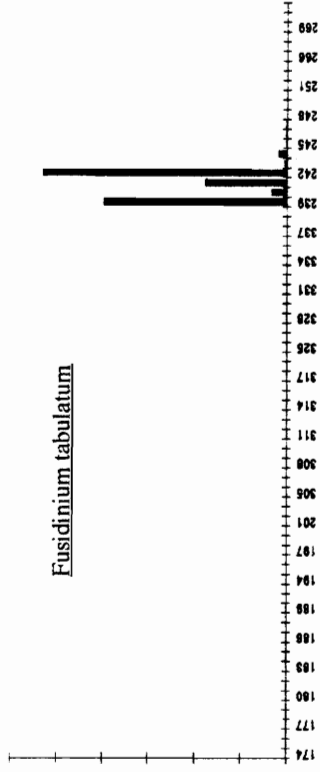
80

Wetzeliella samlandica



60

Fusidinium tabulatum



45

Spinidinium macmurdoense





35

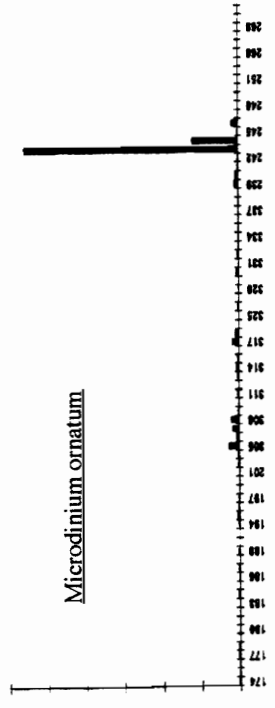
Glaphyrocysta ordinata



80

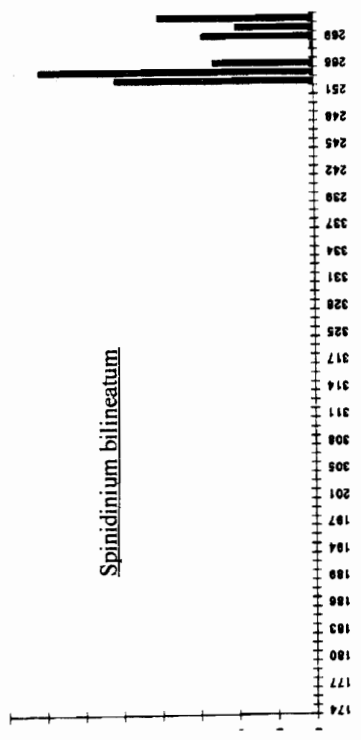
60

Microdinium ornatum



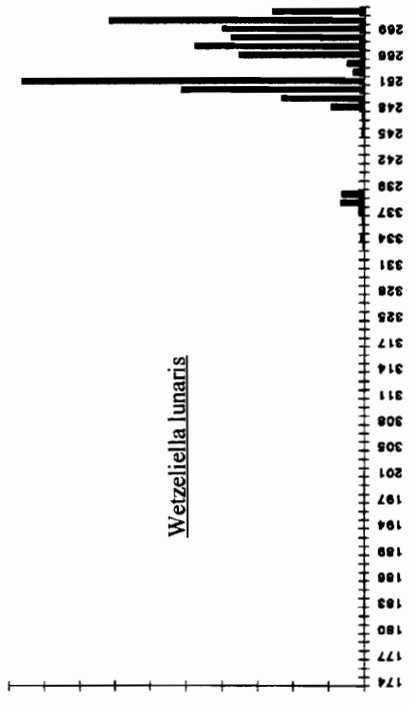
129

Spinidinium bilineatum



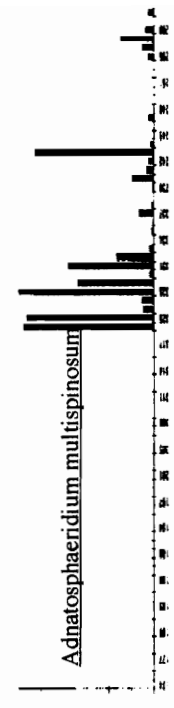
100

Wetzeliella lunaris



30

Adnatosphaeridium multispinosum



**APPENDIX B:**

**MODERN ANALOG FORTRAN MODEL X, Z  
MODEL OUTPUT DATA**

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C      DINOFLAGELLATE MODEL X
C      A DINOFLAGELLATE ECOSYSTEM DYNAMICS MODEL
C      5 DEGREES WARMER THAN BASIC YELLOW SEA MODEL
C      BY J. C. CAWLEY, 1996
C      X23  PHYTOPLANKTON POTENTIAL (NUTRIENT)
C      X1-X10  SPECIES 1-10
C      X11-22  CYSTS TO SEDIMENT
C
C      IMPLICIT REAL (L,M)
C      DIMENSION X(23),XD(23)
C      DIMENSION AC(12)
C      COMMON TDY
C      REAL M41,M42,M43
C      OPEN (6,FILE='CON')
C      OPEN (7,FILE='C:OUT.DAT')
C SET INITIAL CONDITIONS
DATA X /1.,1.,1.,1.,1.,1.,1.,1.,1.,1.,1.,1.,
+1.,1.,1.,1.,1.,1.,1.,1.,1.,1.,500./
DATA AC /1.,1.,1.,1.,1.,1.,1.,1.,1.,1.,1.,1./
D=05.
TEMP=19.
SAL=2.
C TA=TOTAL CYSTS IN SEDIMENT
TA=1.
N=4
T=0.
TMAX=1.0
C DT IS 0.001 YEAR
DT=.001
C IOCTR IS THE NUMBER OF ITERATIONS PER OUTPUT
IOCTR=83.33333333
C IOUT IS THE COUNTER FOR THE OUTPUT INTERVAL
IOUT=IOCTR+1
C WRITE COLUMN HEADINGS
WRITE (6,1)D
WRITE (7,1)D
1  FORMAT(' (DEPTH=' ,F5.1, ' METERS)')
C
WRITE (6,2)
WRITE (7,2)
2  FORMAT(' TIME,    SP1    SP2    SP3    SP4    SP5    SP6    SP7
+SP8    SP9    SP10')
C
C BEGINNING OF CALCULATION LOOP
3  CONTINUE
IF(IOUT.LT.IOCTR)GO TO 5
WRITE (6,4)T, (X(I),I=1,10)
WRITE (7,4)T, (X(I),I=1,10)
4  FORMAT(' ',F5.2,10F7.0)
IOUT=0
5  IOUT=IOUT+1
C CHECK FOR END OF SIMULATION
IF(T.GE.TMAX)GO TO 99
CALL RKS(X,XD,T,DT,N,D,TA,AC,TEMP,SAL)
T=T+DT
GO TO 3
99  STOP
END
CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC
SUBROUTINE DFQS(X,XD,T,N,D,TA,AC,TEMP,SAL)

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      IMPLICIT REAL (L,M)
      DIMENSION XD(23),X(23)
      DIMENSION MONTH(14),S(14)
      DIMENSION AC(12)
C     COMMON TDY
C
C
C THIS SUBROUTINE CONTAINS THE DIFFERENTIAL EQUATIONS
C
C STATEMENTS BELOW ARE USED TO FLUCTUATE TIME FOR SEASONS
      TWY=T-AINT(T)
C   TDY IS TIME WITHIN YEAR IN DAYS
      TDY=TWY*365.
C   TR IS TWY IN RADIANS
      TR=(TWY)*6.28318
C   MONTH IS THE MIDDLE DAY OF EACH MONTH
      DATA MONTH/0.,15.,46.,74.,105.,136.,166.,196.,
+227.,258.,288.,319.,349.,365./
C
C STATEMENTS BELOW VARY TEMP AS A SEASONAL SINE FUNCTION
C   AMT IS THE ANNUAL MEAN WATER TEMPERATURE
      IF(D.LT.25.)AMT=15.
      IF((D.GE.25.) .AND. (D.LT.75.))AMT=20
      IF(D.GE.75.)AMT=25.
C   TAMP IS THE AMPLITUDE OF THE TEMPERTATURE CURVE
      IF(D.LE.25.)TAMP=10.0
      IF((D.GT.25.) .AND. (D.LE.50.))TAMP=5.0
      IF((D.GT.50.) .AND. (D.LE.75.))TAMP=4.0
      IF(D.GT.75.)TAMP=4.0
C   TPS IS THE PHASE SHIFT OF TEMP CURVE IN RADIANS
      TPS=1.75585
      TEMP=AMT+SIN(TR-TPS)*TAMP
      FLAG=1
      IF (TEMP.LT.18.) FLAG=0.
C
C STATEMENTS BELOW VARY SALINITY THROUGH TIME
      DATA S/1.2,1.1,1.1,0.8,0.6,0.8,0.8,1.1,
+1.4,1.4,1.0,0.8,1.1,1.2/
      DO 101 I=1,14
101      IF(MONTH(I).GT.TDY)GOTO 102
102      HFA=(TDY-MONTH(I-1))/(MONTH(I)-MONTH(I-1))
      HMC=S(I)-S(I-1)
      SAL=(HFA*HMC)+S(I-1)
      IF(D.LT.15.)SAL=(SAL*1.4)
      IF((D.GE.15.) .AND. (D.LE.25.))SAL=(SAL*2.0)
      IF((D.GT.25.) .AND. (D.LE.50.))SAL=(SAL*2.5)
      IF((D.GT.50.) .AND. (D.LE.75.))SAL=(SAL*3.0)
      IF(D.GT.75.)SAL=3.4
C   INPUTS, Z IS DEPENDANT ON DEPTH IN METERS
      Z1=150.
      IF(D.GT.100.)Z1=100.
      IF((D.LE.100.) .AND. (D.GT.50.))Z1=150.
      IF((D.LE.50.) .AND. (D.GT.25.))Z1=300.
      IF(D.LE.25.)Z1=750.
C
C THIS SECTION IS FOR SPECIES 1, A NEARSHORE 'BLOOMER'
C   THIS IS SPECIES 1 GROWTHRATE:
      VL23=1.0*X(1)
      IF((TAMP.LT.4.) .OR. (SAL.GE.2.5))X(1)=1.
      IF((TEMP.GE.20.) .AND. (SAL.GT.1.))X(1)=(1.05*X(1))

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      IF((TEMP.LT.15.) .AND. (X(1).GE.10.)) X(1)=(0.95*X(1))
C   THESE ARE CONDITIONS OF SPECIES 1 ENCYSTMENT:
      IF(X(23).LT.50.)X(11)={(0.1*X(1))+X(11)}
      IF((TEMP.LT.23.02) .AND. (TEMP.GT.23.))X(11)={(0.1*X(1))+X(11)}
C   THIS IS WHAT HAPPENS IF SPECIES 1 RUNS OUT OF NUTRIENTS:
      IF(X(23).LT.50.)X(1)=1.
C
C   THIS SECTION IS FOR SPECIES 10, AN OCEANIC FORM
C   THIS IS SPECIES 10 GROWTHRATE
      V1023=1.0*X(10)
      IF((TEMP.LT.18.) .OR. (SAL.GT.2.0))X(10)=(1.02*X(10))
      IF((TEMP.LT.18.) .OR. (SAL.LT.2.0))X(10)=1.
C   THESE ARE CONDITIONS OF SPECIES 10 ENCYSTMENT
      IF((TEMP.LT.18.02) .AND. (FLAG.EQ.1.))X(20)=(0.50*X(10))+
+X(20)
C   THIS IS WHAT HAPPENS IF SPECIES 10 RUNS OUT OF NUTRIENTS
      IF(X(23).LT.50.)X(10)=1.
C
C   THIS SECTION IS FOR SPECIES 2, A COOLER NEARSHORE BLOOMER
C   THIS IS SPECIES 2 GROWTHRATE
      V223=1.0*X(2)
      IF(((TEMP.GT.15.) .AND. (TEMP.LT.20.)) .AND. (SAL.GT.1.))X(2)=
+(1.05*X(2))
      IF(((TEMP.LT.15.) .OR. (TEMP.GT.20.)) .OR. (SAL.LE.1.)))
+X(2)=1.
      IF(TAMP.LT.2.0)X(2)=1.
C   THESE ARE CONDITIONS OF SPECIES 2 ENCYSTMENT
      IF(X(23).LT.50)X(12)=(0.1*X(2))+X(12)
      IF((TEMP.LT.15.2) .AND. (TEMP.GE.15.0))X(12)=(0.05*X(2))+X(12)
      IF((TEMP.GT.17.08) .AND. (TEMP.LT.18.))X(12)=(0.05*X(2))+X(12)
C   THIS IS WHAT HAPPENS IF SPECIES 2 RUNS OUT OF NUTRIENTS
      IF (X(23).LT.50.)X(2)=1.
C
C   THIS SECTION IS FOR SPECIES 9, AN OCEANIC FORM
C   THIS IS SPECIES 9 GROWTHRATE
      V923=1.0*X(9)
      IF((TEMP.GE.18.) .AND. (SAL.GT.3.2))X(9)=1.03*X(9)
      IF((TEMP.LT.18.) .OR. (SAL.LT.3.2))X(9)=1.
C   THESE ARE THE CONDITIONS OF SPECIES 9 ENCYSTMENT
      DO 111 I=1,14
111   IF(MONTH(I).GE.TDY)X(19)=(0.05*X(9))+X(19)
C   THIS IS WHAT HAPPENS IF SPECIES 9 RUNS OUT OF NUTRIENTS
      IF(X(23).LT.50.)X(9)=1.
C
C   THIS SECTION IS FOR SPECIES 3, AN OCEANIC FORM, ASSOC SF 4
C   THIS IS SPECIES 3 GROWTHRATE
      V323=1.0*X(3)
      IF((TEMP.GT.20.) .AND. (SAL.GT.3.0))X(3)=(X(3)+20.)
      IF(X(4).LE.100.)X(3)=(1.01*X(4))
      IF(X(4).GT.100.)X(3)=(X(3)*0.99)
      IF((TEMP.LE.18.) .OR. (SAL.LE.3.0))X(3)=10.
      IF(TAMP.LT.2.5)X(3)=(X(3)/100)
C   THESE ARE THE CONDITIONS OF SPECIES 3 ENCYSTMENT
      IF(X(23).LT.50.)X(13)=0.1*X(3)+X(13)
      IF(X(4).GE.100.)X(21)=(X(3)*0.1)+X(21)
C   THIS IS WHAT HAPPENS IF SPECIES 3 RUNS OUT OF NUTRIENTS
      IF (X(23).LT.50.)X(3)=1.
C
C   THIS SECTION IS FOR SPECIES 4, AN OCEANIC FORM
C   THIS IS SPECIES 4 GROWTHRATE

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      IF((TEMP.GT.18.) .AND. (SAL.GT.3.0))X(4)=(X(4)+30.)
      IF(X(3).LE.100.)X(4)=1.02*X(4)
      IF(X(3).GT.100.)X(4)=X(4)*0.90
      IF((TEMP.LE.18.) .OR. (SAL.LE.3.0))X(4)=10.
      IF(TAMP.LT.2.5)X(4)=(X(4)/10)
C     THESE ARE THE CONDITIONS OF SPECIES 4 ENCYSTMENT
      IF(X(23).LT.50.)X(14)=(0.1*X(4))+X(14)
      IF(X(3).GE.100.)X(22)=(X(4)*0.1)+X(22)
C     THIS IS WHAT HAPPENS IF SPECIES 4 RUNS OUT OF NUTRIENTS
      IF (X(23).LT.50.)X(4)=1.
C
C     THIS SECTION IS FOR SPECIES 8, A MID-DEPTH COSMOPOLITAN FORM
C     THIS IS SPECIES 8 GROWTHRATE
      V823=1.0*X(8)
      IF((TEMP.GT.18.) .AND. (SAL.GT.2.0))X(8)=1.02*X(8)
      IF((TEMP.GT.18.) .AND. (TAMP.LT.2.))X(8)=1.005*X(8)
      IF((TEMP.LE.18.) .OR. (SAL.LE.2.0))X(8)=1.
C     THESE ARE THE CONDITIONS OF SPECIES 8 ENCYSTMENT
      IF(X(23).LT.50.)X(18)=(0.1*X(8))+X(18)
      IF((TDY.GE.319.) .AND. (TDY.LT.319.3))X(18)=(0.5*X(8))+X(18)
C     THIS IS WHAT HAPPENS IF SPECIES 8 RUNS OUT OF NUTRIENTS
      IF(X(23).LT.50.)X(8)=1.
C
C     THIS SECTION IS FOR SPECIES 7, A MIDDEPTH FORM
C     THIS IS SPECIES 7 GROWTHRATE
      V723=1.0*X(7)
      IF(((SAL.GT.1.5) .AND. (TEMP.GT.17.)) .AND. (X(8).LT.10.))
      +X(7)=1.01*X(7)
      IF(X(8).GE.500.)X(7)=5.
      IF((SAL.LE.1.5) .OR. (TEMP.LE.17.))X(7)=1.
C     THESE ARE CONDITIONS OF SPECIES 7 ENCYSTMENT
      IF(X(23).LT.50.)X(17)=(0.1*X(7))+X(17)
      IF((TDY.GE.200.) .AND. (TDY.LT.200.3))X(17)=(0.1*X(7))+X(17)
C     THIS IS WHAT HAPPENS IF SPECIES 7 RUNS OUT OF NUTRIENTS
      IF (X(23).LT.50.)X(7)=1.
C
C     THIS SECTION IS FOR SPECIES 6, A MIDDEPTH FORM
C     THIS IS SPECIES 6 GROWTHRATE
      V623=1.0*X(6)
      IF((SAL.LT.2.5) .OR. (X(8).GT.100.))X(6)=(.75*X(8))
      IF((SAL.GT.2.5) .AND. (X(8).LT.100.))X(6)=(1.008*X(6))
C     THESE ARE THE CONDITIONS OF SPECIES 6 ENCYSTMENT
      IF(X(23).LT.50.)X(16)=(0.1*X(6))+X(16)
      IF((TDY.GE.200.) .AND. (TDY.LT.200.3))X(16)=(0.1*X(6))+X(16)
C     THIS IS WHAT HAPPENS IF SPECIES 6 RUNS OUT OF NUTRIENTS
      IF(X(23).LT.50.)X(6)=1
C
C     THIS SECTION IS FOR SPECIES 5, A MID DEPTH COOLER FORM
C     THIS IS SPECIES 5 GROWTHRATE
      V523=1.0*X(5)
      IF(((TEMP.GT.15.) .AND. (TEMP.LT.18.)) .AND. ((SAL.GT.1.5) .AND.
      +(SAL.LT.3.0)))X(5)=1.01*X(5)
      IF((TEMP.LE.15.) .OR. (SAL.LE.1.5))X(5)=10.
      IF((TEMP.GE.18.) .OR. (SAL.GE.3.0))X(5)=1.
C     THESE ARE THE CONDITIONS OF SPECIES 5 ENCYSTMENT
      IF(X(23).LT.50.)X(15)=(0.1*X(5))+X(15)
      IF((TDY.GE.300.) .AND. (TDY.LT.300.3))X(15)=(0.1*X(5))+X(15)
C     THIS IS WHAT HAPPENS IF SPECIES 5 RUNS OUT OF NUTRIENTS
      IF(X(23).LT.50.)X(5)=1.
C

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```

C DIFFERENTIAL EQUATIONS
  XD(23)=Z1-V123-V223-V323-V423-V523-V623-V723-
+V823-V923-V1023
  XD(1)=V123-(1.0*X(1))
  XD(2)=V223-(1.0*X(2))
  XD(3)=V323-(1.0*X(3))
  XD(4)=V423-(1.0*X(4))
  XD(5)=V523-(1.0*X(5))
  XD(6)=V623-(1.0*X(6))
  XD(7)=V723-(1.0*X(7))
  XD(8)=V823-(1.0*X(8))
  XD(9)=V923-(1.0*X(9))
  XD(10)=V1023-(1.0*X(10))
C
C THE FOLLOWING ARE THE BASELINE CHANGES FOR ENCYSTMENT
  XD(11)=0.
  XD(12)=0.
  XD(13)=0.
  XD(14)=0.
  XD(15)=0.
  XD(16)=0.
  XD(17)=0.
  XD(18)=0.
  XD(19)=0.
  XD(20)=0.
  XD(21)=0.
  XD(22)=0.
C
C TA EQUALS TOTAL CYSTS IN SEDIMENT
  TA=X(11)+X(12)+X(13)+X(14)+X(15)+X(16)+X(17)+
+X(18)+X(19)+X(20)+X(21)+X(22)
C
C AC EQUALS THE PERCENT OF CYST TYPE IN TOTAL
  AC(1)=X(11)/TA
  AC(2)=X(12)/TA
  AC(3)=X(13)/TA
  AC(4)=X(14)/TA
  AC(5)=X(15)/TA
  AC(6)=X(16)/TA
  AC(7)=X(17)/TA
  AC(8)=X(18)/TA
  AC(9)=X(19)/TA
  AC(10)=X(20)/TA
  AC(11)=X(21)/TA
  AC(12)=X(22)/TA
  RETURN
  END
CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC
SUBROUTINE RKS (X, XD, T, DT, N, D, TA, AC, TEMP, SAL)
DIMENSION X(23), XD(23)
DIMENSION AC(12)
DIMENSION XP(23), XQ(23), XR(23), XPD(23), XQD(23), XRD(23)
DT2=DT/2
T2=T-DT2
TT=T+DT
CALL DFQS (X, XD, T, N, D, TA, AC, TEMP, SAL)
DO 1 I=1, 23
1 XP(I)=X(I)+XD(I)*DT2
CALL DFQS (XP, XPD, T2, N, D, TA, AC, TEMP, SAL)
DO 2 I=1, 23

```

```
2   XQ(I)=X(I)+XPD(I)*DT2
    CALL DFQS(XQ,XQD,T2,N,D,TA,AC,TEMP,SAL)
    DO 3 I=1,23
3   XR(I)=X(I)+XQD(I)*DT
    CALL DFQS(XR,XRD,TT,N,D,TA,AC,TEMP,SAL)
    DO 4 I=1,23
4   X(I)=X(I)+(XD(I)+2*XPD(I)+2*XQD(I)+XRD(I))*(DT/6.)
    RETURN
    END
```



(DEPTH= 60.0 METERS) MODEL X OUTPUT

TIME,	SP1	SP2	SP3	SP4	SP5	SP6	SP7	SP8	SP9	SP10
0.00	1.	1.	1.	1.	1.	1.	1.	1.	1.	1.
0.08	1.	1.	10.	10.	10.	1.	1.	1.	1.	1.
0.17	1.	1.	10.	10.	10.	1.	1.	1.	1.	1.
0.25	1.	13.	10.	10.	10.	1.	1.	1.	1.	1.
0.33	1.	720.	10.	10.	1.	1.	1.	1.	1.	1.
0.41	52.	1.	10.	10.	1.	1.	2.	1.	1.	1.
0.50	2961.	1.	10.	10.	1.	1.	4.	1.	1.	1.
0.58	1.	1.	10.	10.	1.	1.	1.	1.	1.	1.
0.66	1.	1.	10.	10.	1.	2.	3.	7.	1.	7.
0.75	1.	3.	10.	10.	1.	25.	3.	34.	1.	34.
0.83	1.	159.	10.	10.	2.	1.	1.	1.	1.	1.
0.91	1.	1.	10.	10.	10.	1.	1.	1.	1.	1.
1.00	1.	1.	10.	10.	10.	1.	1.	1.	1.	1.

(DEPTH= 70.0 METERS)

TIME,	SP1	SP2	SP3	SP4	SP5	SP6	SP7	SP8	SP9	SP10
0.00	1.	1.	1.	1.	1.	1.	1.	1.	1.	1.
0.08	1.	1.	10.	10.	10.	1.	1.	1.	1.	1.
0.17	1.	1.	10.	10.	10.	1.	1.	1.	1.	1.
0.25	1.	13.	10.	10.	10.	1.	1.	1.	1.	1.
0.33	1.	720.	10.	10.	1.	1.	1.	1.	1.	1.
0.41	52.	1.	10.	10.	1.	1.	2.	1.	1.	1.
0.50	2961.	1.	10.	10.	1.	1.	4.	1.	1.	1.
0.58	1.	1.	10.	10.	1.	1.	1.	1.	1.	1.
0.66	1.	1.	10.	10.	1.	2.	3.	7.	1.	7.
0.75	1.	3.	10.	10.	1.	25.	3.	34.	1.	34.
0.83	1.	159.	10.	10.	2.	1.	1.	1.	1.	1.
0.91	1.	1.	10.	10.	10.	1.	1.	1.	1.	1.
1.00	1.	1.	10.	10.	10.	1.	1.	1.	1.	1.

(DEPTH= 80.0 METERS)

TIME,	SP1	SP2	SP3	SP4	SP5	SP6	SP7	SP8	SP9	SP10
0.00	1.	1.	1.	1.	1.	1.	1.	1.	1.	1.
0.08	1.	1.	1127.	270.	1.	2.	2.	5.	12.	5.
0.17	1.	1.	1609.	270.	1.	4.	3.	27.	135.	27.
0.25	1.	1.	1819.	270.	1.	103.	3.	138.	1560.	138.
0.33	1.	1.	96.	127.	1.	1.	1.	1.	1.	1.
0.41	1.	1.	188.	186.	1.	1.	1.	1.	1.	1.
0.50	1.	1.	276.	220.	1.	1.	1.	1.	2.	1.
0.58	1.	1.	63.	95.	1.	1.	1.	1.	1.	1.
0.66	1.	1.	223.	202.	1.	1.	1.	1.	1.	1.
0.75	1.	1.	63.	95.	1.	1.	1.	1.	1.	1.
0.83	1.	1.	170.	176.	1.	1.	1.	1.	1.	1.
0.91	1.	1.	1.	1.	1.	1.	1.	1.	1.	1.
1.00	1.	1.	63.	95.	1.	1.	1.	1.	1.	1.

(DEPTH= 60.0 METERS) MODEL Y OUTPUT

TIME,	SP1	SP2	SP3	SP4	SP5	SP6	SP7	SP8	SP9	SP10
0.00	1.	1.	1.	1.	1.	1.	1.	1.	1.	1.
0.08	1.	57.	10.	10.	1.	2.	1.	1.	1.	1.
0.17	1.	3265.	10.	10.	1.	4.	1.	1.	1.	1.
0.25	1.	1.	1.	1.	1.	1.	1.	1.	1.	1.
0.33	13.	1.	10.	10.	1.	1.	2.	1.	1.	1.
0.41	720.	1.	10.	10.	1.	5.	5.	7.	1.	7.
0.50	1.	1.	10.	10.	1.	2.	2.	4.	1.	4.
0.58	1.	1.	114.	141.	1.	1.	1.	1.	1.	1.
0.66	1.	1.	1.	1.	1.	1.	1.	1.	1.	1.
0.75	1.	1.	96.	127.	1.	1.	1.	1.	1.	1.
0.83	1.	9.	10.	10.	1.	1.	2.	2.	1.	2.
0.91	1.	488.	10.	10.	2.	1.	4.	1.	1.	1.
1.00	1.	29.	10.	10.	1.	2.	1.	1.	1.	1.

(DEPTH= 70.0 METERS)

TIME,	SP1	SP2	SP3	SP4	SP5	SP6	SP7	SP8	SP9	SP10
0.00	1.	1.	1.	1.	1.	1.	1.	1.	1.	1.
0.08	1.	57.	10.	10.	1.	2.	1.	1.	1.	1.
0.17	1.	3265.	10.	10.	1.	4.	1.	1.	1.	1.
0.25	1.	1.	1.	1.	1.	1.	1.	1.	1.	1.
0.33	13.	1.	10.	10.	1.	1.	2.	1.	1.	1.
0.41	720.	1.	10.	10.	1.	5.	5.	7.	1.	7.
0.50	1.	1.	10.	10.	1.	2.	2.	4.	1.	4.
0.58	1.	1.	114.	141.	1.	1.	1.	1.	1.	1.
0.66	1.	1.	1.	1.	1.	1.	1.	1.	1.	1.
0.75	1.	1.	96.	127.	1.	1.	1.	1.	1.	1.
0.83	1.	9.	10.	10.	1.	1.	2.	2.	1.	2.
0.91	1.	488.	10.	10.	2.	1.	4.	1.	1.	1.
1.00	1.	29.	10.	10.	1.	2.	1.	1.	1.	1.

(DEPTH= 80.0 METERS)

TIME,	SP1	SP2	SP3	SP4	SP5	SP6	SP7	SP8	SP9	SP10
0.00	1.	1.	1.	1.	1.	1.	1.	1.	1.	1.
0.08	1.	1.	1127.	270.	1.	2.	2.	5.	12.	5.
0.17	1.	1.	1609.	270.	1.	4.	3.	27.	135.	27.
0.25	1.	1.	1819.	270.	1.	103.	3.	138.	1560.	138.
0.33	1.	1.	1.	1.	1.	1.	1.	1.	1.	1.
0.41	1.	1.	133.	154.	1.	1.	1.	1.	1.	1.
0.50	1.	1.	1.	32.	1.	1.	1.	1.	1.	1.
0.58	1.	1.	114.	141.	1.	1.	1.	1.	1.	1.
0.66	1.	1.	1.	32.	1.	1.	1.	1.	1.	1.
0.75	1.	1.	96.	127.	1.	1.	1.	1.	1.	1.
0.83	1.	1.	1.	1.	1.	1.	1.	1.	1.	1.
0.91	1.	1.	114.	141.	1.	1.	1.	1.	1.	1.
1.00	1.	1.	1.	32.	1.	1.	1.	1.	1.	1.

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C     DINOFLAGELLATE MODEL Z
C     A DINOFLAGELLATE ECOSYSTEM DYNAMICS MODEL
C     5 DEGREES WARMER DEPTH CHANGE AND CYSTS (VERSION WITH FRONT)
C     BY J. C. CAWLEY, 1996
C     X23 PHYTOPLANKTON POTENTIAL (NUTRIENT)
C     X1-X10 SPECIES 1-10
C     X11-22 CYSTS TO SEDIMENT
C
C     IMPLICIT REAL (L,M)
C     DIMENSION X(23),XD(23)
C     DIMENSION AC(12)
C     COMMON TDY
C     REAL M41,M42,M43
C     OPEN (6,FILE='CON')
C     OPEN (7,FILE='C:OUT.DAT')
C SET INITIAL CONDITIONS
DATA X /1.,1.,1.,1.,1.,1.,1.,1.,1.,1.,1.,1.,1.,
+1.,1.,1.,1.,1.,1.,1.,1.,1.,1.,500./
DATA AC /1.,1.,1.,1.,1.,1.,1.,1.,1.,1.,1.,1./
D=05.
TEMP=19.
SAL=2.
C TA=TOTAL CYSTS IN SEDIMENT
TA=1.
N=4
T=0.
TMAX=40.
C DT IS 0.001 YEAR
DT=.001
C IOUTr IS THE NUMBER OF ITERATIONS PER OUTPUT
IOUTr=2000
C IOUT IS THE COUNTER FOR THE OUTPUT INTERVAL
IOUT=IOUTr+1
C WRITE COLUMN HEADINGS
WRITE(6,1)
WRITE(7,1)
1  FORMAT(' TIME,  SP1 SP2 SP3 SP4 SP5 SP6 SP7 SP8 SP9 SP10 SP11
+SP12')
C
C BEGINNING OF CALCULATION LOOP
3  CONTINUE
IF(IOUT.LT.IOUTr)GO TO 2
WRITE(6,5)D
WRITE(7,5)D
5  FORMAT(' TIME (DEPTH=',F5.1,')')
WRITE(6,4)T,(AC(I),I=1,12)
WRITE(7,4)T,(AC(I),I=1,12)
4  FORMAT(' ',F5.2,12F5.2)
D={D+5.0)
AC(1)=0.
AC(2)=0.
AC(3)=0.
AC(4)=0.
AC(5)=0.
AC(6)=0.
AC(7)=0.
AC(8)=0.
AC(9)=0.
AC(10)=0.
TA=0.

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      IOUT=0
2     IOUT=IOUT+1
C     CHECK FOR END OF SIMULATION
      IF (T.GE.TMAX) GO TO 99
      CALL RKS (X, XD, T, DT, N, D, TA, AC, TEMP, SAL)
      T=T+DT
      GO TO 3
99    STOP
      END
CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC
      SUBROUTINE DFQS (X, XD, T, N, D, TA, AC, TEMP, SAL)
      IMPLICIT REAL (L,M)
      DIMENSION XD(23), X(23)
      DIMENSION MONTH(14), S(14)
      DIMENSION AC(12)
C     COMMON TDY
C
C
C     THIS SUBROUTINE CONTAINS THE DIFFERENTIAL EQUATIONS
C
C     STATEMENTS BELOW ARE USED TO FLUCTUATE TIME FOR SEASONS
      TWY=T-AINT(T)
C     TDY IS TIME WITHIN YEAR IN DAYS
      TDY=TWY*365.
C     TR IS TWY IN RADIANS
      TR=(TWY)*6.28318
C     MONTH IS THE MIDDLE DAY OF EACH MONTH
      DATA MONTH/0.,15.,46.,74.,105.,136.,166.,196.,
+227.,258.,288.,319.,349.,365./
C
C     STATEMENTS BELOW VARY TEMP AS A SEASONAL SINE FUNCTION
C     AMT IS THE ANNUAL MEAN WATER TEMPERATURE
      IF (D.LT.25.) AMT=17.
      IF ((D.GE.25.) .AND. (D.LT.75.)) AMT=18
      IF (D.GE.75.) AMT=25.
C     TAMP IS THE AMPLITUDE OF THE TEMPERTATURE CURVE
      IF (D.LE.25.) TAMP=10.0
      IF ((D.GT.25.) .AND. (D.LE.50.)) TAMP=7.0
      IF ((D.GT.50.) .AND. (D.LE.75.)) TAMP=6.0
      IF (D.GT.75.) TAMP=4.0
C     TPS IS THE PHASE SHIFT OF TEMP CURVE IN RADIANS
      TPS=1.75585
      TEMP=AMT+SIN(TR-TPS)*TAMP
      FLAG=1
      IF (TEMP.LT.18.) FLAG=0.
C
C     STATEMENTS BELOW VARY SALINITY THROUGH TIME
      DATA S/1.2,1.1,1.1,0.8,0.6,0.8,0.8,1.1,
+1.4,1.4,1.0,0.8,1.1,1.2/
      DO 101 I=1,14
101    IF (MONTH(I).GT.TDY) GOTO 102
102    HFA=(TDY-MONTH(I-1))/(MONTH(I)-MONTH(I-1))
      HMC=S(I)-S(I-1)
      SAL=(HFA*HMC)+S(I-1)
      IF (D.LT.15.) SAL=(SAL*1.4)
      IF ((D.GE.15.) .AND. (D.LE.25.)) SAL=(SAL*1.8)
      IF ((D.GT.25.) .AND. (D.LE.75.)) SAL=(SAL*2.0)
      IF (D.GT.75.) SAL=3.4
C     INPUTS, Z IS DEPENDANT ON DEPTH IN METERS
      Z=150.

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IF(D.GT.75.)Z1=100.
IF((D.LE.75.).AND.(D.GT.25.))Z1=300.
IF(D.LE.25.)Z1=750.
C
C THIS SECTION IS FOR SPECIES 1, A NEARSHORE 'BLOOMER'
C THIS IS SPECIES 1 GROWTHRATE:
V123=1.0*X(1)
IF((TAMP.LT.4.).OR.(SAL.GE.2.5))X(1)=1.
IF((TEMP.GE.20.).AND.(SAL.GT.1.))X(1)=(1.05*X(1))
IF((TEMP.LT.15.).AND.(X(1).GE.10.))X(1)=(0.95*X(1))
C THESE ARE CONDITIONS OF SPECIES 1 ENCYSTMENT:
IF(X(23).LT.50.)X(11)={(0.1*X(1))+X(11)}
IF((TEMP.LT.23.02).AND.(TEMP.GT.23.))X(11)={(0.1*X(1))-X(11)}
C THIS IS WHAT HAPPENS IF SPECIES 1 RUNS OUT OF NUTRIENTS:
IF(X(23).LT.50.)X(1)=1.
C
C THIS SECTION IS FOR SPECIES 10, AN OCEANIC FORM
C THIS IS SPECIES 10 GROWTHRATE
V1023=1.0*X(10)
IF((TEMP.LT.18.).OR.(SAL.GT.2.0))X(10)=(1.02*X(10))
IF((TEMP.LT.18.).OR.(SAL.LT.2.0))X(10)=1.
C THESE ARE CONDITIONS OF SPECIES 10 ENCYSTMENT
IF((TEMP.LT.18.02).AND.(FLAG.EQ.1.))X(20)={(0.50*X(10))+
-X(20)}
C THIS IS WHAT HAPPENS IF SPECIES 10 RUNS OUT OF NUTRIENTS
IF(X(23).LT.50.)X(10)=1.
C
C THIS SECTION IS FOR SPECIES 2, A COOLER NEARSHORE BLOOMER
C THIS IS SPECIES 2 GROWTHRATE
V223=1.0*X(2)
IF(((TEMP.GT.15.).AND.(TEMP.LT.20.)).AND.(SAL.GT.1.))X(2)=
+(1.05*X(2))
IF(((TEMP.LT.15.).OR.(TEMP.GT.20.)).OR.(SAL.LE.1.)))
+X(2)=1.
IF(TAMP.LT.2.0)X(2)=1.
C THESE ARE CONDITIONS OF SPECIES 2 ENCYSTMENT
IF(X(23).LT.50)X(12)=(0.1*X(2))+X(12)
IF((TEMP.LT.15.2).AND.(TEMP.GE.15.0))X(12)=(0.05*X(2))+X(12)
IF((TEMP.GT.17.08).AND.(TEMP.LT.18.))X(12)=(0.05*X(2))+X(12)
C THIS IS WHAT HAPPENS IF SPECIES 2 RUNS OUT OF NUTRIENTS
IF(X(23).LT.50.)X(2)=1.
C
C THIS SECTION IS FOR SPECIES 9, AN OCEANIC FORM
C THIS IS SPECIES 9 GROWTHRATE
V923=1.0*X(9)
IF((TEMP.GE.18.).AND.(SAL.GT.3.2))X(9)=1.03*X(9)
IF((TEMP.LT.18.).OR.(SAL.LT.3.2))X(9)=1.
C THESE ARE THE CONDITIONS OF SPECIES 9 ENCYSTMENT
DO 111 I=1,14
111 IF(MONTH(I).GE.TDY)X(19)={(0.05*X(9))+X(19)}
C THIS IS WHAT HAPPENS IF SPECIES 9 RUNS OUT OF NUTRIENTS
IF(X(23).LT.50.)X(9)=1.
C
C THIS SECTION IS FOR SPECIES 3, AN OCEANIC FORM, ASSOC SP 4
C THIS IS SPECIES 3 GROWTHRATE
V323=1.0*X(3)
IF((TEMP.GT.20.).AND.(SAL.GT.3.0))X(3)={X(3)-20.}
IF(X(4).LE.100.)X(3)={1.01*X(4)}
IF(X(4).GT.100.)X(3)={X(3)*0.99}
IF((TEMP.LE.18.).OR.(SAL.LE.3.0))X(3)=10.

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      IF(TAMP.LT.2.5)X(3)=(X(3)/100)
C   THESE ARE THE CONDITIONS OF SPECIES 3 ENCYSTMENT
      IF(X(23).LT.50.)X(13)=0.1*X(3)+X(13)
      IF(X(4).GE.100.)X(21)=(X(3)*0.1)+X(21)
C   THIS IS WHAT HAPPENS IF SPECIES 3 RUNS OUT OF NUTRIENTS
      IF (X(23).LT.50.)X(3)=1.
C
C   THIS SECTION IS FOR SPECIES 4, AN OCEANIC FORM
C   THIS IS SPECIES 4 GROWTHRATE
      V423=1.0*X(4)
      IF((TEMP.GT.18.) .AND. (SAL.GT.3.0))X(4)=(X(4)+30.)
      IF(X(3).LE.100.)X(4)=1.02*X(4)
      IF(X(3).GT.100.)X(4)=X(4)*0.90
      IF((TEMP.LE.18.) .OR. (SAL.LE.3.0))X(4)=10.
      IF(TAMP.LT.2.5)X(4)=(X(4)/10)
C   THESE ARE THE CONDITIONS OF SPECIES 4 ENCYSTMENT
      IF(X(23).LT.50.)X(14)=(0.1*X(4))+X(14)
      IF(X(3).GE.100.)X(22)=(X(4)*0.1)+X(22)
C   THIS IS WHAT HAPPENS IF SPECIES 4 RUNS OUT OF NUTRIENTS
      IF (X(23).LT.50.)X(4)=1.
C
C   THIS SECTION IS FOR SPECIES 8, A MID-DEPTH COSMOPOLITAN FORM
C   THIS IS SPECIES 8 GROWTHRATE
      V823=1.0*X(8)
      IF((TEMP.GT.18.) .AND. (SAL.GT.2.0))X(8)=1.02*X(8)
      IF((TEMP.GT.18.) .AND. (TAMP.LT.2.))X(8)=1.005*X(8)
      IF((TEMP.LE.18.) .OR. (SAL.LE.2.0))X(8)=1.
C   THESE ARE THE CONDITIONS OF SPECIES 8 ENCYSTMENT
      IF(X(23).LT.50.)X(18)=(0.1*X(8))+X(18)
      IF((TDY.GE.319.) .AND. (TDY.LT.319.3))X(18)=(0.5*X(8))+X(18)
C   THIS IS WHAT HAPPENS IF SPECIES 8 RUNS OUT OF NUTRIENTS
      IF(X(23).LT.50.)X(8)=1.
C
C   THIS SECTION IS FOR SPECIES 7, A MIDDEPTH FORM
C   THIS IS SPECIES 7 GROWTHRATE
      V723=1.0*X(7)
      IF(((SAL.GT.1.5) .AND. (TEMP.GT.17.)) .AND. (X(8).LT.10.))
      +X(7)=1.01*X(7)
      IF(X(8).GE.500.)X(7)=5.
      IF((SAL.LE.1.5) .OR. (TEMP.LE.17.))X(7)=1.
C   THESE ARE CONDITIONS OF SPECIES 7 ENCYSTMENT
      IF(X(23).LT.50.)X(17)=(0.1*X(7))+X(17)
      IF((TDY.GE.200.) .AND. (TDY.LT.200.3))X(17)=(0.1*X(7))+X(17)
C   THIS IS WHAT HAPPENS IF SPECIES 7 RUNS OUT OF NUTRIENTS
      IF (X(23).LT.50.)X(7)=1.
C
C   THIS SECTION IS FOR SPECIES 6, A MIDDEPTH FORM
C   THIS IS SPECIES 6 GROWTHRATE
      V623=1.0*X(6)
      IF((SAL.LT.2.5) .OR. (X(8).GT.100.))X(6)=(.75*X(8))
      IF(SAL.GT.2.5 .AND. (X(8).LT.100.))X(6)=(1.008*X(6))
C   THESE ARE THE CONDITIONS OF SPECIES 6 ENCYSTMENT
      IF(X(23).LT.50.)X(16)=(0.1*X(6))+X(16)
      IF((TDY.GE.200.) .AND. (TDY.LT.200.3))X(16)=(0.1*X(6))+X(16)
C   THIS IS WHAT HAPPENS IF SPECIES 6 RUNS OUT OF NUTRIENTS
      IF(X(23).LT.50.)X(6)=1
C
C   THIS SECTION IS FOR SPECIES 5, A MID DEPTH COOLER FORM
C   THIS IS SPECIES 5 GROWTHRATE
      V523=1.0*X(5)

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      IF((TEMP.GT.15.).AND.(TEMP.LT.18.)).AND.((SAL.GT.1.5).AND.
+ (SAL.LT.3.0))X(5)=1.01*X(5)
      IF((TEMP.LE.15.).OR.(SAL.LE.1.5))X(5)=10.
      IF((TEMP.GE.18.).OR.(SAL.GE.3.0))X(5)=1.
C   THESE ARE THE CONDITIONS OF SPECIES 5 ENCYSTMENT
      IF(X(23).LT.50.)X(15)=(0.1*X(5))+X(15)
      IF((TDY.GE.300.).AND.(TDY.LT.300.3))X(15)=(0.1*X(5))+X(15)
C   THIS IS WHAT HAPPENS IF SPECIES 5 RUNS OUT OF NUTRIENTS
      IF(X(23).LT.50.)X(5)=1.
C
C   DIFFERENTIAL EQUATIONS
      XD(23)=Z1-V123-V223-V323-V423-V523-V623-V723-
+V823-V923-V1023
      XD(1)=V123-(1.0*X(1))
      XD(2)=V223-(1.0*X(2))
      XD(3)=V323-(1.0*X(3))
      XD(4)=V423-(1.0*X(4))
      XD(5)=V523-(1.0*X(5))
      XD(6)=V623-(1.0*X(6))
      XD(7)=V723-(1.0*X(7))
      XD(8)=V823-(1.0*X(8))
      XD(9)=V923-(1.0*X(9))
      XD(10)=V1023-(1.0*X(10))
C
C   THE FOLLOWING ARE THE BASELINE CHANGES FOR ENCYSTMENT
      XD(11)=0.
      XD(12)=0.
      XD(13)=0.
      XD(14)=0.
      XD(15)=0.
      XD(16)=0.
      XD(17)=0.
      XD(18)=0.
      XD(19)=0.
      XD(20)=0.
      XD(21)=0.
      XD(22)=0.
C
C   TA EQUALS TOTAL CYSTS IN SEDIMENT
      TA=X(11)+X(12)+X(13)+X(14)+X(15)+X(16)+X(17)+
+X(18)+X(19)+X(20)+X(21)+X(22)
C
C   AC EQUALS THE PERCENT OF CYST TYPE IN TOTAL
      AC(1)=X(11)/TA
      AC(2)=X(12)/TA
      AC(3)=X(13)/TA
      AC(4)=X(14)/TA
      AC(5)=X(15)/TA
      AC(6)=X(16)/TA
      AC(7)=X(17)/TA
      AC(8)=X(18)/TA
      AC(9)=X(19)/TA
      AC(10)=X(20)/TA
      AC(11)=X(21)/TA
      AC(12)=X(22)/TA
      RETURN
      END
CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC
      SUBROUTINE RKS(X,XD,T,DT,N,D,TA,AC,TEMP,SAL)
      DIMENSION X(23),XD(23)

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DIMENSION AC(12)
DIMENSION XP(23), XQ(23), XR(23), XPD(23), XQD(23), XRD(23)
DT2=DT/2
T2=T+DT2
TT=T+DT
CALL DFQS(X, XD, T, N, D, TA, AC, TEMP, SAL)
DO 1 I=1, 23
1  XP(I)=X(I)+XD(I)*DT2
   CALL DFQS(XP, XPD, T2, N, D, TA, AC, TEMP, SAL)
   DO 2 I=1, 23
2  XQ(I)=X(I)+XPD(I)*DT2
   CALL DFQS(XQ, XQD, T2, N, D, TA, AC, TEMP, SAL)
   DO 3 I=1, 23
3  XR(I)=X(I)+XQD(I)*DT
   CALL DFQS(XR, XRD, TT, N, D, TA, AC, TEMP, SAL)
   DO 4 I=1, 23
4  X(I)=X(I)+(XD(I)+2*XPD(I)+2*XQD(I)+XRD(I))*(DT/6.)
   RETURN
END

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MODEL Z OUTPUT

TIME,	SP1	SP2	SP3	SP4	SP5	SP6	SP7	SP8	SP9	SP10	SP11	SP12
TIME (DEPTH= 5.0)	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00
0.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00
TIME (DEPTH= 10.0)	0.90	0.00	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.08	0.00	0.00
2.00	0.90	0.00	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.08	0.00	0.00
TIME (DEPTH= 15.0)	0.89	0.00	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.09	0.00	0.00
4.00	0.89	0.00	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.09	0.00	0.00
TIME (DEPTH= 20.0)	0.89	0.00	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.09	0.00	0.00
6.00	0.89	0.00	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.09	0.00	0.00
TIME (DEPTH= 25.0)	0.89	0.00	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.09	0.00	0.00
8.00	0.89	0.00	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.09	0.00	0.00
TIME (DEPTH= 30.0)	0.87	0.01	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.10	0.00	0.00
10.00	0.87	0.01	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.10	0.00	0.00
TIME (DEPTH= 35.0)	0.85	0.01	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.10	0.00	0.00
12.00	0.85	0.01	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.10	0.00	0.00
TIME (DEPTH= 40.0)	0.84	0.02	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.11	0.00	0.00
14.00	0.84	0.02	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.11	0.00	0.00
TIME (DEPTH= 45.0)	0.83	0.02	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.12	0.00	0.00
16.00	0.83	0.02	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.12	0.00	0.00
TIME (DEPTH= 50.0)	0.82	0.02	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.12	0.00	0.00
18.00	0.82	0.02	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.12	0.00	0.00
TIME (DEPTH= 55.0)	0.81	0.03	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.13	0.00	0.00
20.00	0.81	0.03	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.13	0.00	0.00
TIME (DEPTH= 60.0)	0.79	0.04	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.13	0.01	0.00
22.00	0.79	0.04	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.13	0.01	0.00
TIME (DEPTH= 65.0)	0.78	0.05	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.14	0.01	0.00
24.00	0.78	0.05	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.14	0.01	0.00
TIME (DEPTH= 70.0)	0.77	0.05	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.14	0.01	0.00
26.00	0.77	0.05	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.14	0.01	0.00
TIME (DEPTH= 75.0)	0.77	0.06	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.14	0.01	0.00
28.00	0.77	0.06	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.14	0.01	0.00
TIME (DEPTH= 80.0)	0.61	0.04	0.06	0.08	0.00	0.00	0.00	0.00	0.00	0.12	0.00	0.06
30.00	0.61	0.04	0.06	0.08	0.00	0.00	0.00	0.00	0.00	0.12	0.00	0.06
TIME (DEPTH= 85.0)	0.52	0.04	0.09	0.13	0.00	0.00	0.00	0.00	0.00	0.11	0.00	0.09
31.99	0.52	0.04	0.09	0.13	0.00	0.00	0.00	0.00	0.00	0.11	0.00	0.09
TIME (DEPTH= 90.0)	0.45	0.03	0.11	0.17	0.00	0.00	0.00	0.00	0.00	0.10	0.00	0.12
33.99	0.45	0.03	0.11	0.17	0.00	0.00	0.00	0.00	0.00	0.10	0.00	0.12
TIME (DEPTH= 95.0)	0.40	0.03	0.13	0.20	0.00	0.00	0.00	0.00	0.00	0.10	0.00	0.13
35.99	0.40	0.03	0.13	0.20	0.00	0.00	0.00	0.00	0.00	0.10	0.00	0.13
TIME (DEPTH=100.0)	0.36	0.03	0.14	0.22	0.00	0.00	0.00	0.00	0.00	0.09	0.00	0.15
37.99	0.36	0.03	0.14	0.22	0.00	0.00	0.00	0.00	0.00	0.09	0.00	0.15
TIME (DEPTH=105.0)	0.32	0.03	0.15	0.24	0.00	0.00	0.00	0.00	0.00	0.09	0.00	0.16
39.99	0.32	0.03	0.15	0.24	0.00	0.00	0.00	0.00	0.00	0.09	0.00	0.16

**APPENDIX C:**

**PLANTS 77 FORTRAN CLUSTER ANALYSIS PROGRAM**

```

      DEBUG SUBCHK
C   DEBUG TRACE
C   AT 4839
C   TRACE ON
C   AT 12345
C   TRACE OFF
C   END DEBUG
C*****
*
C   PROGRAM PLANTS2 : PERFORMS CLUSTER ANALYSIS OF INPUT DATA.
*
C   WRITTEN BY PLANTS, REVISED BY ARNIE MILLER & TOM ROUNDS      *
C   MODIFIED BY BRET BENNINGTON                                  *
C
C   FILEDEFS : 04 = INPUT OF PLOTTING AND CONTROL PARAMETERS      *
C   05 = INPUT OF DATA SET                                       *
C   06 = OUTPUT OF TEXT OF RESULTS                                *
C   07 = OUTPUT OF DATA SET (+/- INVERS. OR TRANSF.)           *
C   10 = OUTPUT OF PLOTTING INSTRUCTIONS                          *
C
C*****
*
C THE INPUT DATA WILL USUALLY BE IN THE FOLLOWING FORM:          *
C
C       N ROWS : FOR SAMPLES(I),I=1,N_SAMPLES                    *
C       BY  M COLUMNS : FOR SPECIES(J),J=1,M_SPECIES            *
C
C*****
*
C CONTROL CARD PARAMETER: ITEST (USED TO SPECIFY Q OR R MODE)
*
C ITEST = +1 : NO INVERSION OF DATA MATRIX (Q-MODE)             *
C ITEST = -1 : INVERSION OF DATA MATRIX (R-MODE)               *
C ITEST = 00 : ENDS RUN OF PROGRAM (THIS MUST BE ON THE VERY LAST
*
C       CARD FOR THE PROGRAM TO END SUCESSFULLY)                 *
C       ALSO MAY BE USED TO CONVERT/INVERT DATA W/O
CALCULATING*
C       SIMILARITY COEFFICIENTS & W/O PLOTTING IF USED WITH     *
C       NCOEFF -1                                                *
C
C SIM. COEFF. CODE: MCOEFF                                       *
C MCOEFF = 01 : NO TRANS. + JACCARD COEFF. (P/A)                 *
C MCOEFF = 02 : NO TRANS. + CZEK. (DICE) COEFF.                 *
C MCOEFF = 03 : PERCENT TRANS. + CZEK. COEFF.                   *
C MCOEFF = 04 : PERCENT MAX. TRANS. + CZEK. COEFF.              *
C MCOEFF = 05 : LOG10 TRANS. + CZEK. COEFF.                     *
C MCOEFF = 06 : NO TRANS. + HOMOGEN. FUNCTION                   *
C MCOEFF = 07 : PERCENT TRANS. + HOMOGEN. FUNCTION               *
C MCOEFF = 08 : PERCENT MAX. TRANS. + HOMOGEN. FUNCTION         *
C MCOEFF = 09 : LOG10 TRANS. + HOMOGEN. FUNCTION                *
C MCOEFF = 10 : D'BLE TRANS. IF INVERSION ELSE PERCENT TRANS., +
DICE *
C       D'BLE TRANS = PERCENT TRANS. FOLLOWED BY PERCENT      *

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```

C          TRANS. AFTER INVERSION.          *
C MCOEFF = 11 : PERCENT TRANS + FREEMAN-TUKEY TRANS. + CZEK.
COEFF. *
C MCOEFF = 12 : PERCENT TRANS. + CHORD DISTANCE COEFF. (CRD)      *
C
C NEW SIM. COEFF CODE : NCOEFF          *
C NCOEFF = -1 :TERMINATES RUN AFTER DATA TRANSFORMATION AND/OR
*
C          INVERSION WITHOUT CALCULATING SIMILARITY COEFFICIENTS *
C          AND WITHOUT PRODUCING A DENDROGRAM IF USED WITH      *
C          ITEST 00          *
C NCOEFF = 01 TO 09 : SAME AS MCOEFF          *
C*****
*
  INTEGER TITLE(20),FORMT(20),CSIM(7),LABEL(12),BLANK
  INTEGER M,N,ITEST,MCOEFF,NCOEFF,VAL1,VAL2,D,I,J,K,MT,NX,

*NPASS,IM1,LVROW,LVCOL,NBROW,NBCOL,NXM1,JADJ,NROWS,NUM,KOLT
,KOLF.
  *NROWT,IX,IXM1,NUMX,IC,NM1,NP1,IBOOK,INDEX1,INDEX2,NP,IPASS
  REAL
  PERC,PMAX,PQ,PNO,V,W,VALMIN,VALMAX,SMAX,COMBO,HEIGHT,LENGTH
,
  *X,Y,YSCALE,CHARHT,XNUM,XSYM,YPOS,CONST
  REAL MULTTOT,SQTOTI,SQTOTK,CCOS,CRD
  INTEGER ROWCD(200,12),TEMCD(200,12),NUMBR(200)
  INTEGER COLCD(200,12)
  REAL DATA(200,200),OUT(200,3),SIMCO(200,200),BOOK(200,200),
  *SUM(200),T(200),PLOCX(200),PLOCY(200),TMP(200,200)
  EQUIVALENCE (SIMCO,BOOK)
  DATA BLANK,IPASS/0,0/
4839 CALL PLOTS(0.0,50)
C*****
*
C READ IN TITLE HEADING (UP TO 80 CHARACTERS)
4840 READ (5,100) TITLE
  WRITE (6,100) TITLE
100 FORMAT(20A4)
C*****
*
C READ IN EXECUTION-TIME FORMAT FOR INPUT DATA
4841 READ (5,100) FORMT
  WRITE (6,100) FORMT
C*****
*
C READ IN N (#ROWS/SAMPLES) & M (#COLS/SPECIES) AND FIND MAX. OF N
OR M
4842 READ (5,101) N,M
  WRITE(6,101) N,M
101 FORMAT(I3.2X,I3)
  D=N
  IF(M.GT.N) D=M
C*****
*

```

```

C DIMENSIONALIZE ARRAYS
C FIND OUT HOW TO DIMENSIONALIZE ARRAYS AT EXECUTION TIME IF
POSSIBLE.
C*****
*
C INITIALIZE COLUMN AND ROW CODE ARRAYS
  DO 1234 J=1,M
  DO 5678 K=1,12
  COLCD(J,K)=BLANK
5678 CONTINUE
1234 CONTINUE
  DO 8765 I=1,N
  DO 4321 K=1,12
  ROWCD(I,K)=BLANK
4321 CONTINUE
8765 CONTINUE
C*****
*
C READ IN COLUMN CODES (12-A4 FIELDS EACH) FOR SPECIES, 1 TO A CARD.
23 DO 3 J=1,M
24 READ (5,100) (COLCD(J,K),K=1,12)
3 CONTINUE
C*****
*
C READ IN ROW CODES (TWO 4-CHAR. FIELDS EACH) (USUALLY SAMPLES)
C AND THE DATA MATRIX (N-ROWS BY M-COLUMNS)
25 DO 4 I=1,N
26 READ (5,100) (ROWCD(I,K),K=1,12)
27 READ (5,FORMAT) (DATA(I,J),J=1,M)
C READ (5,FORMAT) (ROWCD(I,K),K=1,2),(DATA(I,J),J=1,M)
4 CONTINUE
12345 CONTINUE
C*****
*
C READ IN HEIGHT AND LENGTH OF DENDROGRAM PLOT IN INCHES, ALONG
WITH
C THE VALUE OF "NEWPEN", THE PLOT LINE THICKNESS (1-NARROW TO 5-
WIDE).
C THEN CHECK FOR ILLEGAL VALUES AND SET TO MAXIMUM IF FOUND.
  READ (4,104) HEIGHT,LENGTH,NP
104 FORMAT(F4.1,1X,F4.1,1X,I1)
  IF(HEIGHT.GT.34.0) HEIGHT=34.0
C*****
*
C READ IN CONTROL CARD PARAMETER (ITEST), AND SIM. COEFF. CODE
(MCOEFF)
  READ (4,102) ITEST,MCOEFF,CONST
102 FORMAT(I2,1X,I2,1X,F7.3)
C*****
*
C TEST FOR ILLEGAL MCOEFF VALUE
  IF(MCOEFF.GE.1.OR.MCOEFF.LE.12) GO TO 835
  WRITE(6,105) MCOEFF
105 FORMAT('SIM COEFF. VALUE OF ',I2,' IS ILLEGAL.')

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      GO TO 999
C*****
*
C TEST FOR TRANSFORMATION TYPE (IF ANY)
835  IF(MCOEFF.EQ.1.OR.MCOEFF.EQ.2.OR.MCOEFF.EQ.6) GO TO 825
      IF(MCOEFF.EQ.4.OR.MCOEFF.EQ.8) GO TO 1226
      IF(MCOEFF.EQ.5.OR.MCOEFF.EQ.9) GO TO 1227
      IF(MCOEFF.EQ.11) GO TO 1228
C*****
*
C CONVERTS RAW DATA TO PERCENTAGES : (MCOEFF=03,07,10,12)
1225  PERC=0.
      DO 89 I=1,N
      DO 87 J=1,M
      PERC=PERC+DATA(I,J)
87   CONTINUE
      DO 88 K=1,M
      DATA(I,K)=(100*DATA(I,K)/PERC)
88   CONTINUE
      PERC=0.
89   CONTINUE
      GO TO 825
C*****
*
C PERFORM ARCSIN FREEMAN TUKEY TRANSFORMATION (MCOEFF=11)
1228  DO 404 I=1,N
      MULTTOT=0.0
      DO 408 J=1,M
      MULTTOT=MULTTOT+DATA(I,J)
408  CONTINUE
      DO 414 J=1,M
      DATA(I,J)=.5*(ASIN(SQRT(DATA(I,J)/(MULTTOT+1))))+
$     ASIN(SQRT((DATA(I,J)+1)/(MULTTOT+1)))
414  CONTINUE
404  CONTINUE
      IF (IPASS.GT.0) GO TO 830
420  GO TO 825
C*****
*
C COMPUTES A PERCENT MAXIMUM DATA TRANSFORMATION :
(MCOEFF=04,08)
1226  PMAX=0.
      DO 96 J=1,M
      DO 97 I=1,N
      IF(DATA(I,J).GT.PMAX) PMAX=DATA(I,J)
97   CONTINUE
      DO 98 I1=1,N
      DATA(I1,J)=(DATA(I1,J)/PMAX)*100.
98   CONTINUE
      PMAX=0
96   CONTINUE
      GO TO 825
C*****
*

```

```

C COMPUTES A LOG10 TRANSFORMATION (MCOEFF=05.09)
C IF NO CONSTANT IS SPECIFIED A VALUE OF 1.0 WILL BE USED.
C (NOTE : DECILES WILL BE COMPUTED BASED ON A COUNT SIZE OF 300)
1227 IF(CONST.LT.0.0001) CONST=1.0
      DO 111 I=1,N
        DO 112 J=1,M
          DATA(I,J)=ALOG10((DATA(I,J)/30)+CONST)
112 CONTINUE
111 CONTINUE
C*****
*
C TEST FOR DATA INVERSION (TTEST NEGATIVE)
825 IF(TTEST.GT.0) GO TO 830
C*****
*
C INVERT 'DATA' MATRIX FOR R-MODE CLUSTERING.
800 DO 801 I=1,N
      DO 802 J=1,M
        TMP(J,I)=DATA(I,J)
802 CONTINUE
801 CONTINUE
      DO 805 I=1,N
        DO 806 J=1,M
          DATA(J,I)=TMP(J,I)
806 CONTINUE
805 CONTINUE
      DO 810 I=1,N
        DO 811 K=1,12
          TEMCD(I,K)=ROWCD(I,K)
811 CONTINUE
810 CONTINUE
      DO 815 J=1,M
        DO 816 K=1,12
          ROWCD(J,K)=COLCD(J,K)
816 CONTINUE
815 CONTINUE
      DO 821 I=1,N
        DO 822 K=1,12
          COLCD(I,K)=TEMCD(I,K)
822 CONTINUE
821 CONTINUE
      MT=M
      M=N
      N=MT
      IPASS = IPASS + 1
      IF(MCOEFF.EQ.10) GO TO 1225
C*****
C WRITE OUT DATA MATRIX (TRANSFORMED IF REQUESTED) TO FILEDEF
UNIT 07
830 WRITE (7,100) TITLE
      IF(MCOEFF.EQ.1.OR.MCOEFF.EQ.2.OR.MCOEFF.EQ.6) WRITE(7,301)
      IF(MCOEFF.EQ.3.OR.MCOEFF.EQ.7.OR.MCOEFF.EQ.12) WRITE(7,302)
      IF(MCOEFF.EQ.4.OR.MCOEFF.EQ.8) WRITE(7,303)
      IF(MCOEFF.EQ.10.AND.ITEST.LT.0) WRITE(7,313)

```

```

IF(MCOEFF.EQ.10.AND.ITEST.GT.0) WRITE(7,302)
IF(MCOEFF.EQ.5.OR.MCOEFF.EQ.9) WRITE(7,310) CONST
IF(MCOEFF.EQ.11) WRITE(7,315)
IF(MCOEFF.EQ.13) WRITE(7,315)
301 FORMAT(' NO TRANSFORMATION')
302 FORMAT(' PERCENT TRANSFORMATION')
303 FORMAT(' PERCENT-MAXIMUM TRANSFORMATION')
313 FORMAT(' DOUBLE TRANSFORMED')
315 FORMAT(' PERCENT AND FREEMAN-TUKEY TRANSFORMED')
310 FORMAT(' LOG-10 TRANSFORMATION, CONSTANT = ',F7.3)
WRITE(7,304)
304 FORMAT('DATA IS IN A (10(F7.3,1X)) FORMAT')
WRITE (7,200)
200 FORMAT(' ')
DO 5 J=1,M
WRITE (7,100) (COLCD(J,K),K=1,12)
5 CONTINUE
DO 6 I=1,N
WRITE(7,100) (ROWCD(I,K),K=1,12)
WRITE(7,201) (DATA(I,J),J=1,M)
201 FORMAT(10(F7.3,1X))
6 CONTINUE
WRITE (7,200)
C*****
*
C TEST FOR END OF PROGRAM (USE TO CONVERT/INVERT DATA W/O
PERFORMING
C CLUSTER ANALYSIS)
READ(4,102) ITEST,NCOEFF,CONST
IF(ITEST.EQ.0.AND.NCOEFF.LT.0) GO TO 9999
IF(CONST.LT.0.0001) CONST=1.0
C*****
*
C INITIALIZE MATRICES.
DO 8 I=1,N
SUM(I)=0.
T(I)=1.
NUMBR(I)=1
DO 9 J=1,M
SIMCO(I,J)=0.
9 CONTINUE
8 CONTINUE
C*****
*
C TEST FOR SIM. COEFF. TYPE (JACCARD, CZEKANOWSKI, HQM, OR CRD)
IF(MCOEFF.EQ.1) GO TO 125
IF(MCOEFF.GE.6.AND.MCOEFF.LT.10) GO TO 126
IF(MCOEFF.GE.12) GO TO 450
C*****
*
C COMPUTE 'CZEKANOWSKI' COEFFICIENTS AND STORE IN LOWER
TRIANGULAR
C (LESS DIAGONAL) PART OF MATRIX 'SIMCO'. (MCOEFF=02.03,04.05,10.11)
DO 10 I=1,N

```



```

    DO 20 J=1,M
20  SUM(I)=SUM(I)+DATA(I,J)
    DO 30 K=1,I
    IF (I.EQ.K)GOTO 30
    W=0.
    DO 40 J=1,M
    VALMIN=AMIN1(DATA(I,J),DATA(K,J))
40  W=W+VALMIN
    SIMCO(I,K)=(2.*W)/(SUM(I)+SUM(K))
30  CONTINUE
10  CONTINUE
    GO TO 15
C*****
*
C COMPUTE CHORD DISTANCE COEFFICIENTS AND STORE IN LOWER
TRIANGULAR
C (LESS DIAGONAL) PART OF MATRIX 'SIMCO'. (MCOEFF=12)
450 DO 455 I=1,N
    DO 460 K=1,I
    IF (I.EQ.K)GOTO 460
    MULTTOT=0.
    SQTOTI=0.
    SQTOTK=0.
    DO 465 J=1,M
    MULTTOT=(DATA(I,J)*DATA(K,J))
    SQTOTI=SQTOTI+DATA(I,J)**2
    SQTOTK=SQTOTK+DATA(K,J)**2
465  CONTINUE
    CCOS=MULTTOT/SQRT(SQTOTI*SQTOTK)
    CRD=SQRT(2.*(1-CCOS))
C-- ADJUST CRD TO 0 TO 1 LOW TO HIGH SCALE
    SIMCO(I,K)=1-CRD/SQRT(2.0)
460  CONTINUE
455  CONTINUE
    GO TO 15
C*****
*
C HQM : FREQUENCY MODULATED RELATIVE HOMOGENEITY FUNCTION
(HALL 1967A)
C (MCOEFF=06,07,08,09)
126 DO 11 I=1,N
    DO 21 J=1,M
    SUM(I)=SUM(I)+DATA(I,J)
21  CONTINUE
    DO 31 K=1,I
    IF (I.EQ.K) GO TO 31
    W=0.0
    DO 41 J=1,M
    VALMIN=AMIN1(DATA(I,J),DATA(K,J))
    VALMAX=AMAX1(DATA(I,J),DATA(K,J))
    W=W+VALMIN
    V=V+VALMAX
41  CONTINUE
    SIMCO(I,K)=(W+(W*W/V))/(SUM(I)+SUM(K))

```

```

31 CONTINUE
11 CONTINUE
GO TO 15
C*****
*
C COMPUTE JACCARD COEFF (MCOEFF=01)
125 DO 95 I=1,N
    DO 94 K=1,I
    IF(I.EQ.K)GO TO 94
    PQ=0.
    PNO=0.
    DO 93 J=1,M
    IF(DATA(K,J).GT.0..AND.DATA(I,J).GT.0.)PQ=PQ+1.
    IF(DATA(K,J).GT.0..AND.DATA(I,J).EQ.0.)PNO=PNO+1.
    IF(DATA(K,J).EQ.0..AND.DATA(I,J).GT.0.)PNO=PNO+1.
93 CONTINUE
SIMCO(I,K)=PQ/(PNO+PQ)
94 CONTINUE
95 CONTINUE
C*****
*
C ARRAY 'BOOK', WHICH OCCUPIES SAME LOCATIONS IN MEMORY AS
ARRAY 'SIMCO'
C IN ORDER TO CONSERVE SPACE, WILL BE USED FOR BOOKKEEPING
PURPOSES.
C ONLY THE UPPER TRIANGULAR (INCLUDING DIAGONAL) PART OF 'BOOK'
WILL BE
C USED, SO VALUES CONTAINED IN 'SIMCO' WILL BE UNDISTURBED. EACH
COLUMN
C IN ARRAY 'BOOK' CONTAINS THE NUMBERS OF THE SAMPLES
CLUSTERED
C TOGETHER IN THE OPPOSITE COLUMN IN 'SIMCO'.
15 DO 16 I=1,N
16 BOOK(1,I)=FLOAT(N-I)+1.
    NX=N
    NPASS=1
C*****
*
C 'SMAX' WILL CONTAIN THE LARGEST CORRELATION COEFFICIENT IN THE
CURRENT
C 'SIMCO' MATRIX. THE ROW AND COLUMN NUMBER OF THAT VALUE ARE
RECORDED
C IN 'LVROW' AND 'LVCOL', RESPECTIVELY. 'SMAX' IS INITIALIZED AT THE
C BEGINNING OF THE SEARCH TO A VALUE LOWER THAN THE FIRST
CORRELATION
C COEFFICIENT ENCOUNTERED. THE SAMPLES OR CLUSTERS WITH THESE
HIGHEST
C VALUES WILL THEN BE CLUSTERED.
45 SMAX=SIMCO(2,1)-1.
DO 50 I=2,NX
    IM1=I-1
    DO 51 J=1,IM1
    IF(SMAX.GE.SIMCO(I,J)) GO TO 51
    SMAX=SIMCO(I,J)

```

```

LVROW=I
LVCOL=J
51 CONTINUE
50 CONTINUE
C*****
*
C SAMPLE (OR CLUSTER) NUMBERS ARE CLUSTERED AND THEIR
SIMILARITY
C COEFFICIENT IS RECORDED IN ARRAY 'OUT' FOR LATER PRINTOUT AND
PLOT.
COMBO=T(LVROW)+T(LVCOL)
NBCOL=N-LVCOL+1
NBROW=N-LVROW+1
OUT(NPASS,1)=BOOK(1,NBCOL)
OUT(NPASS,2)=BOOK(1,NBROW)
OUT(NPASS,3)=SMAX
C*****
*
C SIMILARITY COEFFICIENTS ARE CALCULATED FOR THE NEW CLUSTER &
COMPUTED
C AND STORED IN 'SIMCO'. 'SIMCO' IS THEN COMPACTED BY REMOVING
THE ROW
C AND COLUMN OF THE SAMPLE WITH THE LARGER NUMBER WHICH WAS
CLUSTERED.
C ARRAY 'BOOK' IS ALSO UPDATED TO REFLECT THE CHANGE DUE TO THE
NEW
C CLUSTERING.
DO 60 I=1,NX
IF(I.EQ.LVCOL) GO TO 60
IF(I.LT.LVCOL) SIMCO(LVCOL,I)=
*(T(LVCOL)*SIMCO(LVCOL,I)+T(LVROW)*SIMCO(LVROW,I))/COMBO
IF(I.GT.LVCOL.AND.I.LT.LVROW) SIMCO(I,LVCOL)=
*(T(LVCOL)*SIMCO(I,LVCOL)+T(LVROW)*SIMCO(LVROW,I))/COMBO
IF(I.GT.LVROW) SIMCO(I,LVCOL)=
*(T(LVCOL)*SIMCO(I,LVCOL)+T(LVROW)*SIMCO(I,LVROW))/COMBO
60 CONTINUE
T(LVCOL)=COMBO
NXM1=NX-1
DO 70 I=LVROW,NXM1
JADJ=0
DO 75 J=1,I
IF (J.EQ.LVROW) JADJ=1
SIMCO(I,J)=SIMCO(I+1,J+JADJ)
75 CONTINUE
70 CONTINUE
C*****
*
C ARRAY 'NUMBR' CONTAINS THE NUMBER OF SAMPLES COMPRISING
EACH CLUSTER.
NROWS=NUMBR(LVCOL)
NUM=NUMBR(LVROW)
KOLT=NBCOL
KOLF=NBROW
DO 501 I=1,NUM

```

```

NROWT=NROWS+I
BOOK(NROWT,KOLT)=BOOK(I,KOLF)
501 BOOK(I,KOLF)=0.
IF(LVROW.EQ.NX)GO TO 535
DO 505 I=LVROW,NXM1
IX=N-I+1
IXM1=IX-1
NUMX=MAX0(NUMBR(I),NUMBR(I+1))
DO 515 IC=1,NUMX
BOOK(IC,IX)=BOOK(IC,IXM1)
515 CONTINUE
505 CONTINUE
535 NUMBR(LVCOL)=NUMBR(LVCOL)+NUMBR(LVROW)
IF(LVROW.EQ.NX) GO TO 545
DO 510 I=LVROW,NXM1
510 NUMBR(I)=NUMBR(I+1)
545 NUMBR(NX)=0
DO 550 I=1,NX
550 SIMCO(NX,I)=0.
NPASS=NPASS+1
NX=NX-1
IF (NX.EQ.1) GO TO 80
GO TO 45
80 CONTINUE
NM1=N-1
C*****
*
C WRITE OUT RESULTS AS TEXT FILE TO FILEDEF UNIT 06
WRITE(6,100) TITLE
WRITE(12,100) TITLE
WRITE(12,'(I4,2X,"SAMPLES")') N
WRITE(14,100) TITLE
2001 FORMAT(I5)
IF(ITEST.EQ.1) WRITE(6,305)
305 FORMAT(' Q-MODE')
IF(ITEST.EQ.-1) WRITE(6,306)
306 FORMAT(' R-MODE')
IF(MCOEFF.EQ.1) WRITE(6,307)
307 FORMAT(' JACCARD COEFFICIENT')
IF(MCOEFF.GE.2.AND.MCOEFF.LE.5) WRITE(6,308)
308 FORMAT(' CZEKANOWSKI COEFFICIENT')
IF(MCOEFF.GE.6.AND.MCOEFF.LE.9) WRITE(6,309)
309 FORMAT(' HOMOGENEITY FUNCTION')
IF(MCOEFF.EQ.1.OR.MCOEFF.EQ.2.OR.MCOEFF.EQ.6) WRITE(6,301)
IF(MCOEFF.EQ.3.OR.MCOEFF.EQ.7) WRITE(6,302)
IF(MCOEFF.EQ.4.OR.MCOEFF.EQ.8) WRITE(6,303)
IF(MCOEFF.EQ.5.OR.MCOEFF.EQ.9) WRITE(6,310) CONST
WRITE(6,200)
WRITE(6,202)
202 FORMAT('SIM. COEFF. VALUES FOR DENDRO. BRANCHES:')
DO 90 I=1,NM1
VAL1=IFIX(OUT(I,1))
VAL2=IFIX(OUT(I,2))
WRITE(6,203) OUT(I,1),(ROWCD(VAL1,K),K=1,12),OUT(I,2),

```

```

*(ROWCD(VAL2,K),K=1,12),OUT(1,3)
203  FORMAT(F4.0,'=',12A4,5X,F4.0,'=',12A4,5X,F5.3)
90  CONTINUE
    WRITE(6,200)
    WRITE(6,204)
204  FORMAT('DENDRO. BRANCH CODES IN DENDROGRAM ORDER')
    DO 91 I=1,N
        IBOOK=IFIX(BOOK(I,N))
        WRITE(12,'(F4.0)') BOOK(I,N)
        WRITE(14,205) I,(ROWCD(IBOOK,K),K=1,12)
    C   WRITE(6,205) BOOK(I,N),(ROWCD(IBOOK,K),K=1,12)
205  FORMAT(I4,' ',12A4)
91  CONTINUE
    WRITE(6,200)
C*****
C PLOT THE DENDROGRAM. (SENDS PLOTTING INSTRUCTIONS TO FILEDEF
UNIT 10)
C 34 INCHES IS THE MAXIMUM WORKING HEIGHT FOR THIS ROUTINE ON
THE
C VERSATEC PLOTTER, WHILE THE LENGTH MAY BE CHOSEN AS DESIRED.
C THE VALUE OF "NEWPEN" DETERMINES THE PLOT LINE THICKNESS.
    CALL NEWPEN(NP)
C-----
C THIS DRAWS THE AXIS.
    DATA CSIM /4HCOEF,4HFICI,4HENT ,4HOF S,4HIMIL,4HARIT,4HY /
    CALL SAXIS(0.0,(HEIGHT+0.1),CSIM,25,LENGTH,0.0,0.1,0.1,
*(LENGTH/10),(LENGTH/10))
C-----
C THIS SCALES THE DENDROGRAM.
    NP1=N+1
    YSCALE=HEIGHT/FLOAT(N)
    XNUM=LENGTH+0.2
    XSYM=LENGTH+0.9
    CHARHT=0.2
    IF(YSCALE.LT.0.25) CHARHT=0.075
    IF(YSCALE.LT.0.25) XSYM=LENGTH+0.7
    DO 601 I=1,N
        IBOOK=IFIX(BOOK(I,N))
        Y=FLOAT(NP1-I)
        PLOCX(IBOOK)=LENGTH
        PLOCY(IBOOK)=Y*YSCALE
    C   IF(YSCALE.LT.0.15) GO TO 601
C-----
C THIS NUMBERS AND LABELS THE BRANCHES OF THE DENDROGRAM.
    YPOS=PLOCY(IBOOK)-(CHARHT/2)
    CALL NUMBER(XNUM,YPOS,CHARHT,BOOK(I,N),0.0,0)
    DO 71 K=1,12
        LABEL(K)=ROWCD(IBOOK,K)
71  CONTINUE
    CALL SYMBOL(XSYM,YPOS,CHARHT,LABEL,0.0,48)
601 CONTINUE
C-----
C THIS DRAWS THE DENDROGRAM'S BRANCHES.
    NM1=N-1

```

```

DO 610 I=1,NM1
X=LENGTH*OUT(I,3)
INDEX1=IFIX(OUT(I,1))
INDEX2=IFIX(OUT(I,2))
CALL PLOT(PLOCX(INDEX1),PLOCY(INDEX1),3)
CALL PLOT(X,PLOCY(INDEX1),2)
CALL PLOT(X,PLOCY(INDEX2),2)
CALL PLOT(PLOCX(INDEX2),PLOCY(INDEX2),2)
PLOCX(INDEX1)=X
PLOCY(INDEX1)=(PLOCY(INDEX1)+PLOCY(INDEX2))/2.
610 CONTINUE
CALL PLOT(PLOCX(INDEX1),PLOCY(INDEX1),3)
CALL PLOT(0.0,PLOCY(INDEX1),2)
C*****
C TEST CONTROL CARD VALUE FOR END OF PROGRAM OR FOR RE-
EXECUTION OF
C PROGRAM (IF REQUESTED)
IF(ITEST.EQ.0)GO TO 999
CALL PLOT(LENGTH+3.,0.,-3)
GO TO 825
C END PROGRAM
*****
999 CONTINUE
CALL PLOT(LENGTH+3.,0.,999)
RETURN
9999 END

```

## VITA: Jon C. Cawley

### EDUCATION:

1994-1996

**Virginia Polytechnic Institute and State Univ., Blacksburg, VA**  
*Masters of Science.* Defended July 29, 1996.

\*Major Advisors: R. K. Bambach / D.M. McLean / B. Parker

\*Topic: Dynamic Systems Analysis of Fossil Dinoflagellates from the Atlantic Coastal Plain, USA.

Aug. 1993

**SD School of Mines and Technology/NJ State Museum**

*Summer Field Course:* Field Paleontology and Biostratigraphy

Crow Creek (Lakota Sioux) Lands, South Dakota

\*Collected and Documented Cretaceous Vertebrates (Mosasaurs, Plesiosaurs, Turtles).

\*Conducted stratigraphic measurement and sampling of Cretaceous Pierre Shale sections.

\*Graduate level project: Palynological lab processing and investigation of fossil spores and pollen from Pierre Shale members; paper submitted.

1981-1986

**The Pennsylvania State University, State College, PA**

*Bachelor of Science*

\*Major: Earth Science (Geology) Minor: Writing

\*Earth and Mineral Sciences Interest House member

- Chairman of academic committee, 1 year

- Creator and Editor of bi-weekly house newsletter. 1983-1986

- Field trip Officer for PSU Geosciences Club.

\*PSU Museum of Anthropology, Intern (worked with European Paleolithics)

1983-1985

**Undergraduate Thesis:**

*The Bradford Oil Basin: A Regional History of Oil Technology*, 1986.

(39 thousand words)

### EXPERIENCE:

August, 1995

**Pennsylvania Registered Professional Geologist # PG-001529-G**

Sept. 1993

**NICET Certification in Engineering Technology, Level 1**

Certification number: 084120

1989-Present

**CQS Inc., Harrisburg, PA**

*Development and Technical Partner*

Develop and implement laboratory testing procedures; perform aggregate tests to PA and AASHTO specifications; develop environmental consulting programs; oversee field projects. Studies in Geomorphology and land use. Project work in limestone for sorbent purposes, field and lab projects in Alkali Silica Reactivity and use of fly ash as pozzolon.

\*Corporate Officer, November 1990; AASHTO lab accreditation, July 1991.

\*Co-author of PACA Resolution on Alkali Silica Reactivity. 1992.

\*Named in Whos Who Registry of Business Leaders, 1994/1995

1989-1994

**Temple University Geology Department, Philadelphia, PA**  
*Manager*

Organization of teaching labs and field trips; substitute lab instructor; responsibility for department equipment and vehicles; maintenance and operation of thin section lab, photography lab and darkroom; maintenance and expansion of teaching collections; design of geologic displays; liaison between vendors and department for purchasing of equipment; general support to faculty, staff and students.

\*Member of Dean's Committee on Hazardous Chemicals

- Author of CAS Guidelines on Lab Training and Practice.

\*Member of Dean's Security Task Force Committee.

1986-1989

**Tethys Consultants Inc./ Mid-Atlantic Testing, Harrisburg, PA**  
*Staff Geologist*

Performed field geology; data reduction; report preparation and editing; implemented water sampling programs; established SARA III chemical inventories for industry. Assisted and implemented drilling programs in coal waste and limestone for cogeneration and aggregate use. Co-developed and set up Mid-Atlantic aggregates testing lab; performed aggregates tests to PA and AASHTO standards. Performed carbonate measurements for limestone in cogeneration.

\*AASHTO accreditation for laboratory, April 1988 (1st in the U.S.)

#### UNIVERSITY TEACHING AND AWARDS

May, 1996

**Virginia Polytechnic Institute -**  
**1996 Tillman Excellence-in-Teaching Award.**

May, 1995

**Virginia Polytechnic Institute -**  
**1995 Tillman Excellence-in-Teaching Award.**

**Virginia Tech, Paleontology 4304** Spring Semesters 1994 and 1995.

*Lab Instructor*

(Upper-level class included invertebrate, vertebrate, and botanical Paleontology for majors.)

**Virginia Tech, Physical Geology 1104** Fall Semester 1994, Summer 1995. *Lab Instructor*

(Chosen for pilot program of interactive teaching for Physical Geology to nonmajors, Fall 1994.)

**Virginia Tech, Historical Geology 1114** Summer 94, Fall and Spring 95/96. *Lab Instructor*

(Topics included land processes, fossils, time relationships. For majors/nonmajors.)

#### ACTIVITIES:

1995-1996

**Sigma Gamma Epsilon - Earth Sciences National Honor Society.**  
Virginia Tech (Alpha Mu) Chapter. Blacksburg, VA

(Advisor for undergraduate Earth Sciences Honor Society. Placed emphasis on interdisciplinary communication between the different Earth Science departmental majors on campus.)

