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ECOSYSTEM PROCESS MODELING OF SUBMERGED AQUATIC VEGETATION IN THE LOWER CHESAPEAKE BAY

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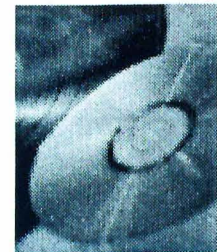
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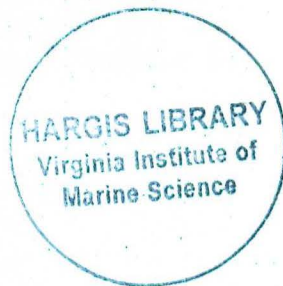
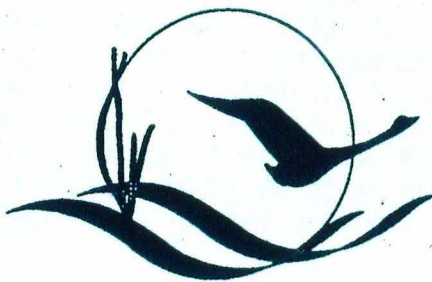


Ecosystem Process Modeling of Submerged Aquatic Vegetation in the Lower Chesapeake Bay

by

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Abstract: The modeling studies reported here emphasize the polyhaline eelgrass habitats of the lower Chesapeake Bay, as a part of a larger program of ecosystem modeling sponsored by the Living Resources Subcommittee of the Chesapeake Bay Program. The report covers progress made through April 1993. The primary focus of the polyhaline SAV modeling studies has been on light-dependent eelgrass productivity and water quality parameters which affect the submarine light climate. Where eutrophication impacts have been assessed, they have been used to drive increases in epiphytes, which block light penetration to eelgrass leaves. Eutrophication of the tidal tributaries diminishes light-dependent eelgrass production and growth and has been implicated as the principal cause for SAV declines throughout the Chesapeake Bay.



EXECUTIVE SUMMARY

The modeling studies reported here emphasize the polyhaline eelgrass habitats of the lower Chesapeake Bay, as a part of a larger program of ecosystem modeling sponsored by the Living Resources Subcommittee of the Chesapeake Bay Program. This report covers progress made through April 1993. The primary focus of the polyhaline SAV modeling studies has been on light-dependent eelgrass productivity and water quality parameters which affect the submarine light climate. Where eutrophication impacts have been assessed, they have been used to drive increases in epiphytes, which block light penetration to eelgrass leaves. Eutrophication of the tidal tributaries diminishes light-dependent eelgrass production and growth and has been implicated as the principal cause for SAV declines throughout the Chesapeake Bay. The implicit assumptions have been that eelgrass is not nutrient-limited in its natural setting and that increased nutrient loadings to the estuary negatively affect SAV growth and survival by promoting the growth of planktonic microalgae and epiphytes.

The successes of eelgrass modeling to date are illustrated by three major findings. First, the current model version yields productivities and plant survival characteristics, both short and long term, which are very similar to those observed in process-oriented studies and in analyses of long term SAV decline within the Chesapeake Bay. The modeling results suggest that the principal environmental variable controlling eelgrass growth and survival in the lower Chesapeake Bay is the availability of light, specifically photosynthetically active radiation. For the water column, suspended inorganic solids and chlorophyll concentration determine the intensity of light reaching the plant canopy but not necessarily the amount available for plant photosynthesis which further depends on epiphytic loads.

Secondly, these modeling studies have demonstrated the importance of epiphytic grazers in controlling epiphyte density on eelgrass leaves. Epiphytes reduce plant photosynthesis by attenuating light and interfering with gas (CO_2 and O_2) exchange at the leaf surface. The absence of or the reduction in the population density of grazers, as shown in modeling sensitivity studies and confirmed in mesocosm studies, can hasten the decline of eelgrass under nutrient loadings typical of present conditions in areas of the lower Chesapeake Bay.

The third finding, particularly suited to elucidation by model simulation analysis, is that variability in the submarine light climate, determined by variable solar irradiance and attenuation coefficients, at daily frequencies can have critical consequences with regard to SAV survival. Model scenarios where light parameters are specified as smoothly varying functions of monthly averages produced results that differ significantly from model scenarios where the light parameters were treated as stochastic variables derived from in situ data; the latter produced results that better reproduced observed conditions. This finding has significance for setting target water quality and habitat restoration criteria, and for deciding the appropriate means of simulating environmental conditions experienced by SAV's in continuing modeling efforts.

Future SAV modeling and simulation efforts will expand along three principal lines. First, the formulation of the light portion of the model will be revised to reflect a more theoretical basis rather than the site specific, data-driven empirical formulations used in the current versions. This will allow better integration with other efforts including the revised 3D hydrodynamic-water quality

model as well as better reflect SAV habitat criteria evaluation of restoration goals. Second, the SAV model will be expanded to include nutrient cycles, sediment oxygen and nutrient fluxes, and water column trophic dynamics. This will form the basis of a generic SAV-Littoral Zone model that can be parameterized for different regions of the Chesapeake Bay and its tributaries. Third, the SAV-Littoral Zone model will be coupled with the water quality and hydrodynamic modeling efforts in tributaries of the lower Bay. This will improve the ability to predict the transport of dissolved and particulate materials important for water quality and better determine the role of littoral zone physical and biological processes in the maintenance of water quality.

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A modeling project of this nature depends on the input from a variety of sources and more often than not the sharing of data often before the originator of those data are able to publish. We particularly wish to thank Robert J. Orth, Kenneth A. Moore, and Hilary A. Neckles for their generous aid in this regard. To our modeling colleagues in Maryland who share in this continuing program, Michael Kemp, Chris Madden and Walter Boynton, thanks for the many ways you have contributed to this effort. We also thank our graduate students who have been directly involved in these studies, William Seufzer and David Fugate.

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INTRODUCTION

For the past several decades and in particular since the mid-1970s, the Chesapeake Bay and its tributaries have been the focus of intense study and directed research to understand the causes and consequences of reduced water quality and the loss of living resources. A long-term goal of these varied research efforts has been to provide the causes of and propose the means to reverse trends in environmental degradation. Research and management programs supported by regional, state, and federal initiatives have included efforts ranging from the large-scale, three-dimensional hydrodynamic and water quality modeling to studies focused on specific habitats (e.g., shallow waters dominated by submerged aquatic vegetation) and populations of economic and/or recreational importance (e.g., oysters, blue crabs and striped bass). Water quality issues have been one general focus, examining the phenomena of low dissolved oxygen, nutrient loadings, and upland land-use practices and management. Living resource issues are another area of research, addressing particular habitats and specific populations, often out of the context of the larger system of which they are a part. This leaves a complex and yet unresolved question:

How and to what extent can the results of research on water quality and living resources, often viewed independently from one another, be integrated to arrive at sound ecological management strategies of the estuary, its tributaries, and multiple resources?

To address this question in part, Wetzel and Hopkinson (1991) suggested that ecosystem modeling and simulation analysis was an appropriate and useful tool for the Chesapeake Bay Program. Ecosystem modeling and simulation analysis has progressed rapidly over the past two decades. Advances in computer technology have played a leading role in this development and at the same time the modeler and model end-user have matured with regard to methods and expectations. While modeling techniques and the use of mathematical models are not new to marine science or environmental management, no concerted effort has been given to integrating the various methods used in the physical, chemical, and biological sciences for studies of the Chesapeake Bay and its tributaries. An integration that is considered by many as both necessary and appropriate to deal with the high degree of complexity of natural systems and the environmental management of large scale systems.

The Living Resources subcommittee of the U.S. E.P.A.'s Chesapeake Bay Program initiated in 1991 an ecosystem modeling program to develop an integrative modeling framework responsive to management needs particularly as they related to living resources of the Chesapeake Bay. The program is a cooperative, integrated modeling effort among the College of William & Mary's Virginia Institute of Marine Science and the University of Maryland laboratories at Solomons (CBL) and Horn Point (HPEL). The program involves three general modeling approaches: 1) Ecological Regression Models, which employ statistical models to establish correlative measures between variables such as nutrient and suspended loads and readily observed environmental variables such as phytoplankton biomass (chlorophyll a) and dissolved oxygen concentration, 2) Ecosystem Process Models, which develop mathematical descriptions of ecological processes and simulate the dynamics in time and space of populations, habitats or entire ecosystems, and 3) Fish Bioenergetics Models, which relate finfish physiological and behavioral responses to food abundance, temperature, and dissolved oxygen. Conceptually, these approaches can be linked through the sharing of common

variables and forcing functions. Analyzed as standalone models, they provide insight and some degree of predictive capability in establishing management criteria and evaluating alternative management scenarios. Together, they provide a powerful tool for analysis of living marine resources in the context of environmental variability and quality.

Within the programmatic goals of ecosystem modeling is the conceptualization, implementation and analysis of three ecosystem process models for spatially and ecologically distinct regions or habitats of the estuary; 1) submerged aquatic vegetation, 2) emergent intertidal marshes and 3) the water column-benthos. While these are somewhat arbitrary distinctions, some environmental management concerns are unique to these areas (e.g. the development of habitat criteria and restoration goals) but as importantly they represent a logical ecological division for model development and validation. Also within the programmatic framework, is the coupling of these models via common forcing functions, environmental water variables and shared state variables with both ecological regression models and fish bioenergetics models.

This report presents results from ecosystem process models of the polyhaline littoral zone habitat dominated by eelgrass (*Zostera marina*) in the lower portions of the Chesapeake Bay. The emphasis of these modeling studies has been on the role of environmental factors, particularly light, in governing the distribution, growth and survival of eelgrass.

BACKGROUND

Ecological Modeling of Complex Systems

Ecosystem models are abstractions or simplifications of complex natural systems and as such are useful tools for scientific analysis; the multitude of interactions is too overwhelming to be perceived intuitively, but may be penetrable within a quantitative simulation framework. Deterministic, numerical simulation models of ecosystems synthesize large amounts of information on individual parts of systems, integrate these data based on a conceptual structure and explore, via simulation analysis, compartmental behavior under various simulated operating conditions. Models are now commonly used to plan and guide research, to identify data weaknesses and gaps, to evaluate management-oriented alternatives, and to provide the basis to formulate hypotheses regarding a system's structure and function. Wetzel and Hopkinson (1991) have recently reviewed ecosystem models of coastal systems with reference specifically to their application to the Chesapeake Bay region.

Models can be classified or characterized according to a number of schemes. Two general categories into which ecological models can be placed are 1) Explanatory Models and 2) Correlative Models (Gold 1977; Gilchrist 1984). Explanatory models attempt to examine causal relations between components of the modeled system based on a conceptual structure or hypothesis and explain cause-effect relationships. Correlative models, on the other hand, statistically describe the relationships between a dependent variable and one or more independent variables using techniques of regression and correlation analysis. These models have no explanatory power and cannot extrapolate relationships beyond the conditions under which they were developed. They can, however, be used to suggest probable causal relationships that can be explored using other methods.

Often, aquatic ecosystem models combine features of both model types, incorporating statistical inferences where theoretical constructs are poorly developed or process-oriented data are lacking.

At the Virginia Institute of Marine Science (VIMS), modeling has been used in conjunction with interdisciplinary research conducted within the Chesapeake Bay and its tributaries. Field and laboratory studies provide necessary information on processes and distributions. Modeling provides an ecosystem context into which these results can be placed and allow for experimentation regarding the sensitivity of an ecosystem to chronic or traumatic perturbations. Poorly understood processes or quantities uncovered by modeling then feed back into a renewed series of observation and experimentation. Models then prove very useful for examining environmental, biological, and ecological relationships and for indicating productive avenues of research (e.g., Wetzel and Neckles 1986). A case study of this process is our research efforts in a tributary of the Chesapeake Bay which has experienced comparatively low levels of anthropogenic impact, the York River.

The York River Regional Ecosystem

In 1991, the Institute (VIMS) and the School of Marine Science, College of William and Mary embarked upon a multidisciplinary program designed to address large scale, complex coastal ecosystems and contemporary management issues. The intent of the program is to provide a professional infrastructure for the integration of knowledge from various marine science disciplines relevant to coastal zone management and the acquisition of basic knowledge necessary for understanding coastal ecosystem dynamics within a large-scale framework. The focus initially is on the York River estuary and its watershed: the York River Regional Ecosystem. The primary organizational and systems analysis tools employed will be the development of ecosystem models coupled with water quality, hydrodynamic and fisheries models. The undertaking is viewed as a long-term effort and commitment of institutional resources augmented by grants and contracts from state and federal agencies, local industry, and the private sector. A brief description of the York River system follows.

Physical-Chemical Characteristics

The York River is a subestuary of the Chesapeake Bay and is formed by the confluence of the Mattaponi and Pamunkey Rivers at West Point, Virginia. Its drainage basin is 69,000 km², of which approximately 70% is forested, 22% is in crop land and pasture, and <2% is classified as urban (Bender 1987). Gloucester County which makes up much of the north shore of the York is one of the most rapidly developing Tidewater counties, experiencing a population increase of 49.3% for the 1980-90 census period. The south shore is somewhat protected for the present from large scale growth and development due to the large military land holdings, the federal Colonial Historic National Park and the York River State park.

Total average freshwater inflow to the river is estimated at 70 m³ sec⁻¹. Average annual rainfall over the watershed is 112 cm which peaks in August but runoff is low due to transpiration and evaporation. The climate is humid temperate with an annual mean temperature of 15°C.

The salinity structure of the York River is influenced mainly by the interaction of freshwater, salt water and tidal energy and to a lesser extent by other forcings (e.g., wind). Periods of stratification/destratification cycle in the York River as a result of energy associated with the spring-

neap tidal cycles. Salinity gradients between the surface and bottom waters tend to be stronger during periods of neap tides and to disappear during spring tide periods. During low flow periods, salt water intrudes 20 to 30 km upriver from West Point. At the mouth, periods of bottom water hypoxia or anoxia generally coincide with stratification events during the summer. The extent to which hypoxia or anoxia in bottom waters of the York River are changing in either duration or spatial coverage is not known. The data for examining this question is not extensive enough nor for sampling reasons designed to address it in terms of an historical data base.

Nutrient concentrations and distributions in the York River show longitudinal and vertical patterns typical of temperate, coastal plain estuaries. Generally, there is a longitudinal gradient in nutrient concentrations that fluctuates in magnitude on a seasonal basis. Concentrations increase upriver reflecting watershed sources and biological processes that modulate in situ concentrations. Sources of nutrients to the York River estuary include inputs at the fall-line, point sources at West Point at the head of the estuary, surface runoff, ground-water inputs, and precipitation.

Overall, nutrient and chlorophyll concentrations in the York River are not indicative of a highly eutrophic system while other indicators suggest some degree of enrichment (e.g., loss of SAV from historically vegetated areas). Based on data available through the late 1970's, Heinle et al. (1980) characterized the York as a 'moderately eutrophic' estuary. Because projected growth in the area (primarily along the northern shoreline and upper watershed) is high for the next decade or longer and there will be concomitant changes in land use (conversion of forested and agricultural lands to rural housing developments and perhaps industry), the likelihood that anthropogenic impacts will increase without proper management is great.

Biological-Ecological Characteristics

Biologically, the York River Regional Ecosystem supports a vast array of species populations and community types that range from tidal freshwater to estuarine-marine dependent. Primary production in the York River Regional Ecosystem is determined by several sources. Emergent wetlands, dominated by various freshwater and salt-tolerant species, are an important component of the estuary and form extensive marshes in the Pamunkey and Mattaponi Rivers, smaller tributaries and creeks, and at the mouth of the York River. Submerged aquatic vegetation, once a dominant autotrophic component in the lower reaches of the York, are now confined to the lower eight to ten kilometers and are dominated by eelgrass (*Zostera marina*). Non-vegetated sediments in the shoal areas (≤ 2 m deep) make up a significant benthic habitat (ca. 38% of the York River surface area) and contribute to primary production by supporting an active autotrophic microflora dominated by benthic diatoms. Phytoplankton, because of the large euphotic water mass, contribute the greatest proportion of primary production to the York River Regional Ecosystem system and are dominated by larger forms (i.e., diatoms) during the spring bloom (January to May) and by smaller forms (nanoplankton and picoplankton) during the summer. Primary production from these sources and allochthonous organic inputs from the watershed support a complex trophic network that includes organisms ranging in size from bacteria to large finfish.

Integrative Modeling and Synthesis Program

The Approach

The York River Regional Ecosystem is conceptualized as being composed of three, interacting large-scale components: uplands, wetlands, and aquatic systems (Figure 1). The interaction between these components is governed primarily by larger-scale hydrologic, meteorologic and anthropogenic natural forcings and human perturbations. Within each of these components are smaller-scale units defined or identified by their ecological/biological structure and organized by the flow of energy, the cycling of essential elements, and the controls imposed by physical, chemical and biological interactions including those imposed by human activity. It is at this fundamental level of ecological organization that our modeling efforts are organized. The development of a single, large-scale, fully-integrated model of the York River Regional Ecosystem, though a goal to work toward, is beyond the scope of the current efforts. The development and implementation of the models proposed here should be viewed as the first necessary steps toward building integrated, large-scale ecosystem models that truly integrate a systems physics, chemistry and biology over both space and time.

The focus of these first efforts has been on the development, validation and simulation analysis of ecosystem process models for specific components of the York River Regional Ecosystem. We have proposed to develop over the first several years conceptual and simulation models of four principal components of the York River Regional Ecosystem: 1) Submerged Aquatic Vegetation (SAV), 2) Emergent Intertidal Marshes, 3) Water Column-Benthos, and, 4) Fisheries. Figure 2 illustrates the general connectivity of the proposed component models and the dominant factors influencing or controlling their interaction.

The models will be time-dependent and spatially-averaged for five geographic regions of the York River Regional Ecosystem that are defined by salinity regime for the wetlands and aquatic components. These are 1) uplands, 2) tidal freshwater (<0.5 ppt salinity), 3) oligohaline (0.5-5 ppt), 4) mesohaline (5-18 ppt), and 5) polyhaline (>18 ppt) or lower estuary/river mouth regions. Each of these geographic regions can be treated as a hydrodynamic unit characterized by one or more of the component models. This segmentation scheme has several advantages in that it

- corresponds to the principal hydrodynamic and physical-chemical regimes of the York River Regional Ecosystem,
- can be adapted to water quality and hydrodynamic modeling efforts,
- includes four sites of the National Estuarine Research Reserve System which assures long term data acquisition, and,
- follows in general a segmentation scheme proposed for similar modeling studies of the Patuxent River watershed in Maryland.

Principal Components of York River Regional Ecosystem

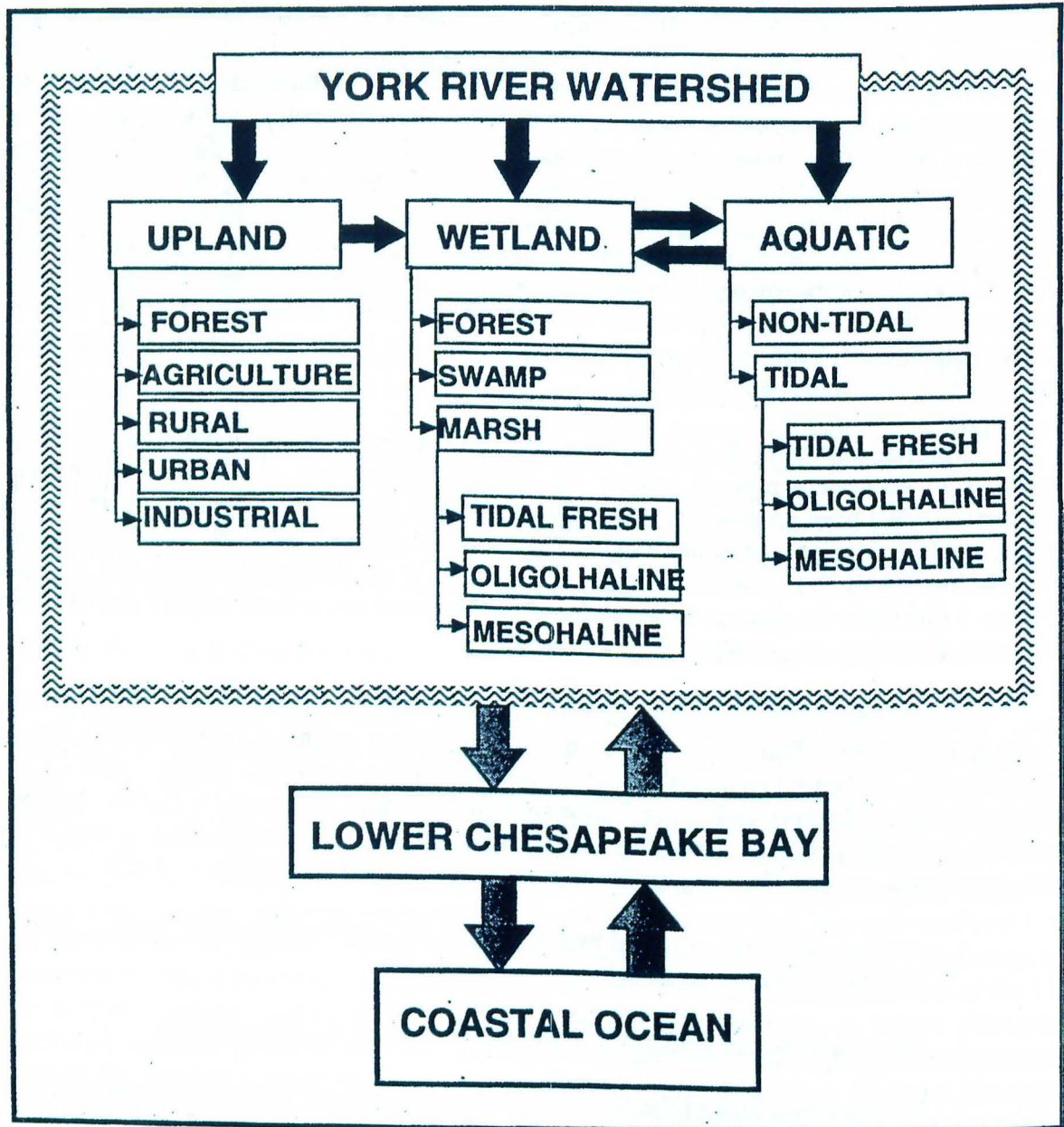


Figure 1. Principal landscape components and habitats comprising the York River Regional Ecosystem. These components form the basis for development of specific ecosystem process models. The ultimate goal is to link the various models in time and space forming a complex landscape model for the watershed.

Conceptual Ecosystem Process Model Interactions

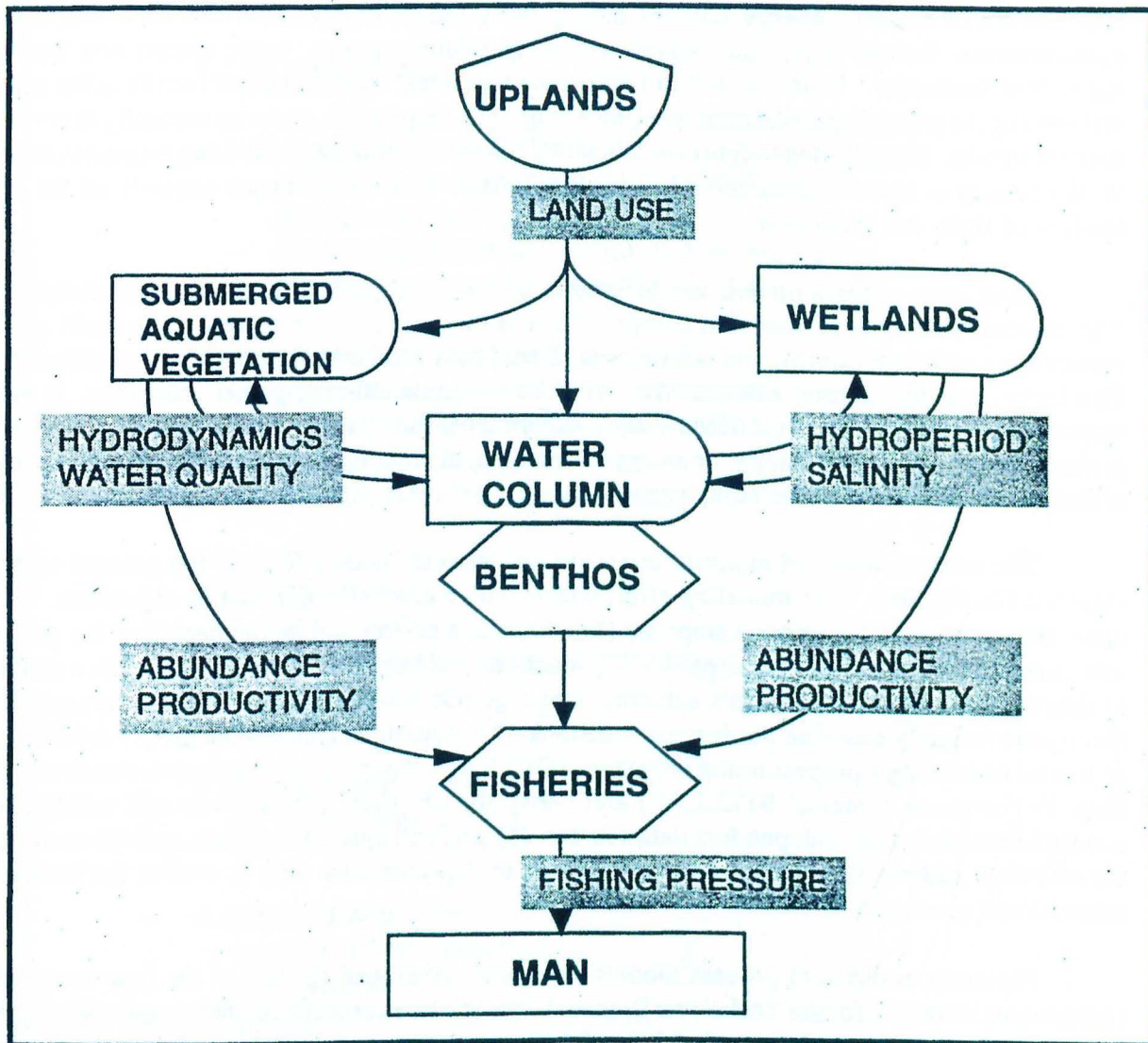


Figure 2. A conceptual diagram illustrating general relationships and couplings between the four principal ecosystem process models currently under development for the Ecosystem Process Modeling Program. The diagram illustrates the importance of land use, physical-hydrodynamic processes and water quality in governing the behavior of Wetlands, Submerged Aquatic Vegetation and the Water Column-Benthos components and their influence relative to abundance and productivity on Fisheries and ultimately human use.

Ecosystem Process Models

Ecosystem process models focus on energy and mass transfers among the components of an ecological system and those physical, chemical and biological variables considered to control these transfers. The state variables are the masses, or stocks, of the modeled components and can represent nutrient stocks, population abundance, or the aggregate density of an entire trophic level. The control of mass or energy transfer among these model components can be a function of environmental factors (e.g., solar irradiance, temperature, salinity, water depth) and density-dependent feedbacks. With respect to biotic state variables, environmental factors often act by influencing the physiologic rate processes, such as growth, respiration, and enzymatically-controlled nutrient uptake. Density-dependent controls act by imposing limits on, or altering processes related to, the density or concentration of organisms or biomass. Examples of such controls are the self-shading of light, and predation.

Ecosystem process models can be used to assess the impacts of organizational complexity (the number and kinds of modeled components and processes), the temporal cycles and spatial variabilities with ecosystems, and the impacts of environmental and anthropogenic perturbations. This latter capability is most often used to address environmental management concerns. In these respects, models form the basis of simulation experiments that are impractical or too expensive to perform in nature. Their veracity, however, is limited to the quality of observation and theory used in their formulation and to the complexity of the processes that they attempt to simulate.

The construction and analysis of ecosystem process models follows the general scheme illustrated in Figure 3. The modeling effort begins with a clearly defined set of objectives. Once these are established, the general steps are (1) construct a conceptual model depicting the general structure and identifying the principal forcing functions and controls, often depicted with a cartoon or diagrammatically using Odum's symbolic language (Odum 1971), (2) formulate the equations that mathematically describe the forcing functions, flux equations and feedbacks, (3) develop the computer code using a programming language (FORTRAN, Pascal, C, etc.) or simulation tool (e.g., High Performance Systems' STELLA[®]) and verify the program, (4) calibrate and validate the simulation model using independent data sources, (5) perform systems analysis, and (6) re-design the model to address the results of model studies, incorporate new data or evolve the model to address new questions.

Presently, ecosystem process models have been developed for two of the four ecosystem components identified for the York River Regional Ecosystem: submerged aquatic vegetation (SAV) and wetland salt marshes characteristic of polyhaline and mesohaline regions of the estuary. For the wetland ecosystem process models (*Spartina alterniflora* marshes), a conceptual model for marsh grass productivity and growth, and distribution relative to specific physical and chemical factors has been developed. For the SAV model, models have been developed, tested and completed simulation analyses with two versions.

Here, the results of SAV model studies are presented as they relate to water quality and the restoration goals set for Chesapeake Bay SAV using other methods (Batiuk et al. 1992; Dennison et al. 1993).

A Generalized Ecosystem Process Modeling Scheme

- **1. LIST OBJECTIVES**
- **2. MODEL CONCEPTUALIZATION**
 - DECIDE COMPARTMENTALIZATION SCHEME (STATE VARIABLES)
 - DETERMINE EXTERNAL FORCING FUNCTIONS
 - DEVELOP INTERACTION MATRIX
- **3. DERIVE MATHEMATICAL STRUCTURE(S)**
 - STATISTICAL - EMPIRICAL RELATIONSHIPS
 - THEORETICAL - A PRIORI CONSTRUCTS
 - FEEDBACK - INFORMATION FLOWS
 - "FORCED" - DATA LOOK-UP TABLES
- **4. PROGRAMMING**
 - DEVELOP CODE
 - APPLY MODELING PACKAGES (*STELLA, MATLAB, ETC*)
 - VERIFICATION
- **5. DIGITAL SIMULATION**
 - VALIDATION
- **6. SYSTEMS ANALYSIS**
 - SENSITIVITY ANALYSIS
 - STABILITY
- **7. RE-DESIGN**

Figure 3. The generalized scheme followed in development of ecosystem process models of the principal components of the Chesapeake Bay and its tributaries. Following this scheme leads to scenario runs that are designed to address management concerns and provide insight to the best management options.

CHESAPEAKE BAY SUBMERGED AQUATIC VEGETATION

Submerged aquatic vegetation (SAV) in Chesapeake Bay and its tributaries has been a focal point of basic research, resource management concerns, and public interest and growing awareness since the late 1970s. Starting in the early 1970s and continuing into the 1980s, SAV began a chronic, bay-wide decline in distribution and abundance (Orth and Moore 1983). In a previous, much-publicized decline during the late 1930s, which was pandemic for the North Atlantic, only eelgrass was greatly affected (Cottam 1935a; 1935b; Cottam and Munro 1954). By contrast, this recent loss of SAV has involved multiple species and appears to be local to the Chesapeake Bay where freshwater, mesohaline, and marine species have been adversely affected, suggesting a bay-wide deterioration in water quality. Meanwhile, other areas of the U.S. East Coast and western Europe had not appeared to be losing SAV during this time period. In the Chesapeake Bay, a "down-river, down-Bay" pattern of loss was demonstrated early in the Chesapeake Bay Program further supporting the hypothesis that changes in water quality were related to SAV decline. At that time, however, this hypothesis could not be supported scientifically because of a paucity of relevant data.

In 1980, the first concerted efforts were begun to scientifically investigate Chesapeake Bay SAV with the ultimate goals of understanding the cause(s) of the declines and the management changes necessary to conserve and restore SAV in the Chesapeake and its tributaries. Over of the next decade, SAV were studied at a variety of scales ranging from physiological to community-level ecological investigations. From the outset, ecosystem process modeling has played a significant role in the research. The results of this long-term SAV modeling effort are presented here and the findings compared to the recently published technical synthesis document on SAV habitat requirements and restoration targets (Batiuk et al. 1992).

SAV Modeling Goals and Objectives

The SAV modeling program goals and objectives fall into two areas: programmatic and specific. For the programmatic area, there are several points which, in general, can be applied to all modeling efforts. They are:

- to provide a conceptual framework in support of SAV research;
- to identify data and specific research needs for better understanding of the controls on, and dynamics of, natural communities;
- to test using simulation analysis the stability characteristics of hypothetical model structures and relationships; and
- to aid in generating alternative hypotheses that better explain the behavior of natural and perturbed systems.

The goals and objectives specific to the SAV modeling program focus on those environmental variables related to water quality that have been shown to be primary factors in

controlling SAV distribution and abundance, depth distribution and primary production. For the studies reported here, they are

- to simulate SAV growth as a function of in situ light, water depth, and temperature as primary factors controlling photosynthesis and growth,
- to evaluate using simulation experiments physical-chemical controls and epiphyte-grazer interactions on long-term SAV survival and stability, and,
- to evaluate using simulation experiments the effects of variable physical-chemical regimes characteristic of in situ conditions on long-term SAV survival and stability.

SAV Management Needs

The 1992 amendments to the Chesapeake Bay Agreement emphasize the need to restore and enhance living resources and their habitats. Toward this goal, links must be forged between resource management, habitat restoration, and pollution reduction and prevention, in part by connecting, scientifically and programmatically, resource management with habitat restoration priorities. Simulation modeling is an important tool with which such linkages can be made. Access to an ecologically-based simulation framework can give Bay Program managers a strong foundation on which to base decisions and establish priorities in the years ahead. The initial support for ecosystem modeling (funded in summer of 1991) represents a reaffirmation of interest by the Bay management community in using these tools to investigate the impact of human perturbations on the production of living resources.

The Ecosystem Process Modeling program is organized around five classes of models: 1) Plankton-Benthos, 2) Littoral Zone/Submerged Aquatic Vegetation, 3) Emergent Intertidal Wetlands, 4) Fish Bioenergetics, and 5) Ecological Regression/Visualization models. This report focuses on submerged aquatic vegetation modeling as a separate component; the future of this modeling effort, however, is the coupling of SAV with planktonic and benthic components in the distinctive littoral zone (shallow, well-mixed waters) environment.

A major goal of the Ecosystem Modeling Program is to link model simulation experiments with definable management endpoints. For instance, modeling studies will address quantitative living resource restoration goals and/or habitat requirements and restoration targets that are currently being developed under the direction and leadership of the Chesapeake Bay Program's Living Resources Subcommittee. In the SAV model presented here, particular attention has been paid to the relationship between environmental variability and biological interactions with the established restoration goals for water clarity. The modeling studies have also addressed possibilities for achieving the Tier II and III targets for SAV restoration with the lower, polyhaline portion of Chesapeake Bay given these water quality standards.

SAV Conceptual Models

Two conceptual and simulation models have been built for SAV communities characteristic of the lower Chesapeake Bay higher salinity habitats. These SAV communities are generally co-dominated by eelgrass, *Zostera marina*, and widgeongrass, *Ruppia maritima*. Eelgrass dominates

deeper areas to a maximum depth of approximately two meters while widgeongrass occupies the shallow, near-shore areas. For the results presented here, the information used in constructing and validating the models reflect available data on the structural and functional ecology of eelgrass-dominated communities.

The first conceptual and simulation model was an ecosystem model designed primarily as a means to summarize available data on SAV productivity, to elucidate sensitive processes where information was lacking, and to help guide a field research effort. The model, illustrated in Figure 4, follows the flow of carbon and includes the major trophic groups characteristic of lower bay eelgrass communities, including two higher trophic level compartments representing blue crabs and large predators such as blue fish, trout and sandbar sharks. Extensive simulation analyses were not performed using this model due to the great uncertainty in many of the parameter estimates used for simulation. Sensitivity analysis with the model, however, clearly indicated the dependency of many compartments on the dynamics of the SAV component, whose functioning was greatly simplified in the model. This led to the second generation conceptual and simulation SAV models designed to better represent SAV growth and environmental controls.

Concurrent with these first modeling efforts, field and laboratory studies on Chesapeake Bay SAV were beginning to identify the principal controls on SAV growth, the response of SAV to various physical, chemical, and biological interactions, and the development of a data base sufficient for model simulation and validation, data collected independent of model parameterization. During the 1980s, sufficient data were collected from monitoring programs, field studies, and laboratory experiments on SAV such that ecosystem process models capable of addressing estuarine water quality issues and the distribution, abundance, and stability of SAV could be developed and tested.

A second conceptual model was developed which focused on eelgrass growth in relation to major physical, chemical, and biological factors, most of which operate by modifying the amount of light available for eelgrass photosynthesis. The conceptual model depicts the factors which control submarine light intensity, as photosynthetically active radiation (PAR), reaching the eelgrass canopy. Three PAR intensities are considered in this model:

- 1) PAR_D: the daily integrated solar irradiance at the water surface determined by atmospheric conditions, latitude, time-of-year, and time-of-day;
- 2) PAR_z: the instantaneous PAR at a given water depth determined by attenuation within the water column which, in turn, is primarily a function of water depth and the suspended particle concentration (both organic and inorganic); and,
- 3) PAR_{vp}: the instantaneous PAR reaching the SAV leaf surface; i.e., PAR_z that is further attenuated by epiphytic fouling on the leaf surface and self-shading by the eelgrass canopy.

Although not explicitly modeled, dissolved inorganic nutrients, both nitrogen and phosphorus, impact the relationship of light, photosynthesis, and carbon flow by stimulating the growth of phytoplankton and epiphytes, reducing PAR available at SAV leaf surfaces, and shifting the balance of inorganic carbon fixation from angiosperm plants (i.e., SAV) to algae. Grazers, in part, may ameliorate this eutrophication effect by cropping both phytoplankton (not included in this

Conceptual Polyhaline SAV Trophic Model

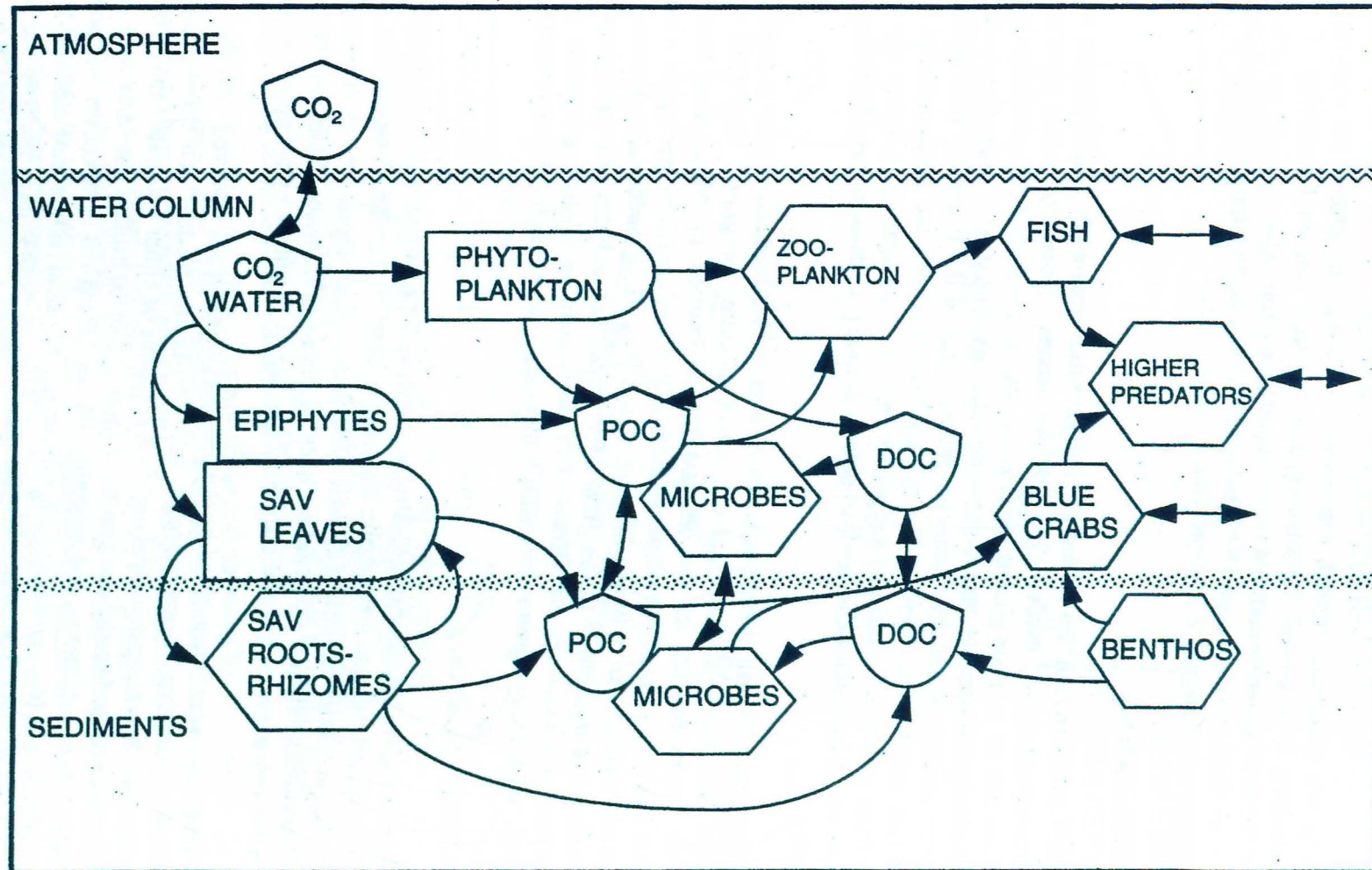


Figure 4. A conceptual model of submerged aquatic vegetation habitat in Chesapeake Bay, illustrating the major primary producers, secondary and higher consumers, and detrital pathways. POC = detrital particulate organic carbon; DOC = dissolved organic carbon.

conceptual model) and epiphyte cover on leaf surfaces, thereby increasing PAR available for SAV photosynthesis and growth.

Using this conceptual model, simulation models were developed for analysis of specific factors controlling SAV growth, distribution, and long-term community stability. For the latest versions of the SAV model (version 4 results reported in a following section), particular attention was given to evaluating the proposed SAV habitat light attenuation requirements for restoration that were developed using empirical methods (Batiuk et al. 1992).

SAV Simulation Models

Since 1984, we have developed and analyzed four simulation models that are based on the aforementioned conceptual model. The first two versions addressed various aspects of physio-chemical controls on eelgrass photosynthesis, the effects and general role of epiphytic grazing on plant productivity and survival, and the long-term stability of eelgrass communities under simulated characteristic environmental conditions for various areas of the lower bay (van Montfrans et al. 1984; Wetzel and Neckles 1986). The site-specific data used for input included information collected from sites that historically supported SAV but no longer did and sites that had healthy populations which had been stable for long periods of time (i.e., decades), as determined from aerial photography.

The third and fourth model versions have been implemented since 1991 as part of the current Chesapeake Bay Program sponsored ecosystem modeling program. The third version is a STELLA® implementation of the eelgrass model reported in Wetzel and Neckles (1986). STELLA® is a commercially available simulation tool well-suited for ecological modeling using time-dependent systems of ordinary differential equations (High Performance Systems 1992). The fourth version is a revision of the original SAV model that includes new environmental data for SAV habitats and addresses the effects of in situ variability relative to specific water quality parameters. With the exception of version 3, all models were programmed in FORTRAN 77.

Eelgrass Simulation Model

Figure 5 gives the compartmentalization scheme and flow structure for the simulation models (versions 1-4) using Odum's symbolic language (Odum 1971). The flows shown as solid lines represent linear, donor-controlled fluxes. The dashed lines represent non-linear, donor and/or recipient controlled fluxes. The dotted lines represent information flows that control negative feedbacks which operate on specific fluxes (indicated on the flows by the open arrow symbol).

The flows are characterized by the principal components involved. All abiotic-biotic and biotic-biotic interactions were modeled with non-linear, feedback-controlled functions. The feedbacks were derived as density-dependent functions of either substrate (donor) concentrations or recipient determined spatial constraints. Linear, donor-controlled functions were used to model processes such as respiration and mortality for the biological components and some physical-chemical exchanges (e.g., air-water CO₂ exchange). Physical factors were modeled with empirical or statistically derived functions. Wetzel and Wiegert (1983) have outlined the approach and general techniques followed here. Similar applications have been reported more recently by Christian and Wetzel (1991).

Compartmental Polyhaline SAV Model

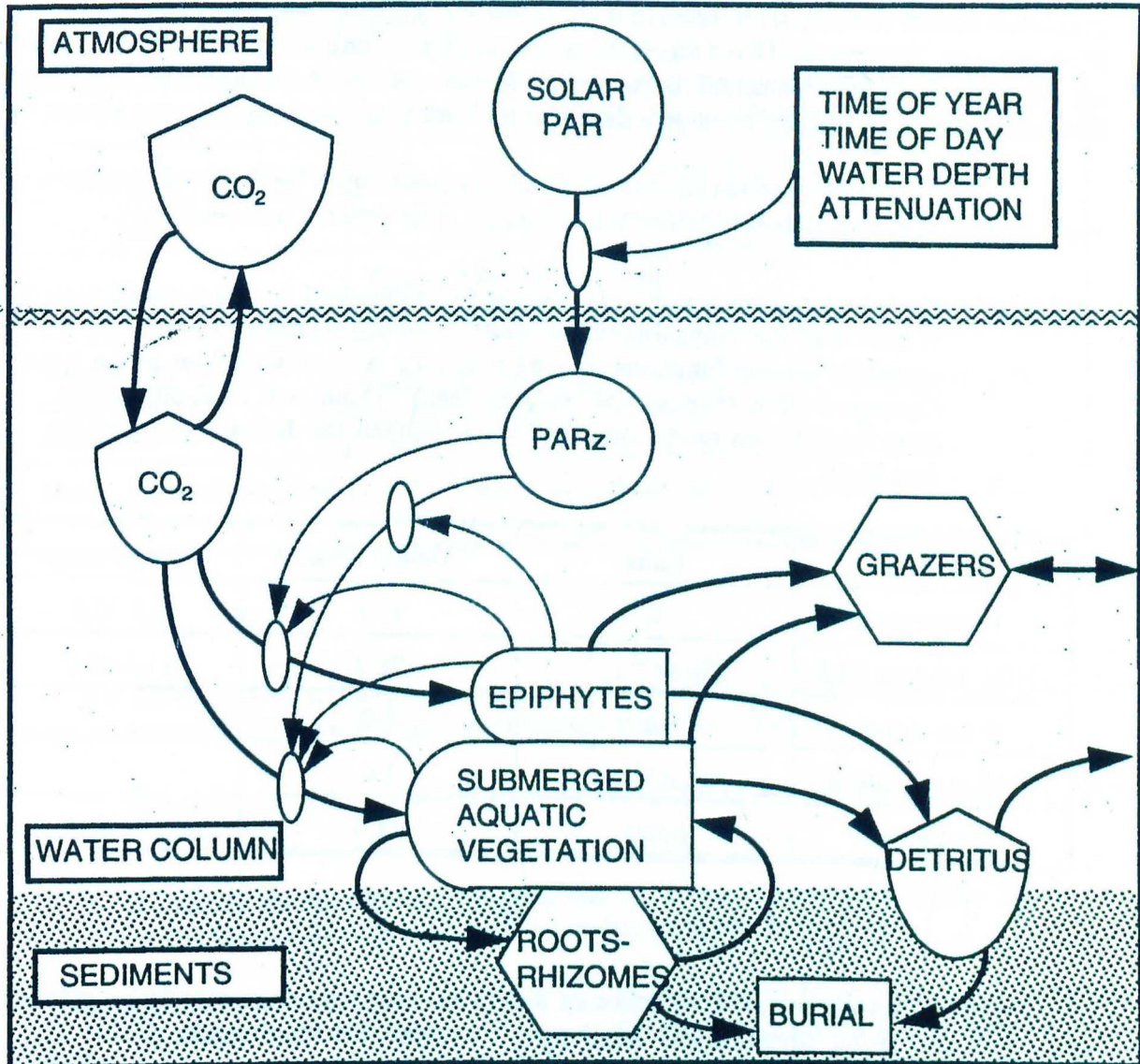


Figure 5. Compartmental and flow structure of the carbon-based SAV model for the lower, polyhaline Chesapeake Bay. The modeled SAV population is eelgrass (*Zostera marina*), which is divided into aboveground leaves and belowground roots-rhizome components. Epiphytic coverage attenuates light and interferes with gas exchange at leaf surfaces. Incident light follows diurnal and seasonal cycles and is attenuated within the water column by water itself (depth) and by particulate inorganic and organic matter and dissolved organic matter. Losses of carbon from the system include burial, accumulation of detrital carbon, and losses through grazers to higher trophic levels. Symbols are after Odum (1971).

Environmental Variables and Forcing Functions

The environmental variables and forcing functions modeled were solar and submarine PAR, PAR attenuation as a function of water column turbidity and epiphytic fouling, tidally variable and fixed water depths relative to the SAV canopy, and daily water temperature and photoperiod. Two simulation model versions are referred to in the following sections, model version 2 and version 4. Model version 2 incorporated fixed inputs for depth and attenuation coefficient designed to represent annually averaged environmental conditions. Model version 4 incorporated stochastically fluctuating inputs based on an eight-year data base for York River shoal habitats (see Moore 1992).

Table 1 gives descriptive statistics for those parameters used for version 2 simulations that were derived from field data for shallow water habitats in the lower Chesapeake Bay.

Table 1. Environmental parameters and forcing functions used for the nominal case of version 2 of the Polyhaline SAV Model. Annual means and ranges are for smoothly varying functions, and were derived from observations in the lower Chesapeake Bay (Wetzel and Neckles 1986). Depth and PAR attenuation were fixed within model runs, and were changed for different model cases (see text).

Variable	Units	Annual mean	Annual range
Temperature	°C	16.2	2.5-30.0
Daily incident PAR	Ein m ⁻² d ⁻¹	28.2	11.5-45.0
Water depth	m	1.0	
PAR attenuation	m ⁻¹	1.0	
Photoperiod	hours	11.8	9.5-14.0

For the non-variable model (version 2), the environmental parameters and forcing functions were simulated using trigonometric functions fit to available in situ data (Table 2). The downwelling PAR attenuation coefficient (K_d) was treated as a constant (annual mean) for different simulation scenarios in model version 2 principally because of a lack of spatial and temporal data. For the variable-forcing model (version 4), the natural variability characteristic of surface incident solar irradiance (PAR_0) and K_d was incorporated by using eight-year (1984-1992) monthly means and ranges measured in situ as part of a bi-weekly monitoring program in shoal areas of the York River estuary (Moore 1992). Version 4 model simulations calculated K_d as a variable with a random component that varied within measured statistical limits. Two temporal scales were explored for K_d variability, seasonal as defined by temperature and used in establishing habitat requirements (Batiuk et al. 1992), and monthly as used for solar irradiance estimation. To derive K_d estimates, three approaches were used: 1) site-specific statistical estimates using seasonal mean values only, 2) site-specific statistical estimates using seasonal means and the standard deviation, and 3) site-

specific variation using monthly ranges, the annual minimum and random variability within the observed range.

Table 2. SAV model version 2 forcing functions, based on environmental data from the lower Chesapeake Bay.

Water temperature	$T = 16.25 - 13.75 \cos[2\pi(\text{day}-25)/365]$
Incident solar irradiance	$PAR_o = 28.25 - 16.75 \cos[2\pi \text{ day}/365]$
Photoperiod	$PP = 11.75 - 2.25 \cos[2\pi \text{ day}/365]$
Tidal water depth	$Tlag = 0.842105263 Lday$ $Z(t) = 1.25 + 0.275 \cos[2\pi(\text{hr}-Tlag)/12]$
Submarine, time-dependent PAR irradiance	$PAR(t, \text{ hourly}) = PAR_o / (0.63662 PP) \cos[\pi(\text{hr}-12)/PP]^+$ $PAR(z) = PAR(t) \exp[-K_d z]$
Note: day = day-of-year (1-365), Tlag = tidal phase lag, Lday = lunar day (0-28).	

For the simulation studies, three sites, all in the York River, were used for data input and parameter estimation for the environmental variables and forcing functions:

1. Guinea Marsh, a shoal monitoring site located at the York River mouth and having stable eelgrass beds;
2. Gloucester Point, a shoal monitoring site, ca. 10 km upriver from Guinea Marsh, and having eelgrass beds that are highly variable from year to year; and,
3. Claybank, a shoal monitoring site ca. 20 km upriver from Guinea Marsh which historically marked the upriver limit for eelgrass, but which has not supported a stable SAV community since the early 1970s.

These factors were incorporated as controls on various pathways of carbon flow, most of which also operated within intrinsic biological limits. The light-related parameters controlled photosynthesis by eelgrass and epiphytes. Temperature controlled the rate coefficients for photosynthesis and respiration by eelgrass and epiphytes, root/rhizome translocation by eelgrass, ingestion and respiration by grazers, and the daily ration of higher level predators, further discussed below.

Biological Processes, Interactions and Controls

The biological processes simulated were photosynthesis by eelgrass and epiphytic microflora, ingestion of eelgrass leaves and epiphytes by grazers, respiration and natural mortality for all biological components, and seasonally controlled immigration-emigration of grazers and higher level predators. The mathematical functions derived to simulate these processes follow the rationale developed originally by Wiegert (1973). Applications for other ecosystem models are given by Wiegert and Wetzel (1979), Christian and Wetzel (1978), van Montfrans et al. (1984), Wetzel and Christian (1984), Wetzel and Neckles (1986), Christian et al. (1988), Neckles and Wetzel (1989) and Christian and Wetzel (1991). Details of the mathematical derivations are given in the following sections.

Eelgrass Photosynthesis and Growth

The processes modeled that determined eelgrass aboveground (shoots, or leaves) and belowground (roots and rhizomes) biomass were photosynthesis, respiration, shoot and root/rhizome translocation, and mortality of shoots and root/rhizome material. Photosynthesis was modeled as a function of a specific rate coefficient (P_{ij}) that was both light- and temperature-dependent, leaf biomass (X_j), and non-linear, negative feedback controls which reduced photosynthesis when either CO_2 or space (SAV density) became limiting, the terms FB_{ij} and FB_{jj} respectively. The general form of the flux equation, F_{ij} , for CO_2 fixation by eelgrass leaves was

$$F_{ij} = P_{ij}X_j [(1 - FB_{ij})(1 - FB_{jj}C_{ij})] \quad (1)$$

The specific rate coefficient for photosynthesis (P_{ij}) was calculated using a rectangular hyperbolic (Monod) function which is dependent on the light-saturated photosynthesis rate, P_{max} at temperature, T , the PAR intensity reaching the leaf surface, PAR_{vp} , and the half-saturating PAR intensity, I_k' :

$$P_{ij} = P_{max}(TEMP) [PAR_{vp}/(I_k' + PAR_{vp})] \quad (2)$$

$P_{max}(TEMP)$ was derived as a linear function of temperature using the data of Wetzel and Penhale (1983) and Evans (1984); for temperatures greater than 25°C, $P_{max}(TEMP)$ was reduced linearly such that at 30°C, $P_{max}(30)$ equaled one-half the rate at 25°C (Equation 3).

$$P_{max}(TEMP) = (0.000162*TEMP + .0041)*[1 - (TEMP-25)/(35-25)] \quad (3)$$

I_k' then was calculated as a linear function of $P_{max}(TEMP)$ based on Penhale (1977) (Equation 4).

$$I_k' = 15220.78*P_{max}(TEMP) - 31.86 \quad (4)$$

The PAR intensity reaching the plant surface (PAR_{vp}) was derived as a function of daily solar irradiance (PARD), water depth (Z), the water column PAR attenuation coefficient (K_d), and PAR absorbance by epiphytes (EPLR). PAR intensity at the top of the plant canopy was calculated as a negative exponential function of PARD, Z, and K_d . PAR attenuation due to epiphytes was derived as a hyperbolic function (Equations 5 and 6) of the ratio of epiphyte (X05) to eelgrass leaf biomass (X03), which approached 75% at maximum epiphytic colonization (Murray 1983). The relationship

was derived from Murray's field and experimental data for leaves with a minimum reported biomass ratio of 0.5. Since new, unepiphytized leaves are produced continuously within the canopy, direct application of this relationship would significantly over-estimate epiphyte-induced PAR attenuation. To account for this, we assumed that 50% of the leaf biomass at any time was composed of young leaves having no significant epiphyte biomass, and therefore we reduced the predicted epiphyte attenuation by half.

$$\text{PAR}_{vp} = (1 - 0.75 \cdot \text{EPLR}) \cdot \text{PAR}_Z \quad (5)$$

$$\text{EPLR} = 0.25 \cdot [(\text{X}_{05}/\text{X}_{03}) - 0.1]^{1/2}, \text{ where } 0.0 \leq \text{EPLR} \leq 1.0 \quad (6)$$

Finally, Sand-Jensen (1977) showed that at high light intensity and high epiphyte biomass, diffusion of bicarbonate limited eelgrass photosynthesis by 30%. We incorporated this effect in a manner analogous to epiphytic PAR attenuation as a hyperbolic function of the epiphyte-leaf biomass ratio (Equation 6).

The final terms in Equation 1, FB_{ij} and FB_{jj} , that potentially limit eelgrass photosynthesis are non-linear feedback functions of CO_2 concentration and leaf biomass, respectively. The first of these terms, FB_{ij} , which is the substrate- or donor-control feedback, was derived as a function of the ambient CO_2 concentration (X_i), the CO_2 concentration below which carbon became limiting (A_{ij}), and the CO_2 concentration (G_{ij}) at which photosynthesis by eelgrass approached zero (Equation 7).

$$\text{FB}_{ij} = 1 - (\text{X}_i - \text{G}_{ij}) / (\text{A}_{ij} - \text{G}_{ij}) \quad (7)$$

The second of these terms, FB_{jj} , which is the recipient- or self-control feedback, was derived as a function of leaf biomass (X_j), the leaf biomass (A_{jj}) above which space or some density- dependent factor (e.g., self-shading, crowding, nutrient limitation) limited growth, and the maximum leaf biomass (G_{jj}) that could be maintained metabolically (i.e., analogous to "carrying capacity" in population models) (Equation 8).

$$\text{FB}_{jj} = (\text{X}_j - \text{A}_{jj}) / (\text{G}_{jj} - \text{A}_{jj}) \quad (8)$$

Both of these density dependent feedbacks are constrained mathematically to assume values in the range 0.0 to 1.0.

The final term, C_{ij} in Equation 1, is a metabolic correction factor which allows for maintenance ($d\text{X}_j/dt = 0$) at maximum standing stock ($\text{X}_j = \text{G}_{jj}$). It was calculated as

$$\text{C}_{ij} = 1.0 - (\text{R}_j / \text{P}_{ij}) \quad (9)$$

where R_j equals the specific rate coefficient for leaf respiration and P_{ij} is as defined previously. The correction term was derived such that the instantaneous rate of CO_2 uptake exactly equalled the rate of metabolic loss when no other factors limited growth at the maximum standing stock.

Natural losses of eelgrass leaf biomass occurred through respiration, mortality, and translocation of organic carbon compounds to roots and rhizomes. Respiration and leaf mortality

were derived as linear functions of leaf biomass and a specific rate coefficient. The specific rate coefficient for respiration was estimated as the sum of a basal rate (Murray 1983; Murray and Wetzel 1987) and a specific rate operating only during the photoperiod that was linearly dependant on the realized rate of photosynthesis (Biebl and McRoy 1971; McRoy 1974). Both rate coefficients were temperature dependent and statistically derived (Biebl and McRoy 1971; Nixon and Oviatt 1972; Murray 1983). The specific rate coefficient for leaf mortality was calculated using a cosine function and ranged from 0.5% to 3.0% per day with the maximum rates occurring in mid-summer (Vaughan 1982).

Translocation was perhaps the least documented process affecting eelgrass dynamics. McRoy (1974) indicated that the maximum potential translocation to roots and rhizomes was 17% of net leaf organic matter production. We set the maximum specific rate coefficient at 17% of net productivity and reduced the rate using negative feedback control functions when either aboveground or belowground compartments became limiting. The derivations are analogous to Equations 3 and 4.

Epiphytes

Epiphytes were modeled as an autotrophic community dominated by microflora (Murray 1983; Murray and Wetzel 1987; Neckles 1990; Neckles et al. 1993). Processes affecting epiphyte dynamics were photosynthesis, respiration, natural mortality, and grazing.

Photosynthesis by the epiphytic community was derived mathematically as for eelgrass. P_{max} and temperature were positively and linearly related up to 25°C (Penhale 1977). Above 25°C, P_{max} declined linearly such that at 30°C, P_{max} was 75% of the rate estimate at 25°C (Penhale 1977). I_k' was linearly related to temperature in the range 10 to 30°C and increased from 50 to 150 $\mu\text{Ein m}^{-2} \text{sec}^{-1}$ (Penhale 1977). PAR intensity reaching the epiphyte community was derived as for eelgrass except the only reduction in light was due to water column attenuation. Similar to eelgrass (1), photosynthesis by epiphytes was derived as a function of a specific rate coefficient given available PAR and temperature, the compartment biomass, and non-linear feedback control functions. For epiphytes, however, the limiting factor for colonization and growth was leaf surface area. We reformulated the density-dependent feedback, FB_{jj} , as the ratio of epiphyte (X_j) to SAV leaf (X_i) biomass rather than epiphyte biomass alone. The relationship between leaf surface area and biomass was statistically derived from our own unpublished data. The feedback function was formulated as

$$FB_{jj} = [(X_j/X_i) - A_{jj}] / (G_{jj} - A_{jj}) \quad (10)$$

where A_{jj} and G_{jj} were biomass ratios. Natural losses due to respiration and mortality were modeled as for eelgrass except the specific rate coefficient for mortality was fixed at 0.5% per day.

Grazers

Grazing on epiphytes were modeled using data available for isopod and amphipod populations typical of lower Chesapeake Bay eelgrass communities. Their dynamics were determined by preferences for, and ingestion of, organic resources, resource-specific assimilation

efficiencies, egestion, respiration, natural mortality, seasonally variable immigration-emigration, and loss via predation by higher trophic levels.

Ingestion of epiphytes by grazers was derived as a function of the preference or selectivity for a specific resource, the maximum potential specific rate of ingestion, the grazer biomass, and non-linear feedback control functions dependent on resource availability and grazer density. The mathematical form of the equation was analogous to Equation 1 for eelgrass photosynthesis. Ingestion of eelgrass leaves was derived in the same manner. The preference values and the maximum potential ingestion rates of eelgrass leaves and epiphytes were derived as functions of temperature and based on data for gammarid amphipods (Zimmerman et al. 1979). As in Equations 3 and 4, the feedback control terms were derived as functions of resource availability and grazer density.

Immigration and emigration were not modeled explicitly. However, Marsh (1973) found few epifaunal organisms in Chesapeake Bay eelgrass communities from November to March when water temperatures are generally below 10°C. Therefore, seasonality in grazer standing stocks was incorporated by restricting epiphytic grazing to periods when water temperature was greater than 10°C. Mid-summer declines in grazer population densities have also been documented and correspond to summer increases in predatory fish densities (Orth and Heck 1980; Diaz and Fredette 1982). Loss of grazers to fish predation was incorporated implicitly in the model as a function of date and temperature and was based on the seasonal pattern of predatory fish abundances in eelgrass meadows of North Carolina (Adams 1976a 1976b) and their reported food preferences, daily rations, and assimilation efficiencies for specific prey (Hoss 1974; Adams 1976c; Peters et al. 1976). Respiration by grazers was modeled as a function of temperature. Grazer mortality was fixed at 1.0% per day.

Numerical Computation and Simulation Analysis

The models were programmed in FORTRAN 77. All time-dependent equations were solved using simple Euler numerical integration (Wiegert and Wetzel 1974). The time step for integration was 1.0 hour. Appendix A contains the complete FORTRAN 77 source code listing for the SAV simulation model.

Simulation analyses consisted of first establishing a "nominal" run by repeated simulations of the model until limit-cycle behavior (within three years) and long-term stability occurred (over five years) and the predicted rates, material fluxes, and standing stocks agreed with available field and/or laboratory data. Various models were then run to explore the effects of selected physical-chemical and biological perturbations on eelgrass dynamics. The nominal case and test cases all were started from the same initial conditions. Two complete series of analyses have been completed to date using two model versions (versions 2 and 4).

Model version 2 simulation analyses addressed physical-chemical controls and biological interactions without incorporating environmental variability and portions of these studies have been reported in the literature (Wetzel and Neckles 1986). The results of these past model studies will be briefly reported here for continuity. The analyses were divided into two series of simulations. The first series included physical (environmental) controls only. The second series addressed the potential for epiphyte-grazer interactions to control eelgrass dynamics by varying the grazing

pressure on epiphytes. Grazing pressure was varied under both nominal and perturbed physical conditions in the model.

A final series of simulation studies using version 2 explored implicitly the effects of water column nutrient enrichment and epiphyte grazing intensity (Neckles 1990). The effects were noted by varying surrogate variables in model simulation runs. In this case, increases in the rate coefficient for epiphyte photosynthesis was used as the surrogate for nutrient enrichment.

Version 4 simulation analyses addressed physical-chemical controls on eelgrass growth, abundance and depth distribution relative to specific water quality parameters. Simulation studies investigated in situ variability of environmental parameters for shoal areas that encompass SAV historical or present distribution patterns. With the incorporation of statistically-defined random variability in the environmental forcings, specifically with regard to the underwater light climate, modeling scenarios explored whether such fluctuations might be important in the natural cycle of SAV productivity and in determining water quality criteria for SAV (Batiuk et al. 1992).

SIMULATION RESULTS AND DISCUSSION

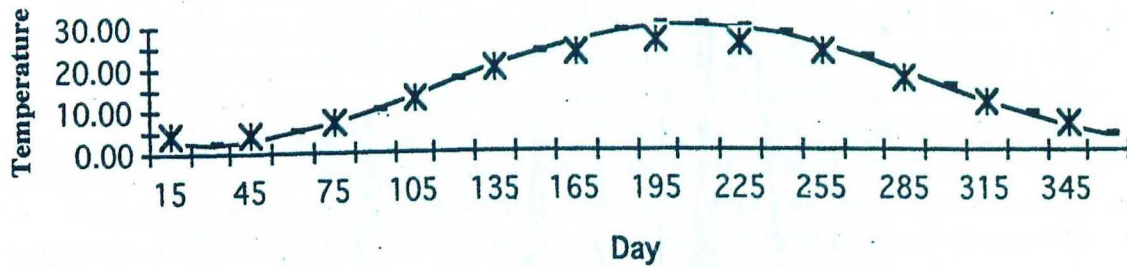
Nominal Simulation Analyses (Version 2: non-variable inputs).

Figure 6 illustrates the measured and simulated values for three of the four principal physical forcing variables in the model, water temperature, solar irradiance, and the downwelling PAR attenuation coefficient. Tidally varying water depth relative to plant canopy height is not shown. The high degree of natural variability is evident in all three and this variability is obviously not completely described by the trigonometric functions used to simulate these variables. Because of the high in situ variability, general lack of data at the time, and poor agreement between the simulated and observed values, the attenuation coefficient was fixed at 1.0 m^{-1} , a value characteristic of all observations from a historically stable eelgrass bed (Moore 1992). Also, for the version 2 nominal simulation, the depth was fixed at 1.0 m (MLW) for comparative purposes, because this represented the predominate depth of in situ sampling within existing eelgrass beds. Therefore, nominal model predictions for the various compartments best represent long term averages and do not address spatial or temporal variability.

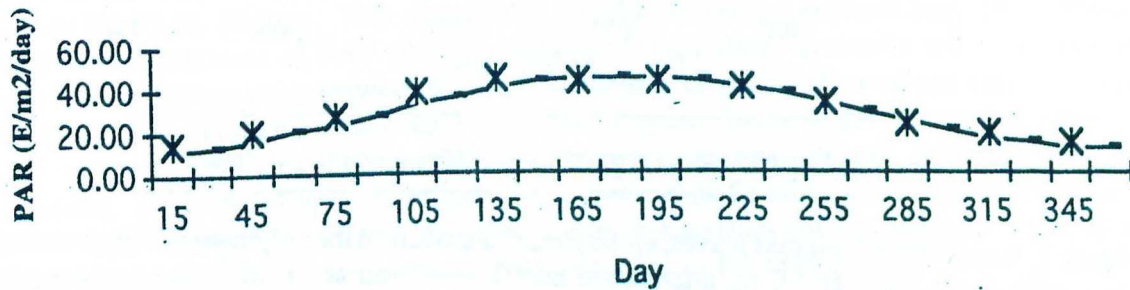
The simulated dynamics for the standing stocks of eelgrass leaves, epiphytes, and grazers under nominal conditions are given in Figure 7. All components demonstrated limit cycle behavior and were stable over time (up to ten years of simulated dynamics). Eelgrass biomass maxima occurred in late April, reaching 143 g C m^{-2} . Epiphyte biomass reached a maximum of 79 g C m^{-2} in late June. The grazers feeding on these epiphytes reached their maximum biomass, 0.93 g C m^{-2} , during the late summer. The epiphyte:eelgrass biomass ratio was 0.04 during the early spring growth of eelgrass and reached 1.5 during the late summer eelgrass decline. These values are consistent with field estimates (Murray 1983; Neckles 1990; Neckles et al. 1993). At maximum epiphyte biomass, PAR reaching the plant epidermis was attenuated 22% by epiphytes and eelgrass photosynthesis was reduced 10% by epiphyte-limited CO_2 diffusion. The maximum sustainable standing stock for the above-ground eelgrass component (leaves) was set to 150 g C m^{-2} in the density-dependant feedback control function (Equation 4), which was the maximum reported field density for

Physical Forcing Functions

A. Water Temperature



B. Solar PAR



C. Attenuation

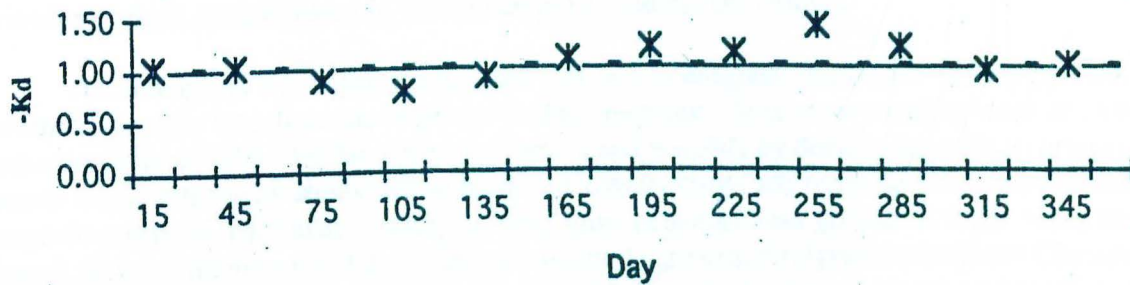


Figure 6. Modeled (dashed line) versus observed (*) physical parameters for the polyhaline SAV model. The in situ values are the monthly means of biweekly observations taken over the period 1985 to 1992 for water temperature and the attenuation coefficient (K_d). The solar PAR data are the monthly means of daily integrated PAR taken at the VIMS over the period 1988 to 1991.

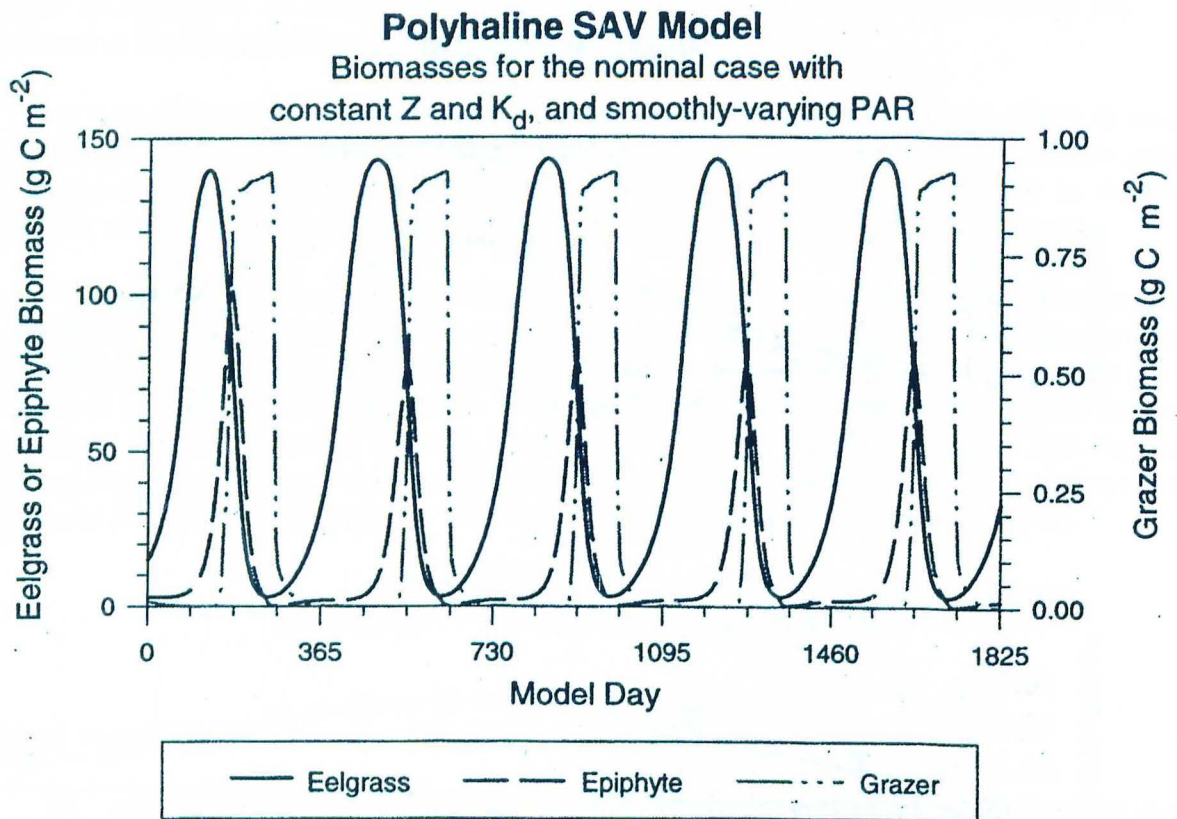


Figure 7. Biomasses of eelgrass leaves, epiphytes, and grazers in the polyhaline SAV model nominal case, with depth fixed at 1.0 m, attenuation coefficient fixed at 1.0 m^{-1} , and using a smoothly varying annual PAR cycle, designed to mimic healthy habitats in the lower Chesapeake Bay.

communities in the lower Chesapeake Bay (Orth and Moore 1983). For the nominal simulation, this feedback function reduced potential eelgrass growth by 88% during periods of maximum biomass, suggesting that some density-dependent factor (e.g., self-shading) may limit SAV growth at certain times of the year.

The physical factors controlling eelgrass production in the simulation are submarine irradiance and temperature. Simulated eelgrass photosynthesis (Figure 8) is light-limited throughout the year and particularly during mid-summer. Temperature (Figure 8B) also contributes to the mid-summer decline as the temperature-dependent respiration rate exceeds the rate of photosynthesis (Figure 8A). Eelgrass is near its equatorward biogeographic extent and the high summer water temperatures ($>25^{\circ}\text{C}$) impose a significant stress on the plant (Wetzel and Penhale 1983; Evans 1984). Together, light limitation and thermal stress contributes to the mid-summer decline in eelgrass production and biomass. Production (Figure 8A) and biomass (Figure 7) do increase in the fall, with the increase in biomass continuing through the following spring.

The overall pattern and standing stock predictions agree with field data but the temporal pattern is out of phase with some data sets; i.e., simulated standing stocks precede in situ observations by ca. 30 days. This discrepancy could have several explanations. First, given the temperature dependence of SAV photosynthesis and respiration, unusually warm or cold years would lead to discrepancies between the temporal pattern of in situ observations versus simulated results based on multi-annual averaged temperature cycle. Second, the factors that control the magnitude and pattern of translocation of organic matter to roots and rhizomes are poorly understood. The assumption of a constant 17% of net carbon fixation is an oversimplification and requires better information. Third, the model does not account for the energetic costs of non-vegetative reproduction; flowering and seed production in the spring would lower or delay the accumulation of above-ground biomass. Fourth, solar irradiance and water column PAR attenuation are highly variable. For model version 2, a non-variable, continuous function was fitted to local data to predict daily solar irradiance, and the PAR attenuation coefficient was fixed at an annual mean value. Both should be modeled as stochastic functions that operate within the range of natural variability, which are addressed in model version 4 studies (see below).

These nominal simulations suggested that in situ eelgrass photosynthesis and growth are governed primarily by submarine irradiance and temperature. Maximum standing stock and annual production appear limited by these physical factors and possibly by density-dependent controls that operate when submarine irradiance is high and temperatures are optimal. Sediment dissolved inorganic nutrients, particularly nitrogen, may limit above-ground growth at these times but, in general, do not limit the current distribution or preclude growth of eelgrass in the lower Chesapeake Bay (Orth 1977). Although physical and chemical factors limit eelgrass growth and production to less than its intrinsic maximum, the simulated dynamics of the model's components demonstrate long term stability and predicted densities agree with field estimates. Qualitatively, the model can be applied as an analytical tool for simulating the effects of various changes in physical-chemical and biological interactions on SAV productivity (Wetzel and Wiegert 1983).

Polyhaline SAV Model

Eelgrass production and respiration in the nominal model case

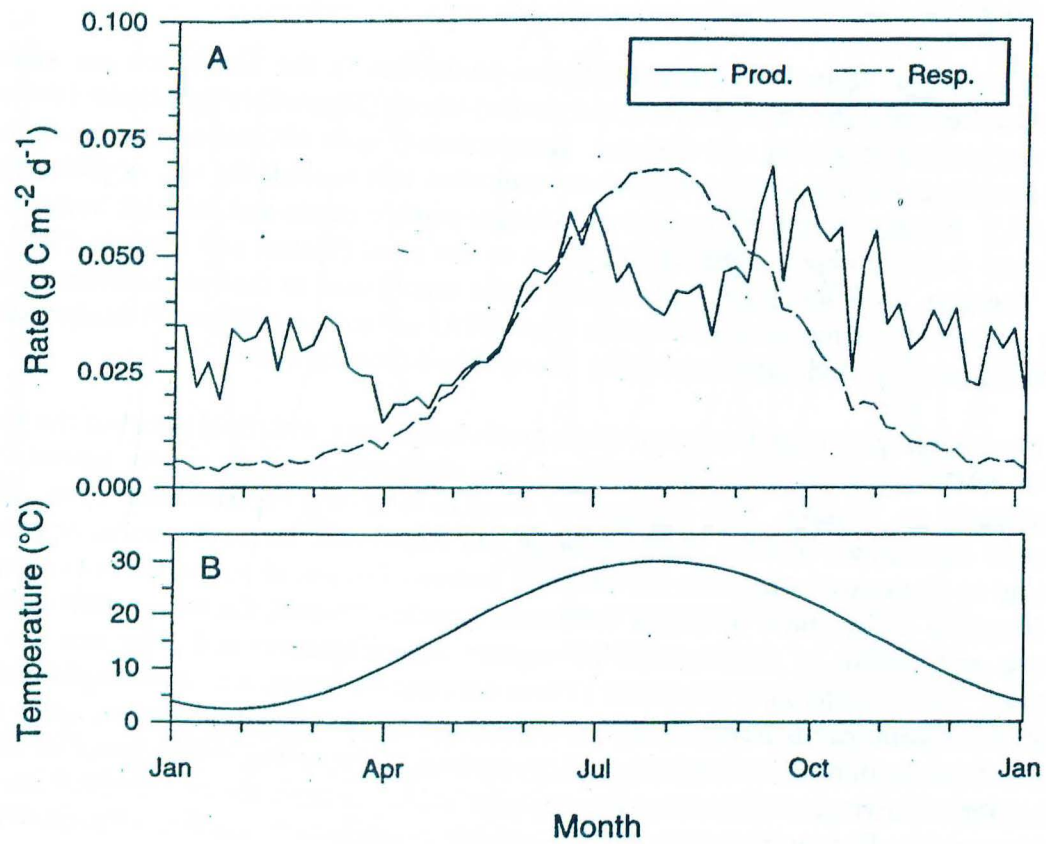


Figure 8. Polyhaline SAV model nominal case with fixed depth and light attenuation coefficient. A. Rates of photosynthesis and respiration in the nominal model case. B. The annual temperature cycle used in the model. While photosynthesis is light-limited much of the year, temperature drives respiration rates higher than photosynthesis during the summer months.

Polyhaline SAV Model

Sensitivity to fixed values of depth and light attenuation

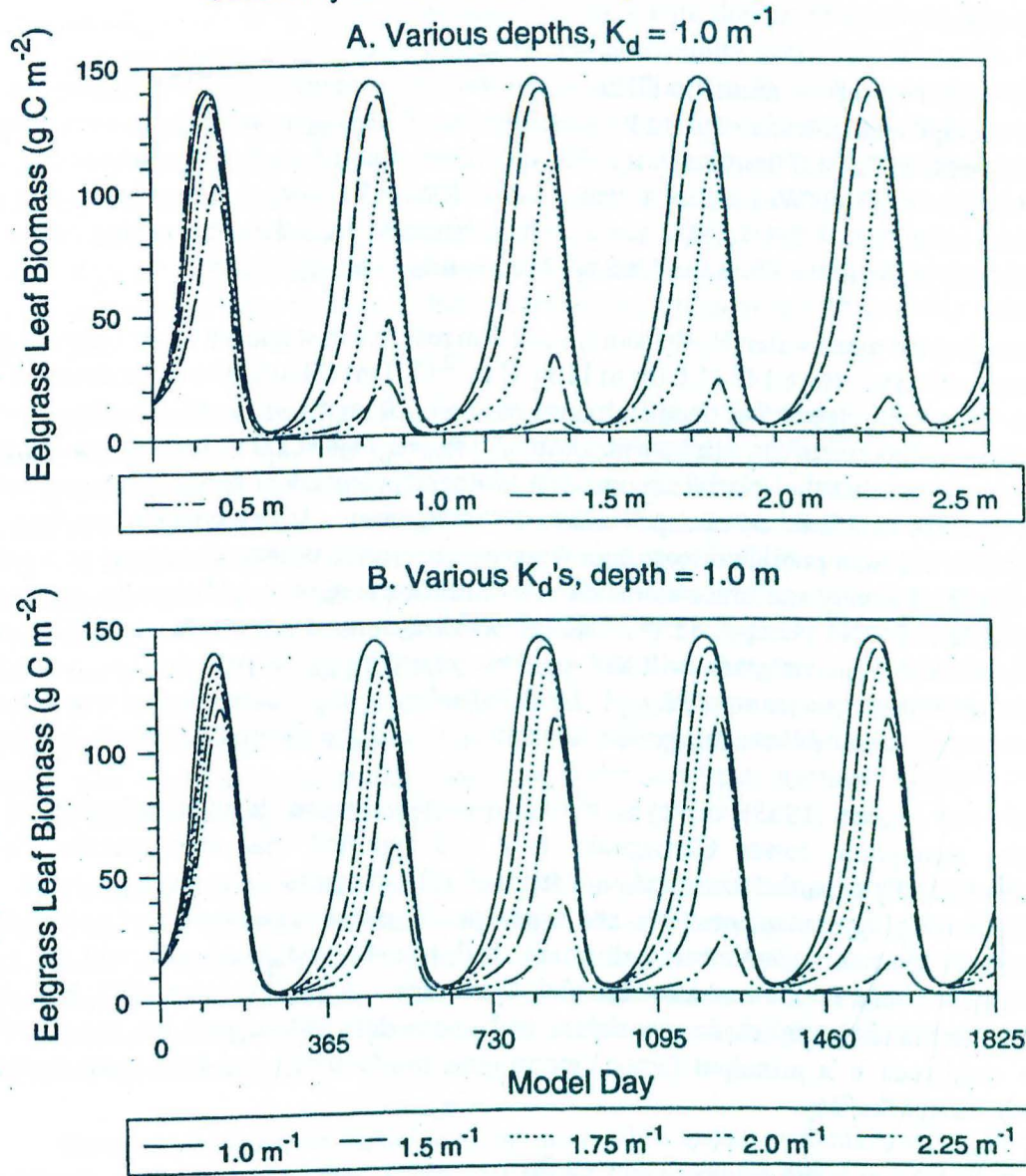


Figure 9. Sensitivity of eelgrass leaf biomass predicted by the polyhaline SAV model to changes in depth and light attenuation coefficient (K_d). A. Various depths with K_d fixed at 1.0 m^{-1} . B. Various K_d 's with depth fixed at 1.0 m.

Model Responses to Physical Regime and Epiphyte-Grazer Interactions

Physical Regimes

The sensitivity of the model to different physical regimes was evaluated by varying water depths and average annual water column PAR attenuation. Changes in water depth were simulated by fixing the depth at 0.5, 1.0 (nominal), 1.5, 2.0, and 2.5 m. For 0.5 m, the upper temperature limit was also increased to 33°C (Wetzel, unpublished data). The PAR attenuation coefficient was varied through fixed values 1.0, 1.5, 1.75, 2.0, and 2.25 m^{-1} , based on field data for existing or historical eelgrass habitats in the lower Chesapeake Bay (Moore, unpublished data).

Changing the mean water depth from 1.0 to 2.0 m resulted in stable but lower eelgrass annual peak biomass estimates, from 143 (1.0 m) to 110 g C m^{-2} (2.0 m) (Figure 9A) and a decrease in the importance of density-dependent feedback control. At 1.0 m fixed depth, density-dependent controls limited eelgrass leaf growth by a maximum 93%, whereas at 2.0 m, growth was limited by 50%. This suggests that the relative importance of density-dependent factors changes not only temporally but also as a function of depth within the community. At fixed depths less than 1.0 m and greater than 2.0 m, a predicted loss of above-ground eelgrass biomass occurred as a result of temperature effects in very shallow water and PAR limitation at depth. Additionally, eelgrass leaf biomass was sensitive to changes in the average water column PAR attenuation (Figure 9B). Simulations indicate that eelgrass will not survive over long time periods (i.e., years) with attenuation coefficients greater than 2.0 m^{-1} . Modeled eelgrass population survival was somewhat less sensitive to K_d than has been suggested for natural population (Wetzel and Penhale 1983).

Orth and Moore (1983) surveyed the relative density and depth distribution of SAV communities throughout lower Chesapeake Bay and reported that monospecific Eelgrass communities typically occurred to a maximum depth of 1.2-1.6 m relative to mean sea level. Long term studies (surveying, transplantations, and experiments) across a continuum of potential SAV sites, from presently non-vegetated to well-established, has shown that eelgrass will not survive where the water column PAR attenuation coefficient averages $>1.5 \text{ m}^{-1}$ (Moore 1992; Batiuk et al. 1992). The simulation analyses are consistent with these data and support the hypothesis that submarine irradiance is a principal factor determining productivity and long term survival of eelgrass in Chesapeake Bay.

Epiphyte-Grazer Interactions

The correspondence of eelgrass biomass and productivity with the annual variation of physical factors suggested that the epiphyte and grazer components had little effect on eelgrass productivity. To test this, additional simulations were run without epiphytes and their grazers. In these simulations, predicted dynamics and standing stocks of eelgrass leaves were nearly identical to those in the nominal simulation. Predicted photosynthetic rates also were similar most of the year, but increased slightly during mid-summer relative to nominal simulations. It can be inferred, then, that in the nominal case, grazing maintains epiphyte density below levels that would limit eelgrass photosynthesis and growth. As a corollary, any perturbation that would allow epiphytes to outgrow their grazers might be expected to have an adverse effect on eelgrass productivity. In the absence of impacts on the epiphyte-grazer components, any perturbation which would limit eelgrass photosynthesis would also adversely effect eelgrass productivity.

The simulated response of eelgrass to changes in epiphytic grazing pressures combined with various levels of water column PAR attenuation is shown in Figure 10. Under nominal physical conditions ($K_d = 1.0 \text{ m}^{-1}$), eelgrass biomass declined with decreased epiphyte grazing pressure. However, even at very low grazing intensity, the community persisted. The effect of reduced grazing and the concomitant increase in epiphyte biomass became more dramatic as PAR attenuation increased. For example, nominal grazing with $K_d \leq 1.75 \text{ m}^{-1}$ resulted in long term SAV stability, whereas grazing pressures less than 25% of those presently observed in the environment caused a predicted diminution or, under the more severe cases, a complete demise of eelgrass leaf biomass where light attenuation coefficients were equal to or greater than 1.5 m^{-1} (Figure 10). Of note, at K_d 's of 1.75 and 2.0 m^{-1} , where modeled eelgrass populations survived under nominal grazing pressures (Figure 9), they failed to survive in combination with severely reduced grazing (Figure 10).

These simulations indicate that under reduced light conditions, additional factors, such as grazing, which control epiphytic growth can impact eelgrass production and community stability. For eelgrass communities typical of the lower Chesapeake Bay which are stressed naturally by sub-optimal submarine irradiance and high, late-summer water temperatures ($>25^\circ\text{C}$), relatively small changes in the factors controlling epiphyte growth may greatly affect community stability, recalling that epiphytes interfere with both light transmission to the leaf surface as well as gas diffusion across the leaf surface. For example, increased water column nutrients and reduced grazing favor epiphyte growth and reduce SAV photosynthesis (Murray 1983; van Montfrans et al. 1984; Borum 1985; Twilley et al. 1985). These simulation experiments suggest that under conditions of reduced grazing intensity and increased PAR attenuation, the SAV community would not survive over longer time frames. One might also expect that conditions which accelerate intrinsic epiphyte growth would also have an adverse impact on eelgrass communities.

To more closely examine the impact of the epiphyte assemblage on eelgrass, Neckles (1990) extended the SAV simulation model using the results of mesocosm experiments. These laboratory experiments investigated plant-epiphyte-grazer interactions relative to eelgrass photosynthesis, growth and community stability. Grazer effects (presence/absence) on epiphyte density and SAV growth were examined under both ambient and nutrient-enriched (three-fold ambient concentration) conditions in mesocosms.

However, because the model does not explicitly include nutrients, a surrogate variable was used for these simulation studies. It has been shown that a primary effect of increased water column nutrient concentration is an increase in both the realized growth rate of microalgae and the leaf-area-specific density of epiphytes (Murray 1983). To simulate this, Neckles (1990) adjusted the model's epiphyte P_{max} upward until predicted epiphyte densities agreed with observed densities from the experimental mesocosms. Increasing P_{max} by two- and three-fold appeared to best represent increased nutrient effects relative to lower bay SAV. The simulations were run for ten-year periods to investigate the effects of eutrophication, epiphytes, and grazers on community stability.

Figure 11 shows model predictions (solid and dashed lines) and mesocosm observations ($\bar{x} \pm \text{SE}$) of the mass ratios of epiphyte to SAV leaf. The upper panels (A & B) in Figure 11 give the results for ambient nutrient conditions with and without grazers present. The observed and model predicted values are remarkably similar for this condition. The outlying observation (panel A during the fall) resulted from an increase in ambient nutrient concentrations during the mesocosm studies

Polyhaline SAV Model

Annual Eelgrass biomass maximum and the interaction of light attenuation and epiphytic grazing

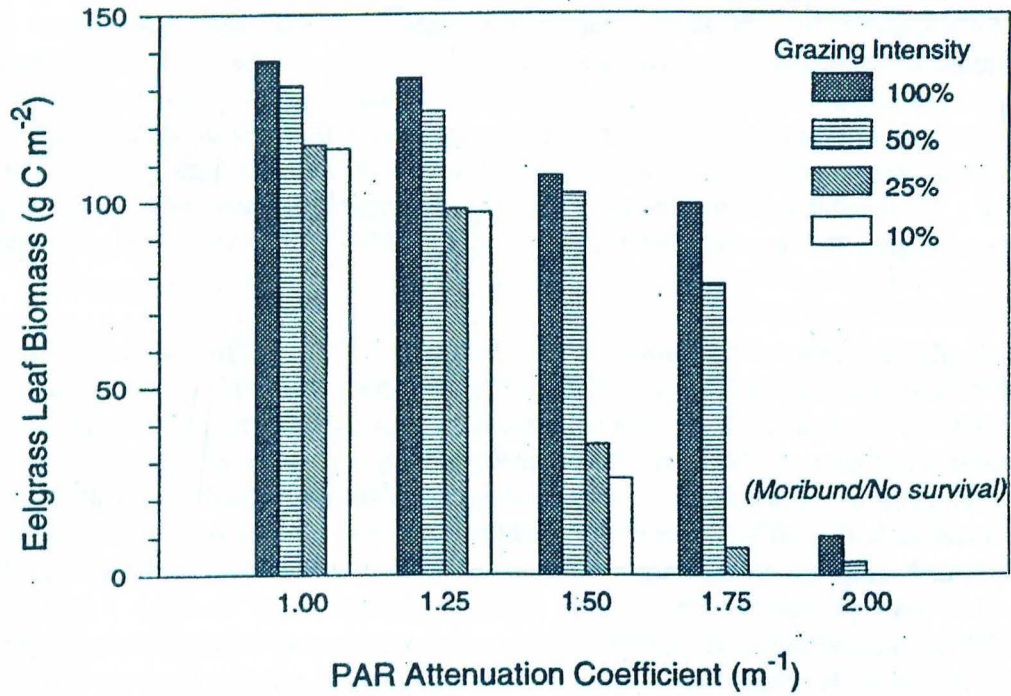


Figure 10. Predicted eelgrass leaf biomass with combinations of attenuation coefficients and grazing pressure (relative to the nominal case) using polyhaline SAV model version 2.

Epiphyte:SAV Biomass Ratios - Grazing and Eutrophication

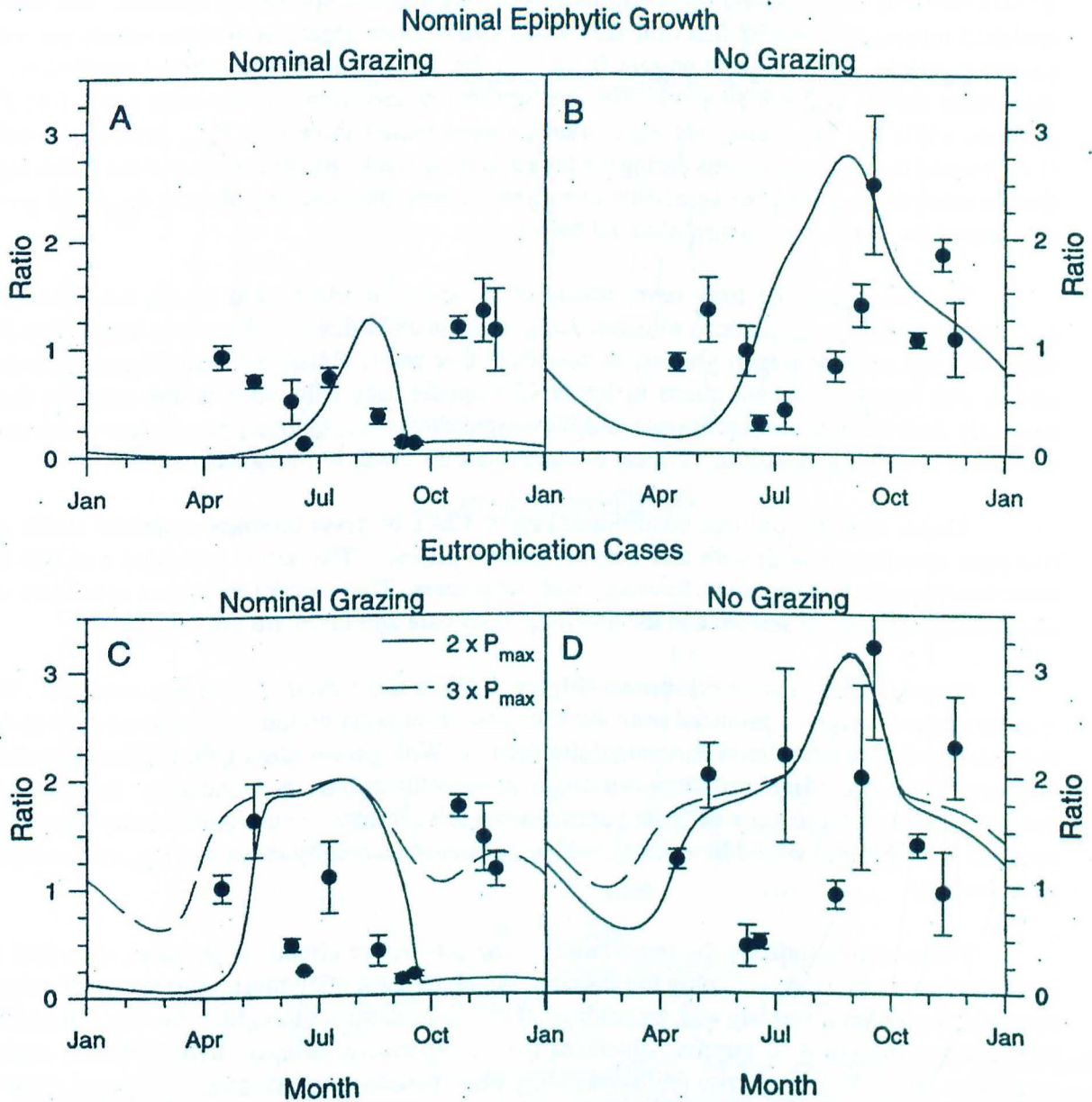


Figure 11. Predicted biomass maximum of eelgrass leaves during model year 5, under combinations of PAR attenuation coefficient (m^{-1}) and relative grazing intensity (percent of the nominal case) in the Polyhaline SAV Model using a fixed depth of 1.0 m.

which the model did not simulate (Neckles 1990). The lower panels (C & D) in Figure 11 give the same comparison except under nutrient enriched conditions. With grazers absent (panel D), model simulations and experimental observations agreed very well except during summer. The nutrient enriched mesocosms during this time developed dense macroalgal populations which the model cannot simulate. With grazers present (panel C), the agreement between model predictions and mesocosm results was not as good. The late summer observations agreed with a two-fold P_{\max} increase while the fall mesocosm experiments agreed with a three-fold P_{\max} increase. Neckles (1990) noted that the mesocosms during the fall experiment had nutrient concentrations much higher than at other times. The poor agreement during the summer resulted in part from the dense growth of macroalgae in the mesocosms as noted before.

To investigate the long term effects of increased nutrients and grazer interactions on community stability, simulations were run using nominal and a three-fold increase in epiphyte P_{\max} both with and without grazers present for periods of five years. Relative to natural grazer densities and in situ nutrient concentrations in lower Chesapeake Bay tributaries where eelgrass did or currently does exist, these conditions would represent extremes. All other parameters of the model were held at nominal conditions typical of contemporary lower bay eelgrass habitats.

Under ambient nutrient conditions (Figure 12A), eelgrass biomass remained stable over five-year simulations both with and without grazers present. The model indicated a ca. 20-25% reduction in peak standing stock, however, without grazers. These model results are consistent with all previous simulation studies and the available field data and literature information.

For enriched nutrient conditions (Figure 12B), a more dramatic pattern emerged. With grazers present, eelgrass persisted over the five-year simulation period, but suffered a 30 to 40% reduction in abundance relative to nominal simulations. With grazers absent, the eelgrass population was not stable over time and demonstrated a protracted decline in abundance. In nature, for conditions such as these, once eelgrass populations reach a threshold minimum density, they would cease to be viable and would be replaced with a different community structure (e.g., macroalgal or planktonic).

These results indicate the importance of the interactive effects of physical, chemical and biological controls for determining the distribution, abundance and long-term stability of eelgrass communities in the lower bay and, by analogy, SAV communities throughout the bay. In addition to the direct influence of physical-chemical (i.e., temperature, salinity, hydrodynamic regime) properties on SAV productivity and community composition, a major control appears to be the intensity of submarine PAR reaching the plant leaf to support photosynthesis. Both grazing and nutrient enrichment appear to act as controls via modification of the submarine light environment.

Effect of Submarine Light Variability on Eelgrass Dynamics (Model Version 4)

Field, laboratory, and simulation studies all indicate that the submarine light environment is fundamental to the depth distribution, productivity, and stability of SAV communities. Understanding the factors attenuating photosynthetically available radiation (PAR) is critical for developing effective SAV management policies and implementing strategies for SAV conservation and enhancement.

Polyhaline SAV Model

Impact of grazing pressure and eutrophication effects
on epiphyte cover as reflected in SAV biomass

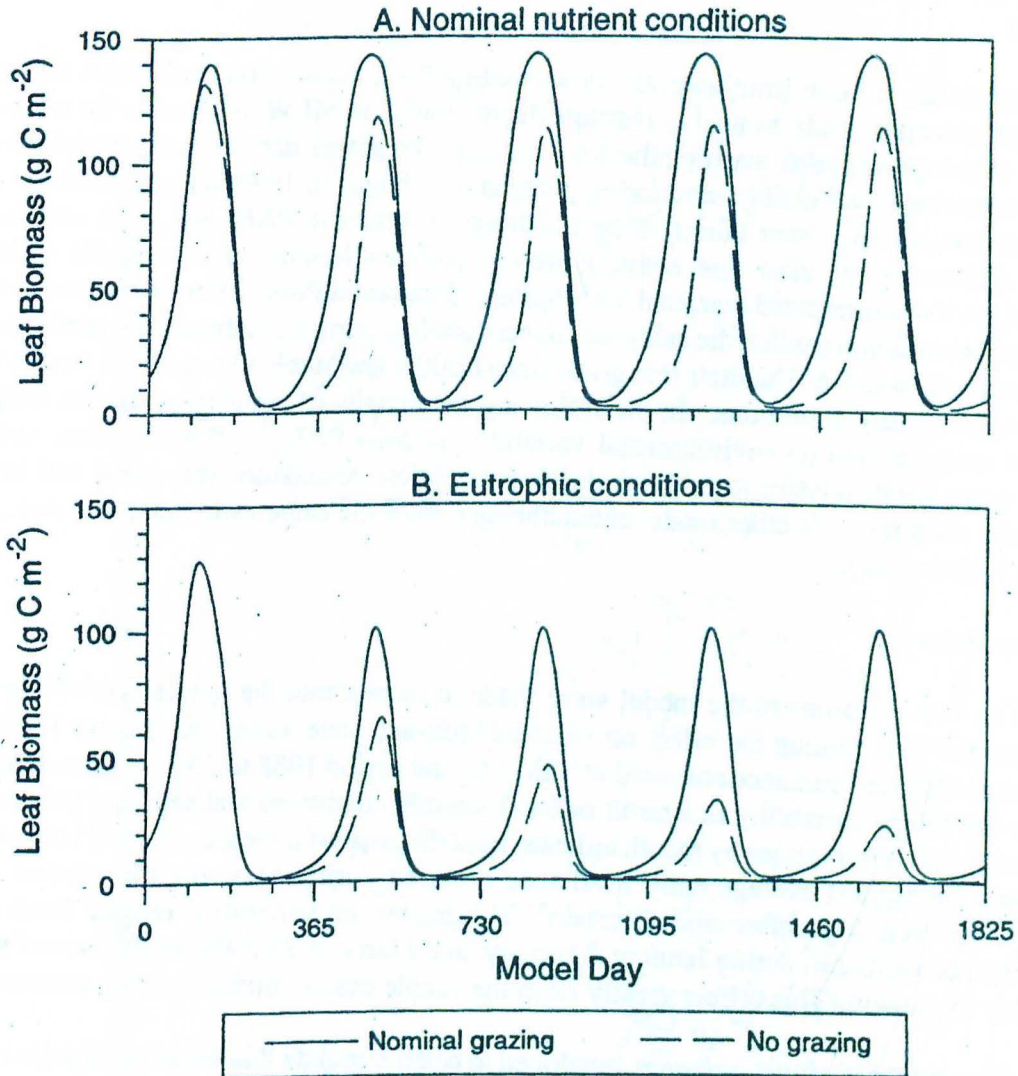


Figure 12. Epiphyte:Eelgrass leaf biomass ratios predicted by the polyhaline SAV model (with fixed depth of 1.0 m) under combinations of grazing and eutrophication conditions. Dots are mean and standard error of ratios from manipulated microcosms (Neckles 1990; Neckles et al. 1993). A. Nominal grazing and epiphytic growth rates. B. Nominal epiphytic growth rate without grazing. C. Nominal grazing rate with enhanced epiphytic growth rate, as a proxy for eutrophication. D. No grazing with enhanced epiphytic growth rate. Figure modified from Neckles 1990.

The simulation model as configured in version 2 treated PAR and the factors attenuating PAR in the water column simply and without regard for environmental variability. The simulation results presented for version 2 modeled solar irradiance as a simple cosine function fitted to local observations. Water depth and the attenuation coefficient were fixed or held constant for a given simulation.

Variability in solar irradiance, the downwelling PAR attenuation coefficient, and variable water depth driven by tides, as well as absolute depth relative to MLW, determine the intensity and temporal variability of PAR reaching the SAV canopy. To gather data to better model submarine PAR, an intensive water quality monitoring program was begun in 1984 in the York River estuary to characterize, among other things, solar irradiance, submarine PAR, and PAR attenuation at selected shallow water sites that either currently (Guinea Marsh) or historically (Claybank) supported SAV or represented marginal SAV habitat (Gloucester Point). For the purposes of model revision and simulation studies, the analyses concentrated on using these three site-specific data sets. These areas represent SAV habitats that grade from healthy and stable to unsuitable for SAV under present water quality conditions. In the following, the results of simulation studies with model version 4 are presented on environmental variability in solar PAR, PAR attenuation, and tidally varying water depth relative to eelgrass depth distribution, abundance (biomass) and long term community stability. All other model characteristics were the same as in version 2 (e.g., grazer-epiphyte interactions).

Solar Irradiance

The first revisions to the model were made to incorporate the natural variability of solar irradiance and determining the effect on selected rates and state variables. Figure 13 shows the daily record of solar irradiance collected at VIMS for the period 1988 to 1991. These data indicate the high day to day variability as a result of local weather conditions and seasonal patterns for the area. Pooling all observations by month indicated that the greatest increase occurred between March and April, the highest average daily irradiance occurred during May and the greatest decrease occurred between September and October. The degree of variability ranged from ca. 50% (coefficient of variation) during January, February, and March to 25-30% (coefficient of variation) from May to August. This differs greatly from the simple cosine model used for version 2.

Two stochastic functions were developed to better simulate this behavior in solar irradiance compared to known variability, specifically based on the VIMS data set. The first description (RANDOM-1) for daily irradiance was based on monthly averages and standard deviations (Equation 11). This daily solar PAR (PARD; $\text{Ein m}^{-2} \text{d}^{-1}$) was derived as

$$\text{PARD} = [\text{AVG} - \text{SD}] + 2 * \text{RAND} * \text{SD} \quad (11)$$

where AVG and SD are the monthly means and standard deviations calculated from the VIMS data set and RAND is a random number in the range 0.0 to 1.0. A second description (RANDOM-2) was based on monthly ranges, in place of standard deviations, of the pooled data and PARD derived as

$$\text{PARD} = \text{RAND} * \text{RANGE} + \text{MIN} \quad (12)$$

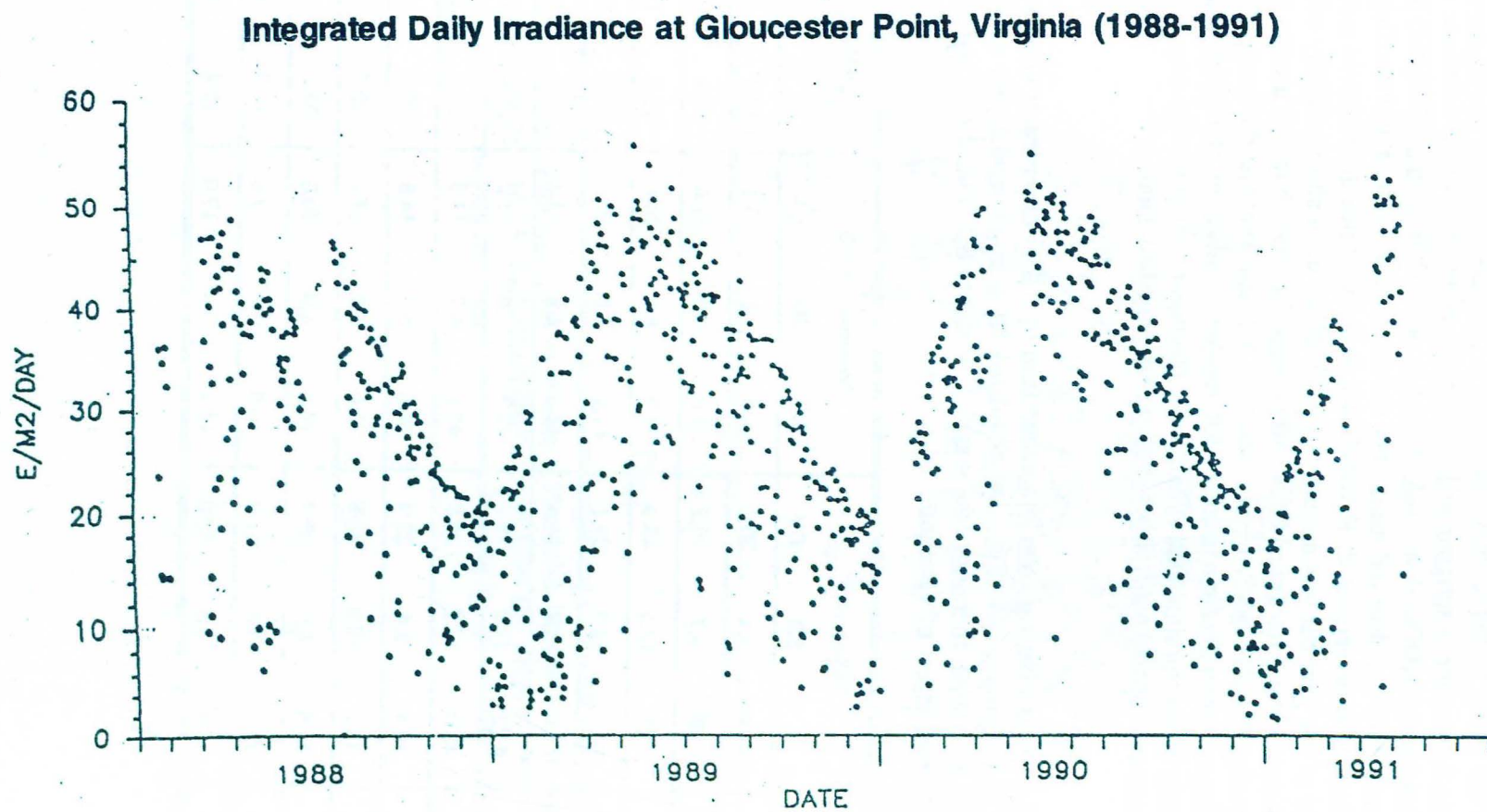


Figure 14. Four years of surface daily PAR irradiance ($\mu\text{Ein m}^{-2} \text{d}^{-1}$) at Gloucester Point, Virginia (1988-1991; data available from the Virginia Institute of Marine Science).

where MIN is the minimum monthly irradiance. Both derivations allowed for intra-annual and interannual variability in daily solar irradiance that is characteristic of lower bay climatic conditions.

Table 3 gives the descriptive statistics for the eight-year VIMS data base and for one-year simulations using RANDOM-1 and RANDOM-2. RANDOM-1 mean monthly PAR predictions compared well with the observed means but the standard deviations and coefficients of variation were significantly underestimated. RANDOM-2 underestimated the observed mean monthly PAR during parts of the year but was a much better predictor of in situ variability overall. Figure 14 gives the observed daily integrated PAR compared to the smooth model used in version 2 and the stochastic model (RANDOM-2) used in version 4. It is apparent that the stochastic model captures the variability and annual pattern in solar PAR very well. However, changing the solar PAR model did not greatly affect the simulated dynamics of eelgrass under otherwise nominal conditions, i.e., with fixed attenuation and water depth (Wetzel, unpublished data).

Table 3. Comparison of solar irradiance ($\text{Ein m}^{-1} \text{d}^{-1}$) observed at Gloucester Point, Virginia (1984-1992) and simulated for a single model year using two stochastic functions (see text). \bar{X} = mean; SD = standard deviation; CV = coefficient of variation.

Month	Observed			Stochastic on SD			Stochastic on range		
	\bar{X}	SD	CV	\bar{X}	SD	CV	\bar{X}	SD	CV
Jan	14.8	7.4	50.0	14.0	3.8	27.1	12.8	6.4	50.0
Feb	19.7	9.7	49.2	19.0	6.0	31.6	18.4	9.5	51.6
Mar	27.7	13.5	48.9	29.9	8.0	26.8	25.7	12.0	46.7
Apr	38.1	14.9	39.1	41.4	7.8	18.8	29.9	13.4	44.8
May	44.4	11.8	26.6	44.8	6.8	15.2	38.6	10.8	28.0
Jun	43.2	12.8	29.6	42.3	5.6	13.2	35.4	13.6	38.4
Jul	43.0	11.2	26.0	42.1	5.7	13.5	33.9	14.1	41.6
Aug	39.4	9.4	23.9	38.6	5.7	14.8	29.7	11.5	38.7
Sep	33.2	10.9	32.8	34.6	5.9	17.1	26.0	11.8	45.4
Oct	23.6	8.9	37.7	24.5	4.9	20.0	22.6	8.0	35.4
Nov	18.2	5.8	31.9	18.8	3.3	17.6	14.0	6.1	43.6
Dec	13.8	5.8	42.0	14.1	2.4	17.0	12.8	5.9	46.1

Daily Solar Irradiance

Comparison of measured and modeled PAR used in SAV simulations

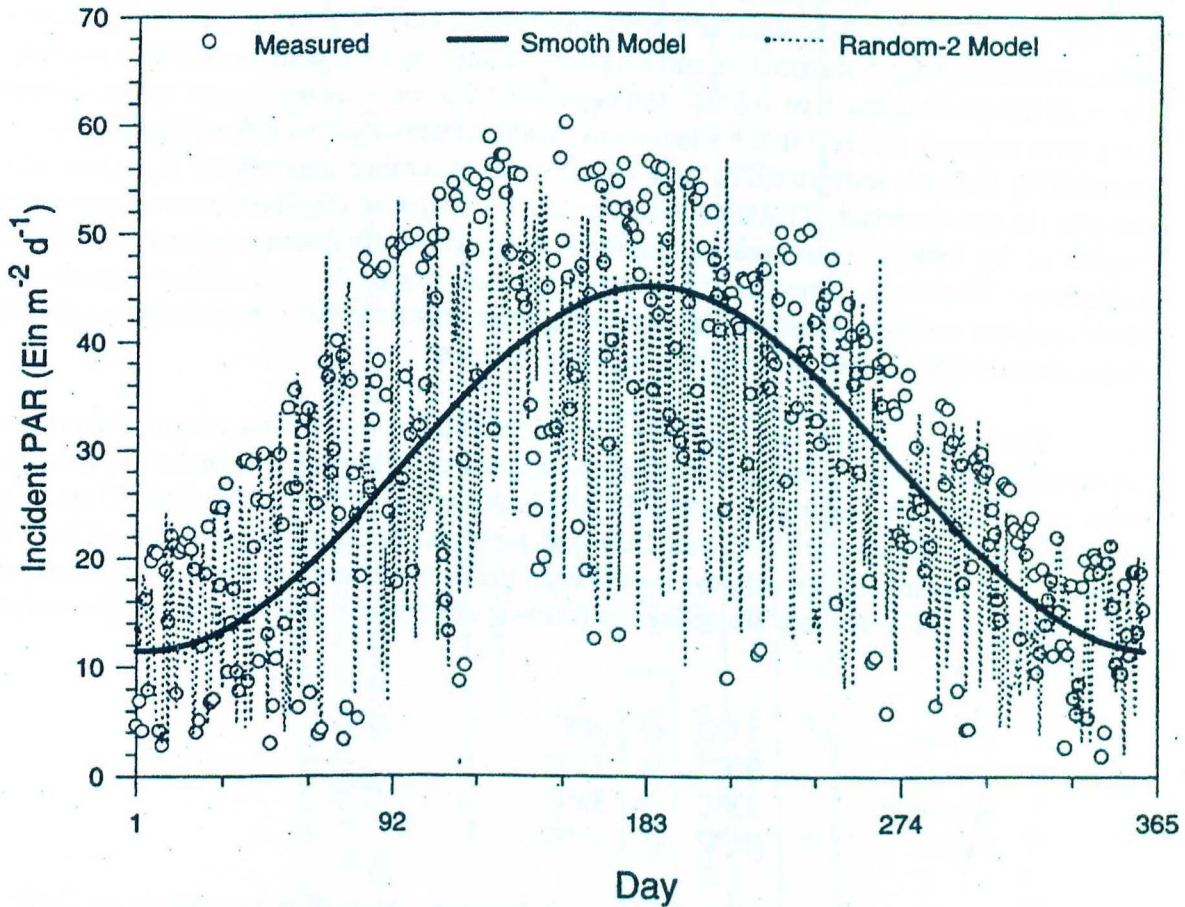


Figure 15. A comparison of daily surface irradiance from a long-term data base at Gloucester Point, Virginia (open circles) with modeled irradiance. The solid line shows the smoothly varying function, $PAR = 28.25 - 16.75 \cos(2\pi DAY/365)$, fitted from the observations. The dashed line shows a stochastically varying function derived from the daily integrated minimum and range from the same data base.

PAR Attenuation

The second component determining submarine PAR intensity in model version 4, and perhaps still the least predictable, is the downwelling PAR attenuation coefficient, K_d . SAV model version 2 treated K_d as a constant for a given simulation. However, these earlier simulation results indicated that the depth distribution and long term stability of SAV were sensitive to changes in K_d . Attenuation coefficients of ca. 2.0 m^{-1} and depths of 2.0 m were the maximum values allowable for long term eelgrass survival in the simulations. Data accumulated from field studies and the shoal monitoring program indicated that these values were higher than maxima for K_d values and water column depths observed in extant eelgrass beds. We proposed that these overestimates could be explained by lack of simulated variability in K_d particularly when coupled with solar PAR variability. That is, inclusion of higher frequency fluctuations in the modeled light environment (both incident and submarine) would result in a greater sensitivity of modeled eelgrass productivity to parameters that determine the in situ light environment.

The impact of time-varying K_d was investigated in two ways. First, temporal averages based on the temperature-delineated growth phases of eelgrass in the lower Chesapeake Bay (Moore 1992) were constructed for three sites in the Shoal Monitoring Program in the York River, Virginia. Guinea Marsh, Gloucester Point, and Claybank represent the spatial continuum of SAV habitat quality, from healthy extant eelgrass meadows of the first site to a presently denuded site upriver at the third site. The seasons are defined as follows; see Moore 1992 for a detailed description of this delineation:

Winter:	13°C	→	0°C	→	9°C
Spring:	9°C	→	23°C		
Summer:	23°C	→	30°C	→	25°C
Fall:	25°C	→	13°C		

The observed seasonal variability within and between these sites is evident in Table 4. In comparison with the earlier simulations (version 2) where K_d was fixed, it can be seen that the Guinea Marsh site has K_d values similar to the SAV model's nominal case value of 1.0 m^{-1} during the spring and fall eelgrass growth periods, while Claybank has K_d 's higher than the predicted maximum value for a sustainable eelgrass community (Batiuk et al. 1992). Seasonal attenuation coefficients at the Gloucester Point site are marginally high for predicted eelgrass survival.

The impact of these differences in site-specific and temporally varying K_d 's is evident in the simulated annual cycle of eelgrass shoot biomass (Figure 15). Simulated eelgrass populations exhibited long term stability under variable K_d forcing for Guinea Marsh and Gloucester Point site-specific data. Despite the introduced variability, the Guinea Marsh case supported a biomass within 90% of that sustained using the fixed conditions and smoothly varying PAR of the nominal case. Similarly, the Gloucester Point site simulation exhibited somewhat reduced, but stable eelgrass population density. Using seasonal patterns at Claybank, however, a simulated eelgrass population rapidly (over three years) declined. Of course, presently there is no eelgrass population at Claybank.

Temporal variability in K_d was further enhanced in another set of simulations, whereby stochastic variations were introduced on a daily basis to the seasonally averaged data (see Table 4). The method for introducing this variability was identical to that used for introducing stochastic

Table 4. Comparison of simulated variability of light attenuation coefficients at three York River, Virginia shoal monitoring sites pooled by month and by temperature-determined seasons. Simulated values are derived from a stochastic functions based on observed (1984-1992) monthly means (\bar{X}) and standard deviations (SD) at each of the sites.

Season	Guinea Marsh				Gloucester Point				Claybank			
	Monthly		Seasonal		Monthly		Seasonal		Monthly		Seasonal	
	\bar{X}	SD	\bar{X}	SD	\bar{X}	SD	\bar{X}	SD	\bar{X}	SD	\bar{X}	SD
Winter	0.985	0.231	1.000	0.105	1.069	0.178	1.095	0.079	1.559	0.387	1.582	0.189
Spring	0.819	0.144	0.813	0.120	1.080	0.229	1.054	0.150	2.103	0.436	2.219	0.249
Summer	1.227	0.249	1.196	0.101	1.508	0.204	1.575	0.100	2.185	0.553	2.374	0.201
Fall	1.221	0.256	1.193	0.176	1.198	0.145	1.148	0.086	1.626	0.314	1.574	0.142

Polyhaline SAV Model

Using seasonally-varying K_d for three sites (York River, Va.)

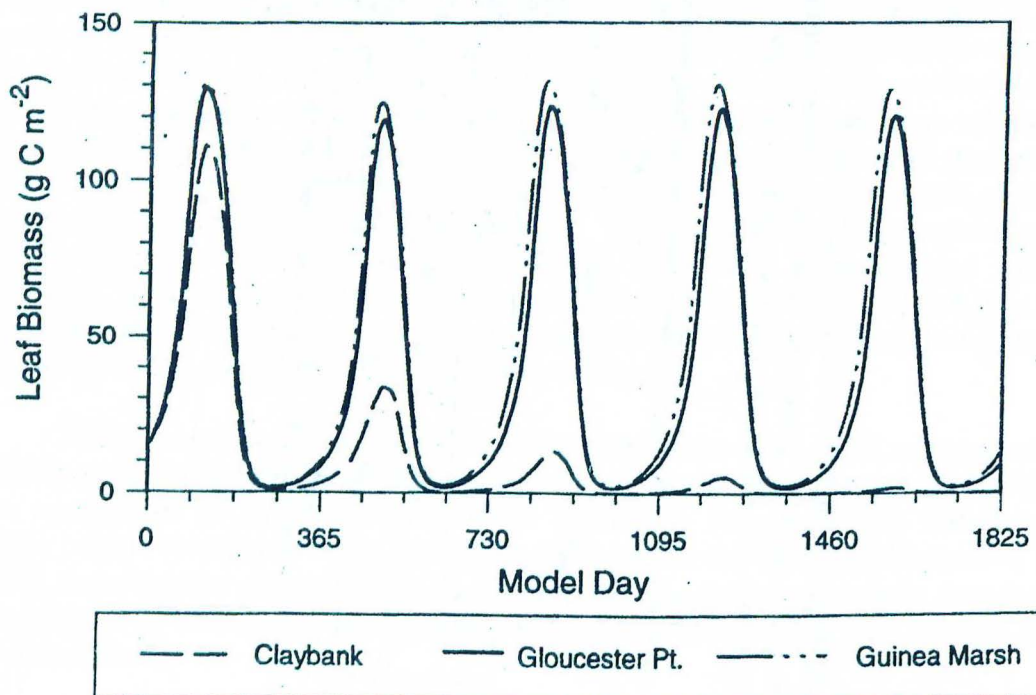


Figure 16. Predicted eelgrass leaf biomass using seasonally varying light attenuation coefficients from three sites in the York River, Virginia. In all cases, depth was fixed at 1.0 m.

variations in the incident PAR data, as described above. Using the Guinea Marsh case as an example, daily random variations about seasonal means imposed little further limitation on simulated eelgrass populations; biomass levels were similar under the imposition of both frequencies of variation in K_d (compare the Guinea Marsh case in Figure 15 with the 1.0 m case in Figure 16).

After examination of the effects of variable versus annual mean K_d on the prediction of eelgrass population survival, it is useful to ask, given these site-specific and temporally varying results, can these simulations reproduce the depth-restricted distributions presently observed, and what will the predicted depth limits be using the restoration criteria for K_d established for SAV habitats (see Batiuk et al. 1992)? Examining the impact of variable K_d for different isobaths provides a test for these seasonally defined restoration criteria. Such tests were performed using K_d 's, including the addition of daily stochastic variations, characteristic of those at the extant Guinea Marsh eelgrass beds (see Table 4). Predicted stable populations were maintained for plant canopies 1.5 m from the surface (Figure 16). Assuming a canopy height of ca. 0.30 m, this corresponds to isobaths of ca. 1.80 m, somewhat shallower than the Tier III target (2.0 m; see Batiuk et al. 1992) for restoration. That is, even though this site has water clarity that is within the restoration criteria, the model does not predict that stable populations can be maintained in waters 2.0 m deep. Presently, eelgrass populations at this site, while dense and apparently healthy, remain restricted to depths ≤ 1.5 m (MLW) (e.g., Orth and Moore 1988; Orth et al. 1992, 1993).

The final issue addressed by SAV modeling during this work period dealt with tidal variations in depth as a third source of high frequency variability associated with eelgrass habitat. In conjunction with the light attenuation coefficient, this is another factor influencing the light available at the leaf surfaces of eelgrass. To examine the potential impacts, relative to the previous model cases with constant depth, a series of simulations were run using the conditions at Guinea Marsh, which, of the three sites used for simulation analysis, has the most productive extant eelgrass population. Each case was run using a stochastically varying K_d based on seasonally grouped statistics as described above (Table 4).

In the face of stochastically varying PAR and K_d and tidally varying depth, eelgrass populations were not stable for plant canopies deeper than 1.0 m below the surface (Figure 17). This is 0.5 m shallower than in simulations lacking tidal variability (Figure 16); a significant difference with respect to both the restoration targets and the potential areal extent of SAV habitat. Again, assuming a ca. 0.30-m canopy height for eelgrass populations in the lower Chesapeake Bay, conditions in this model scenario would meet the Tier II target for maintaining or restoring eelgrass to a depth of 1.0 m, but not the Tier III target of 2.0 m. Thus, for the same water quality conditions, the addition of tidal variability to the model led to a greater light sensitivity for eelgrass. Furthermore, these scenarios suggest that the adopted habitat restoration criterion for K_d may not be sufficient to achieve the Tier III goal.

Key Simulation Findings

The results of several phases of polyhaline SAV modeling, focusing on eelgrass as the dominant species, have provided strong support for the efficacy of such models in addressing management oriented questions regarding water quality and habitat restoration. While future work will enhance and integrate SAV models into larger ecosystem models capable of addressing a broader scope of living resource questions (e.g., relating water quality and habitat restoration goals

Polyhaline SAV Model
Using stochastic PAR and K_d with fixed depth

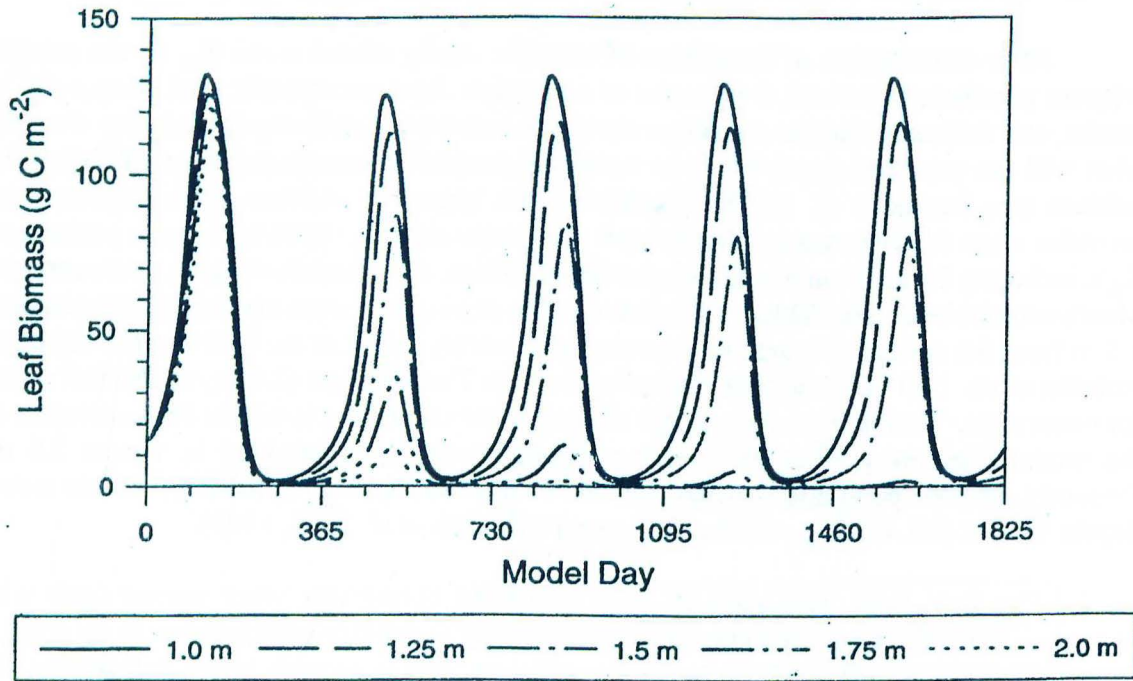


Figure 17. Predicted eelgrass leaf biomass with increasing water depth (mean low water) using stochastically varying light (PAR) and light attenuation coefficients (K_d), the latter derived from data at Guinea Marsh, York River, Virginia.

Polyhaline SAV Model

Using stochastic PAR and K_d with tidally-varying depth

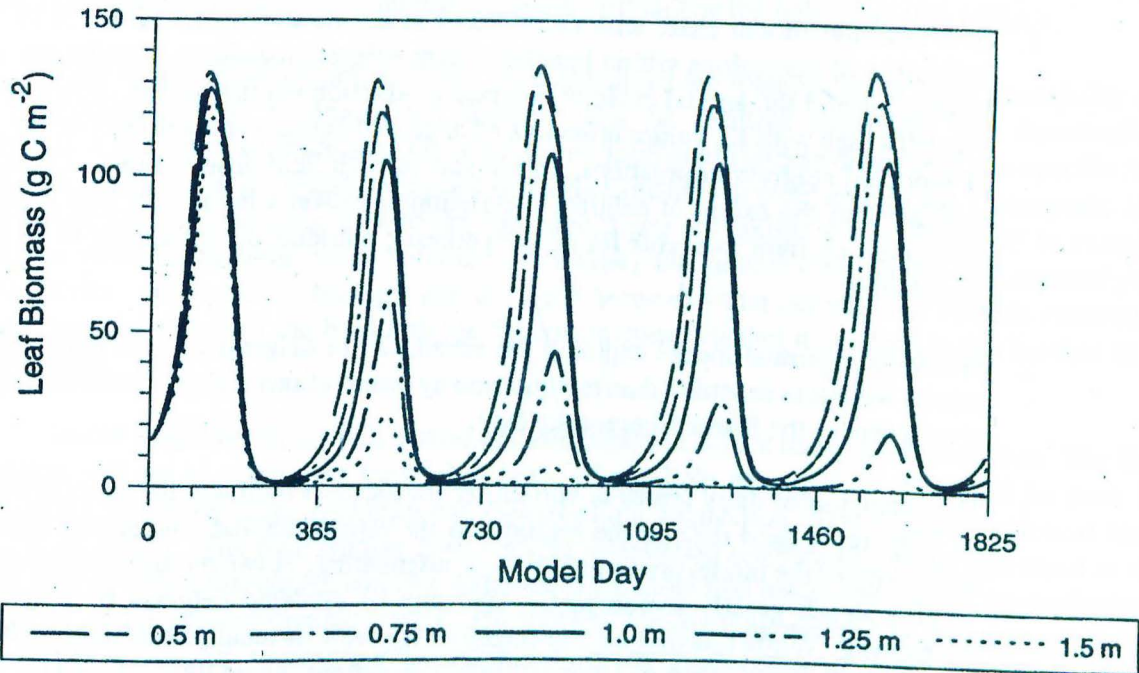


Figure 18. Predicted eelgrass leaf biomass with tidal fluctuations at five mean low water depths using stochastically varying light (PAR) and light attenuation coefficients (K_d), the latter derived from data at Guinea Marsh, York River, Virginia.

to the productivity of living resources), several key findings have arisen from the standalone SAV models presented here. These are summarized here:

1. In a nominal model case, with depth fixed at 1.0 m and K_d fixed at 1.0 m^{-1} , both indicative of conditions within healthy, extant eelgrass meadows, simulations yielded biomasses and the annual cycle of eelgrass production similar to that observed in situ. Further cases with K_d values reflective of degraded habitats showed decline and failure of modeled eelgrass populations, also in accord with field observations. These results indicate that the extent of eelgrass populations in lower Chesapeake Bay is primarily controlled by light available for photosynthesis; nutrients do not appear to be limiting production.
2. While the nominal model captured the sensitivity of eelgrass to light attenuation, the results were less sensitive than is suggested by observations and the distribution patterns of eelgrass in the lower Chesapeake Bay.
3. The inclusion of daily scales of variability for incident light and for attenuation, based on long term field data in the region and its various habitats, increased slightly the sensitivity of the model predictions to light attenuation. However, the addition of tidally varying depth greatly increased the sensitivity of modeled eelgrass biomass to light attenuation. While this may be non-intuitive, it should be recalled that light is attenuated exponentially with depth and that eelgrass populations, while light-limited for much of the year and experience severe thermal stress during the summer, such that the additional stress of low light when high tide is coincident with periods of daily photosynthetic maxima, eelgrass production is sufficiently depressed to lead to loss in biomass. This sensitivity, detected in the modeling effort, may indicate that habitat restoration targets with respect to water clarity may not be sufficient to allow the restoration of eelgrass beds to the 2.0 m Tier III target.
4. Modeling the interactions of eutrophication on epiphyte density, which in turn affects the photosynthesis of SAV, and the control of these epiphytes by grazing activity, indicates that, at present, grazers are sufficiently cropping epiphytes within, though the impact of eutrophication on phytoplankton and its light attenuation was not explicitly addressed. However, any perturbations which reduce the grazer populations or increase algal (planktonic and epiphytic) growth rates will have a negative impact on SAV by allowing phytoplankton and epiphytes to accumulate to levels which would severely reduce SAV photosynthesis.

FUTURE WORK

In our simulation analyses, in corroboration of continuing studies of SAV communities and water quality, the nature of the highly variable light environment is a critical factor in SAV productivity. Often, even in flourishing SAV communities, light criteria can be near critical values for SAV productivity (Moore 1992). The submarine light climate is determined by depth, suspended particulates (including phytoplankton, non-living organic detritus, and inorganic sediments,

measured collectively as TSS), and dissolved organic matter (DOM). In terms of anthropogenic impacts, alteration in the TSS from upstream perturbations or increased nutrient loadings which lead to increased phytoplankton and epiphytic growth will reduce the light available to SAV.

Our previous simulation analyses have examined the relationship of light availability and SAV productivity by incorporating, directly, data on incident irradiance and downwelling attenuation into simulation studies. Obviously, this approach is largely dependent site-specific data and thus is limited in its ability to explain or predict features in other area of the Chesapeake Bay and its tributaries. Also, the data-driven approach limits our ability to use the model to examine complex relations between water column (microalgal) and benthic (microalgal and macrophytic) productivity. Competition for light and nutrients between water column and benthic autotrophs determines the nature of the food web for the community, whether it is based in the benthos or in the pelagial.

In our future work, we will extend our simulation efforts in three main directions. The first direction will be to extend the theoretical basis of our light description. This will be done by calculating incoming solar irradiance and downwelling attenuation from a more theoretical basis (e.g., Kirk 1983). The total diffuse downwelling attenuation coefficient can be described as the linear sum of individual coefficients for water, detritus, inorganic suspended matter, phytoplankton pigments, and dissolved organic matter. Particularly for inorganic sediments and phytoplankton, this would allow for more complex examinations of interactions of these readily-measured parameters, light availability and ecosystem processes in relation to water quality and living resources restoration goals.

The second area of fruitful expansion of our modeling efforts will be the inclusion of nutrient cycling, sediment oxygen and nutrient fluxes, and water column trophic dynamics (including phytoplankton, grazing zooplankton, and predators). Of course, a description of phytoplankton biomass will be necessary in the aforementioned description of light attenuation. Additionally, inclusion of this aspect of the SAV-littoral ecosystem will allow for complex examinations of eutrophication, specifically, the cycles of nitrogen and phosphorus. Also included will be the dynamics of dissolved oxygen, which is intimately couple in cycles of organic production, degradation, and in the fluxes between the sediment and water column.

Thirdly, we are current in the initial stages of integrating a SAV ecosystem process model with the water quality and hydrodynamic modeling efforts of colleagues at VIMS. This was discussed in the Background section, and is focused on the York River in a program called the York River Regional Ecosystem project. The coupling of physical and biological processes two (x, y , depth-integrated) and three dimensions (x, y, z) improves the ability to describe and predict the mass transport of materials important in determining water quality (nitrogen, phosphorus, suspended matter, oxygen), as well as examining the spatial scales associated with biological processes important for water quality and habitat restoration. It can also indicate potentially sensitive locations, based on water flow and sediment suspension characteristics, within the tributaries where improved water quality is crucial for the re-establishment of SAV in ways in which spatially averaged models cannot predict.

While the last of these improvements is a long term effort, because of the mathematically, theoretical and computational complexities, all three of these areas of improvement will have

significant impacts on our understanding of the complex biological interactions within shallow water aquatic SAV habitats. In concert with long term survey data and with process-oriented field and laboratory experiments, the modeling efforts can aid the identification of critical and sensitive aspects of the ecosystem (Figure 4). In addition, the coupling of complex biological processes with water quality and hydrodynamic models should refine and improve predictive and management capabilities.

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APPENDIX

PROGRAM SEAGRASS

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C
C           A SIMULATION MODEL OF ZOSTERA MARINA PHOTOSYN-
C           THESIS, RESPIRATION, MORTALITY AND LOSS DUE TO
C           GRAZING; EPIPHYTE COLONIZATION, PHOTOSYNTHESIS,
C           RESPIRATION, MORTALITY AND LOSS DUE TO GRAZING
C           AND THE ROLE OF EPIPHYTIC GRAZERS (MODELED AFTER
C           THE ISOPOD, IDOTEA BALTICA). THE MODEL IS DRIVEN
C           BY LIGHT, WATER TEMPERATURE, PAR ATTENUATION AND
C           WATER DEPTH (A FUNCTION OF A PROGRESSIVE TIDAL
C           CYCLE WITH A LUNAR PERIOD OF 30 DAYS. THE MODEL RUNS
C           ON THREE TIME SCALES: HOURS, DAYS AND YEARS.
C*****
C           DEFINE PROGRAM ENVIRONMENT (ASSIGN NAMES TO DATA STRUCTURES, ALLOCATE
C           STORAGE SPACE) AND FORMATS FOR MODEL OUTPUT
C*****
C           IMPLICIT DOUBLE PRECISION (A-H,O-Z)
C           IMPLICIT INTEGER (I-N)
C           DOUBLE PRECISION MLW, MSL
C           CHARACTER*1 PARFCT, TIDE
C           CHARACTER*32 FNAME
C           CHARACTER RUNHDR*79, SITE*2

C$LARGE:FLUX,AFLUX,PHYSICAL,TRATES !MS-Fortran-specific compiler instruction
C
C           DIMENSION FUMMY(16),X(7),DX(7),FB(12,500),SFB(12)
C           DIMENSION FLUX(16,500),SIFLUX(16),STFLUX(16),DFLUX(16),TFLUX(16)
C           DIMENSION AFLUX(16,500)
C           DIMENSION DAFLUX(9)
C           DIMENSION DARATE(2)
C           DIMENSION PHYSICAL(7,500),PHYSDAT(7)
C           DIMENSION TRATES(14,500),SRATES(14)

C           EQUIVALENCE (FUMMY(1),A0203),(X(1),X01),(SFB(1),FB0203)
C           EQUIVALENCE (SIFLUX(1),F0203)
C           EQUIVALENCE (PHYSDAT(1),TEMP)
C           EQUIVALENCE (SRATES(1),EPLR)

C           COMMON/FIXED/A0203,G0203,A0303,G0303,A0404,G0404,A0205,G0205,
C           +A0505,G0505,A0306,G0306,A0606,G0606,A0506,G0506

C           COMMON/STATE/X01,X02,X03,X04,X05,X06,X07,DT,NDPRT,JYRMX

C           COMMON/TIME/ITIME(500)

C           COMMON/FLUX/F0203,F0302,F0304,F0402,F0307,F0205,F0502,F0507,
C           +F0306,F0506,F0602,F0607,F0600,F0006,F0102,F0201

C           COMMON/FB/FB0203,FB0303,FB0404,TF0203,FB0205,FB0505,TF0205,
C           +FB0306,TF0306,FB0506,TF0506,FB0606

C           COMMON/PHYSICAL/TEMP,Z,PP,PARD,PARSEC,PARZ,AKZ

C           COMMON/RATES/EPLR,PARVP,VPIK,PM03,T0203,R0302,EPIK,PM05,
C           +T0205,R0502,DIFFL,T0306,T0506,R0602

C           write(6,'(///)')
C           WRITE(6,*) ' Welcome to the Seagrass Ecosystem...'
C           write(6,*) ' PAR, T, Kd, Z --> Zostera, Epiphyta, Grazer'
C           write(6,*) ' --> CO2, Fish'
C           write(6,'(///)')
C           write(6,1000)
C           read(5,'(a79)') RUNHDR
C           1000 format(' ','Give a model run descriptive header (<80 chars)...'/
C           & ',','>')
C           WRITE(6,1001)
C           1001 FORMAT(5X,'ENTER FILENAME FOR DAILY INTEGRATED FLUX OUTPUT '\)
C           READ(5,1002)FNAME

```



```

C 1002 FORMAT(A)
      NFLX = 20
      FNAME='DFLX.PRN'
      OPEN(NFLX,FILE=FNAME,STATUS='NEW')
C      WRITE(6,1003)
C 1003 FORMAT(5X,'ENTER FILENAME FOR DAILY INTEGRATED RATE OUTPUT '\)
C      READ(*,1002)FNAME
      NRAT = 21
      FNAME='DRAT.PRN'
      OPEN(NRAT,FILE=FNAME,STATUS='NEW')
C      WRITE(*,1004)
C 1004 FORMAT(5X,'ENTER FILENAME FOR STANDING STOCK OUTPUT '\)
C      READ(*,1002)FNAME
      FNAME='SS.PRN'
      NSST = 22
      OPEN(NSST,FILE=FNAME,STATUS='NEW')
C      WRITE(*,1005)
C 1005 FORMAT(5X,'ENTER FILENAME FOR PHYSICAL DATA OUTPUT '\)
C      READ(*,1002)FNAME
      FNAME='PHYS.PRN'
      NPHY = 23
      OPEN(NPHY,FILE=FNAME,STATUS='NEW')
C      WRITE(*,1006)
C 1006 FORMAT(5X,'ENTER FILENAME FOR FLUXES AND MASS BALANCE OUTPUT '\)
C      READ(*,1002)FNAME
      FNAME='MB.PRN'
      NMBL = 24
      OPEN(NMBL,FILE=FNAME,STATUS='NEW')
C
C WRITE FILE HEADERS TO STD STOCK FILE
      WRITE(NSST,*) RUNHDR
      WRITE(NSST,200)
C
C TABULAR OUTPUT FORMATS FOR COMPARTMENT BIOMASS, DAILY AND
C SUMMED FLUXES AND FEEDBACK TERMS.
C
C OUTPUT FORMATS FOR STATE VARIABLES
200 FORMAT(3X,'DAY',5X,'CO2-AIR',5X,'CO2-H2O',5X,'PLT-LVS',5X,
+ 'PLT-R/R',5X,'EPIPHY',6X,'GRAZER',5X,'DETRITUS',5X,'EPI/PLT-LVS')
210 FORMAT(1X,I5,2X,7F12.4,5X,F7.4)
C
C OUTPUT FORMAT FOR DAILY INTEGRATED FLUXES
599 FORMAT(I5,9F10.4)
598 FORMAT(I5,2F10.5)
C
C      OUTPUT FORMATS FOR DAILY AND SUMMED FLUXES
C
229 FORMAT(3X,'MODEL PREDICTIONS: SPECIFIC FLUX RATE ',
+ '(GM-C / M2 / HOUR) AT 12:00 HOURS ON GIVEN DAY')
230 FORMAT(3X,' DAY',2X,' F0203 F0302 F0304',
+ ' F0402 F0307 F0205 F0502 F0507')
231 FORMAT(3X,' DAY',2X,' F0306 F0506 F0602',
+ ' F0607 F0600 F0006 F0102 F0201')
232 FORMAT(3X,I5,2X,8F10.4)
233 FORMAT(3X,'MODEL PREDICTIONS: CUMULATIVE FLUX ',
+ 'BY SPECIFIC PATHWAY')
C
C      OUTPUT FORMATS FOR FEEDBACK TERMS
C
239 FORMAT(3X,'MODEL PREDICTIONS: FEEDBACK CONTROL ',
+ 'VALUES AT 12:00 HOURS FOR THE GIVEN DAY')
240 FORMAT(3X,' DAY',2X,' FB0203 FB0303 FB0404',
+ ' TF0203 FB0205 FB0505')
242 FORMAT(3X,' DAY',2X,' TF0205 FB0306 TF0306',
+ ' FB0506 TF0506 FB0606')
241 FORMAT(3X,I5,2X,6F10.4)
C
C      OUTPUT FORMATS FOR COMPARTMENT MASS BALANCE
C
249 FORMAT(10X,'MODEL PREDICTIONS: ANNUAL COMPART',
+ 'MENTAL MASS BALANCE')
250 FORMAT(10X,'COMPARTMENT',10X,'SUM INPUT',10X,'SUM OUTPUT',
+ 10X,'MASS BALANCE')

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251 FORMAT(10X,'X01: CO2-AIR ',6X,F10.4,10X,F10.4,10X,F10.4)
252 FORMAT(10X,'X02: CO2-H2O ',6X,F10.4,10X,F10.4,10X,F10.4)
253 FORMAT(10X,'X03: PLANT-LVS',6X,F10.4,10X,F10.4,10X,F10.4)
254 FORMAT(10X,'X04: PLANT-R/R',6X,F10.4,10X,F10.4,10X,F10.4)
255 FORMAT(10X,'X05: EPIPHYTES',6X,F10.4,10X,F10.4,10X,F10.4)
256 FORMAT(10X,'X06: GRAZERS ',6X,F10.4,10X,F10.4,10X,F10.4)
257 FORMAT(10X,'X07: DETRITUS ',6X,F10.4,10X,F10.4,10X,F10.4)

```

OUTPUT FORMATS FOR PHYSICAL DATA (ANNUAL CYCLES)

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299 FORMAT(3X,'MODEL PREDICTIONS: PHYSICAL DATA USED ',
+'FOR FORCING FUNCTIONS')
300 FORMAT(3X,' DAY',1X,' TEMP ',',',DEPTH ',
+' DAYLT ',',',PAR/DAY ',',',PAR-INC ',
+' PAR-SUB ',',',ATT-COEF ')
301 FORMAT(9X,' (C) ',',', (M) ',',', (HRS) ',',',
+' (E/M2/DY) ',',', (E/M2/S) ',',', (UE/M2/S) ',',', (M-2) ')
302 FORMAT(3X,I5,7F10.3)

```

OUTPUT FORMATS FOR CALCULATED RATE COEFFICIENTS
UNITS ARE MG-C/MG-C/HOUR

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399 FORMAT(3X,'MODEL PREDICTIONS: PHOTOSYNTHESIS ',
+'PARAMETERS AND RATE COEFFICIENTS FOR ZOSTERA MARINA AND ',
+'EPIPHYTES')
400 FORMAT(3X,' DAY',',',ZOSTERA ',',',ZOSTERA ',
+' ZOSTERA ',',',ZOSTERA ',',',ZOSTERA ',',',5X,
+' EPIPHYTE ',',',EPIPHYTE ',',',EPIPHYTE ',',',EPIPHYTE ')
401 FORMAT(8X,' EPLR ',',',PARVP ',',',VPIK ',
+' P-MAX ',',',GPP ',',',RESP ',',',5X,' EPIK ',
+' P-MAX ',',',GPP ',',',RESP ',',',DIFFL ')
402 FORMAT(3X,I5,6F10.5,5X,5F10.5)
403 FORMAT('MODEL PREDICTIONS: RATE COEFFICIENTS ',
+'FOR THE GRAZER COMPARTMENT')
404 FORMAT(3X,' DAY',',',GRAZER ',',',GRAZER ',
+' GRAZER ')
405 FORMAT(8X,' ING:PLTS ',',',ING:EPI ',',',RESP ')
406 FORMAT(3X,I5,3F10.5)

```

GENERAL MODEL STRUCTURE

COMPARTMENTS:

```

X01 CO2-AIR
X02 CO2-WATER
X03 ZOSTERA MARINA LEAVES
X04 ZOSTERA MARINA ROOTS AND RHIZOMES
X05 ALGAL EPIPHYTES
X06 GRAZERS
X07 DETRITUS

```

FLows:

FIJ=FLOW FROM COMPARTMENT I TO J; UNITS GRAMS CARBON/M2/H

TRANSFER COEFFICIENTS:

TIJ, UIJ, OR RIJ=SPECIFIC RATE OF CARGON TRANSFER FROM COMPARTMENT I TO J; UNITS GRAM/GRAM/HOUR
>>>NOTE THAT SOME TERMS REMAIN CONSTANT THROUGHOUT A GIVEN SIMULATION WHEREAS OTHERS ARE CALCULATED WITH EACH ITERATION. EXOGENOUS VARIABLES (E.G. TEMPERATURE, LIGHT) EXERT CONTROL BY AFFECTING THESE VARIABLE RATE COEFFICIENTS.

FEEDBACK CONTROL:

FBIJ=FEEDBACK FUNCTION WHICH REDUCES FLUX FROM COMPARTMENT I TO J WHEN CONCENTRATION OF RESOURCE I BECOMES LIMITING.
FBJJ=DENSITY DEPENDENT FEEDBACK FUNCTION WHICH REDUCES FLUX TO COMPARTMENT J WHEN SPACE (NUTRIENTS, ETC.) BECOMES LIMITING;
CONSTRAINS DENSITY OF COMPARTMENT J BELOW A MAXIMUM MAINTAINABLE LIMIT

INITIALIZE FIXED PARAMETERS


```

    PI = ACOS(-1.00)
    LCB = 0
    LGP = 0
    LGM = 0

C MBM: LAHEY F77L3 FCT TO SEED/PICK A RANDOM NO.
C (MIMICKS MS-FTN, WHICH USES AN INITIAL SEED OF 1.0)
    RAND = RANDS(1.0)

    WRITE(6,3005)
    READ(5,'(F6.2)') AKD
    AKZ = AKD
    IF (AKD .EQ. 0.0) THEN
        WRITE(6,4001)
        READ(5,'(A2)') SITE
        IF (SITE .EQ. 'cb' .or. SITE .EQ. 'CB') LCB = 1
        IF (SITE .EQ. 'gp' .or. SITE .EQ. 'GP') LGP = 1
        IF (SITE .EQ. 'gm' .or. SITE .EQ. 'GM') LGM = 1
    ENDIF
    WRITE(6,3006)
    WRITE(6,3007)
    READ(5,'(F4.2)') MLW
    WRITE(6,3008)
    READ(5,'(F4.2)') TAMP
    WRITE(6,3009)
    READ(5,'(A1)') TIDE
    MSL = MLW + 0.5*TAMP
    WRITE(6,3011)
    READ(5,'(A1)') PARFCT
    WRITE(6,3012)
    READ(5,'(F5.2)') EUTROF
    WRITE(6,3013)
    READ(5,'(F5.2)') GRAZF
    IF (AKD .EQ. 0) WRITE(6,4010) SITE
    WRITE(6,3010) AKD,MLW,TAMP,TIDE
    WRITE(6,3015) PARFCT,EUTROF,GRAZF

3005 FORMAT(' ','Enter Kd (x.xx, 0. For computed values )__:')
3006 FORMAT(' ','Enter nominal depth (m MLW), tidal range (m) and',
    &' whether tidal variations',/
    &,' ',T5,'should be calculated (MSL = MLW + 0.5*range)...')
3007 FORMAT(' ',T15,'Z (m MLW)__:')
3008 FORMAT(' ',T15,'Range (m)__:')
3009 FORMAT(' ',T15,'Tide (Y|N)__:')
3010 FORMAT(' ',Kd = ',F6.2,' 1/m Z(mlw) = ',F4.2,' Range = ',F4.2
    &,' m Tide calc'd= ',A1)
3011 FORMAT(5X,'Pick a (S)mooth or (N)oisy PAR signal__:')
3012 FORMAT(5X,'Eutrophication in this model impacts epiphyte growth.',
    &/', ' ',Assign a factor (0.0-99.9, 1.0=nominal, xx.xx)__:')
3013 FORMAT(5X,'Grazing pressure impacts epiphyte biomass.',
    &/', ' ',Assign a factor (0.0-99.9, 1.0=nominal, xx.xx)__:')
3015 FORMAT(' ',PAR = ',A1,' EUTROF = ',F5.2,' GRAZF = ',F5.2)
4001 FORMAT(' ',Computed Kd's are available for 3 sites in the',/
    & ' ',York River, Virginia. Enter your choice using',/
    & ' ',the initials as indicated:',//
    & ' ',Claybank (cb) Gloucester Point (gp)
    & ' ',Guinea Marsh (gm)',/
    & ' ',
    & ' ',__:')
4010 FORMAT(/,' ',Site = ',a2)

C
C ENVIRONMENTAL EXCHANGE PARAMETERS
C
C----> T0102=0.0
C----> T0201=0.0
C----> T0600=0.0
C----> T0006=0.0
C
C ROOT/RHIZOME TRANSFER COEFFICIENTS
C
C T0304=0.17
C R0402=0.00042
C
C FIXED MORTALITY TERMS (FRACTION OF STANDING STOCK PER DAY)

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C          IDAY IS THE DAY COUNTER
C          LDAY IS THE LUNAR DAY COUNTER
C          IPLT IS THE COUNTER FOR PLOTTING INTERVAL
C          IPRT IS THE COUNTER FOR PRINTING INTERVAL
C          ISTORE IS THE COUNTER FOR STORAGE USED
C          IHR IS THE HOUR COUNTER FOR EACH DAILY ITERATION
C          JCDAY IS THE JULIAN DAY COUNTER
C
C          INITIALIZE FLUX STORAGES TO ZERO
C
C          DO 10 I=1,16
C          STFLUX(I)=0.0
10 CONTINUE
C          DO 11 I=1,16
C          SIFLUX(I)=0.0
11 CONTINUE
C
C          INITIALIZE DAILY INTEGRATION STORAGE ARRAYS TO ZERO
C
C          DO 50 I=1,9
C          DAFLUX (I)=0.0
50 CONTINUE
C          DO 60 I=1,2
C          DARATE(I)=0.0
60 CONTINUE
C
C          COMPUTE FEEDBACK DENOMINATOR TERMS FOR FIXED VALUES OF
C          ALPHA AND GAMMA FOR BOTH DONOR AND RECIPIENT CONTROLS.
C
C          NOTE:  THEY ARE CALCULATED AS THE INVERSE.....
C          NOTE:  FB0505 (I.E. EPIPHYTE SPATIAL LIMITATION) IS A
C          VARIABLE FEEDBACK DEPENDENT ON PLANT LEAF BIOMASS
C          (IMPLICITLY, SURFACE AREA AVAILABLE FOR COLONIZA-
C          TION) AND THE DATA ARE INPUT AS BIOMASS RATIOS.
C
C          CALCULATE FEEDBACK DENOMINATOR TERMS THAT REMAIN CONSTANT
C          THROUGHOUT A GIVEN RUN
C
C          D0203=1./((A0203-G0203))+.1E-15
C          D0303=1./((G0303-A0303))+.1E-15
C          D0404=1./((G0404-A0404))+.1E-15
C          D0205=1./((A0205-G0205))+.1E-15
C          D0505=1./((G0505-A0505))+.1E-15
C          D0306=1./((A0306-G0306))+.1E-15
C          D0506=1./((A0506-G0506))+.1E-15
C          D0606=1./((G0606-A0606))+.1E-15
C
C          PRINT OUT INITIAL CONDITIONS AND RESET IDAY COUNTER
C          CALCULATE INITIAL EPIPHYTE:PLANT LVS RATIO
C          RA0503=(X05/(X03+0.1E-15))
C          WRITE(NSST,210) IDAY,X,RA0503
C          IDAY=1
C
C          MODEL SIMULATION STARTS HERE
C
C          THREE DO LOOPS CONTROL SIMULATION FOR TIME OF DAY, DAY
C          OF YEAR AND NUMBER OF YEARS SIMULATED.
C
C          DO 120 JYEAR=1,JYRMX
C          DO 110 JYDAY=1,IEND
C
C          MS-Fortran random number generator:
C          CALL RANDOM( RAND )
C          MBM: LAHEY F77L3 FCT REF TO GET A PRE-SEEDED RANDOM NO. ONCE A DAY:
C          RAND = RND()
C
C          *****
C          *****          FITTED OR ASSUMED          *****
C          *****          FUNCTIONAL RELATIONSHIPS      *****
C          *****          THAT VARY DAILY (NOT W/IN A DAY) *****
C          *****
C
C          COMPUTE THE FOLLOWING ENVIRONMENTAL VARIABLES TO 'DRIVE'

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C***** AKZ, THE VERTICAL PAR ATTENUATION COEFFICIENT WAS *****
C***** TREATED AS A CONSTANT (ANNUAL AVERAGES) FOR DIFFERENT *****
C***** SIMULATION SCENARIOS IN EARLY MODEL VERSIONS. BASED *****
C***** ON YORK RIVER SHOAL DATA (1984-92), AKZ IS NOW SET *****
C***** SEASONALLY (TEMP BASED USING MOORE'S DEFINITIONS) *****
C***** FOR DIFFERENT TIMES OF THE YEAR AND REPRESENTATIVE OF *****
C***** DIFFERENT HABITATS WITHIN THE LOWER BAY (GUINEA, VIMS *****
C***** AND CLAYBANK DATA BASES). THREE SCENARIOS ARE POSSIBLE *****
C***** FOR EACH AREA: 1) SIMULATIONS USING SEASONAL MEANS; *****
C***** 2) SEASONAL MEANS + OR - 1.0 S.D.; 3) RANDOM VARIATION *****
C***** WITHIN A SEASON USING THE RANGE AND LOWER LIMIT AS FOR *****
C***** PARD (I.E. ((RANDOM# * RANGE) + LOWER LIMIT). ALL ARE *****
C***** POSSIBLE IN THE PROGRAM SECTION THAT FOLLOWS BY *****
C***** "COMMENTING" THE APPROPRIATE EXECUTABLE STATEMENTS. *****
C*****

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C*****

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C
C IF (AKD .EQ. 0.0) THEN

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```

C
C***** THE FOLLOWING SET OF STATEMENTS ARE USED TO *****
C***** SET THE ATTN.COEFF. TO EITHER SEASONAL MEANS *****
C***** OR TO SEASONAL RANGES DETERMINED BY THE LOWER *****
C***** LIMIT, MEASURED RANGE (+- ONE STAND. DEV.) AND *****
C***** AND A RANDOM NO. *****

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```

C
C IF (JCDAY.LE.85 .OR. JCDAY.GE.315) THEN
C SEASON=WINTER
C AKZ=1.596 !CLAYBANK
C AKZ=((RAND*(0.616)))+(1.288)
C ELSE IF (JCDAY.GE.86 .AND. JCDAY.LE.145) THEN
C SEASON=SPRING
C AKZ=2.247 !CLAYBANK
C AKZ=RAND*0.910 + 1.792
C ELSE IF (JCDAY.GE.146 .AND. JCDAY.LE.260) THEN
C SEASON=SUMMER
C AKZ=2.069 !CLAYBANK
C AKZ=RAND*0.804 + 1.951
C ELSE
C SEASON=FALL
C AKZ=1.604 !CLAYBANK
C AKZ=RAND*0.514 + 1.347
C END IF

```

```

C***** THE LAST METHOD FOR ESTIMATING THE ATTN. COEFF. *****
C***** IS DERIVED AS ABOVE FOR PARD USING MONTHLY MEANS *****
C***** THE RANGE IS ESTIMATED AS 2X THE STD. *****

```

```

C
C IF (JCDAY.LE.31) THEN
C AKZ = LGM*(RAND*0.50+0.78)
C & + LGP*(RAND*0.50+0.88)
C & + LCB*(RAND*0.96+1.17)
C ELSE IF (JCDAY.LE.59) THEN
C AKZ = LGM*(RAND*0.50+0.79)
C & + LGP*(RAND*0.56+0.85)
C & + LCB*(RAND*1.38+1.13)
C ELSE IF (JCDAY.LE.90) THEN
C AKZ = LGM*(RAND*0.80+0.49)
C & + LGP*(RAND*0.68+0.77)
C & + LCB*(RAND*1.16+1.09)
C ELSE IF (JCDAY.LE.120) THEN
C AKZ = LGM*(RAND*0.50+0.53)
C & + LGP*(RAND*0.50+0.64)
C & + LCB*(RAND*1.46+1.17)
C ELSE IF (JCDAY.LE.151) THEN
C AKZ = LGM*(RAND*0.66+0.58)
C & + LGP*(RAND*0.88+0.95)
C & + LCB*(RAND*1.56+1.78)
C ELSE IF (JCDAY.LE.181) THEN
C AKZ = LGM*(RAND*1.12+0.62)
C & + LGP*(RAND*0.96+1.40)
C & + LCB*(RAND*3.02+1.11)
C ELSE IF (JCDAY.LE.212) THEN
C AKZ = LGM*(RAND*1.04+0.69)
C & + LGP*(RAND*0.78+1.39)

```



```

& + LCB*(RAND*0.72+1.49)
ELSE IF (JCDAY.LE.243) THEN
AKZ = LGM*(RAND*0.72+0.78)
& + LGP*(RAND*0.70+1.12)
& + LCB*(RAND*0.76+1.53)
ELSE IF (JCDAY.LE.273) THEN
AKZ = LGM*(RAND*0.82+1.00)
& + LGP*(RAND*0.82+1.03)
& + LCB*(RAND*1.56+1.34)
ELSE IF (JCDAY.LE.304) THEN
AKZ = LGM*(RAND*0.90+0.81)
& + LGP*(RAND*0.48+0.96)
& + LCB*(RAND*1.18+1.08)
ELSE IF (JCDAY.LE.334) THEN
AKZ = LGM*(RAND*0.74+0.66)
& + LGP*(RAND*0.32+0.93)
& + LCB*(RAND*0.68+1.06)
ELSE
AKZ = LGM*(RAND*0.82+0.62)
& + LGP*(RAND*0.56+0.72)
& + LCB*(RAND*0.78+0.92)
END IF
C
ENDIF
C*****
C***** SET DO LOOP FOR WITHIN DAY ITERATIONS *****
C*****
C
DO 100 NITER=1,ITER
C
C FEEDBACK CONTROL CALCULATIONS
C RESOURCE CONTROL FB TERMS
C
FB0203=DIM(1.,(DIM(X02,G0203)*D0203))
FB0205=DIM(1.,(DIM(X02,G0205)*D0205))
FB0306=DIM(1.,(DIM(X03,G0306)*D0306))
FB0506=DIM(1.,(DIM((X05/X03),G0506)*D0506))
C
C NOTE: DIM FUNCTION PRODUCES THE POSITIVE DIFFERENCE OF TWO VALUES;
C IF X>Y, DIM(X,Y)=X-Y; ELSE DIM(X,Y)=0.
C
C SPATIAL CONTROL FB TERMS
C
FB0303=(1.-DIM(1.,DIM(X03,A0303)*D0303))
FB0404=(1.-DIM(1.,DIM(X04,A0404)*D0404))
FB0505=(1.-DIM(1.,DIM((X05/(X03+.1E-15)),A0505)*D0505))
FB0606=(1.-DIM(1.,DIM(X06,A0606)*D0606))
C
C TOTAL FEEDBACK CALCULATIONS
C CALCULATE FIJ AND FJJ PRIME VALUES FOR MULTIPLICATIVE
C INTERACTION
C
FBP0203=DIM(1.,FB0203)
FBP0205=DIM(1.,FB0205)
FBP0306=DIM(1.,FB0306)
FBP0506=DIM(1.,FB0506)
C--> FBP0303=DIM(1.,FB0303)
C--> FBP0505=DIM(1.,FB0505)
C--> FBP0606=DIM(1.,FB0606)
C
C CALCULATE REALIZED FLUX PREFERENCE VALUES
C
C COMPUTE THE DENOMINATOR TERM
C AS THE INVERSE
C
PD06=1./((P0306*(1.-FB0306)+P0506*(1.-FB0506))+.1E-15)
C
C COMPUTE FLUX PREFERENCES
C
TP0306=P0306*(1.-FB0306)*PD06
TP0506=P0506*(1.-FB0506)*PD06
C
C*****

```

```

C*****
C*****          FITTED OR ASSUMED          *****
C*****          FUNCTIONAL RELATIONSHIPS    *****
C*****          THAT VARY WITHIN A DAY     *****
C*****
C
C      COMPUTE THE FOLLOWING ENVIRONMENTAL VARIABLES TO 'DRIVE'
C      VARIOUS COMPARTMENTAL FLOWS
C
C          4. PARHR: HOURLY INCIDENT PAR (EINSTEINS/M-2/HOUR-1)
C          5. PARSEC: INSTANTANEOUS INCIDENT PAR (EINSTEINS/M-2/SEC-1)
C          7. TLAG:  TIME LAG FOR TEMPORAL PROGRESSION OF SUCCESSIVE
C                  TIDES OVER A LUNAR CYCLE (PERIOD = 30 DAYS)
C          8. Z   :  WATER DEPTH IN METERS (R; 1.0 TO 1.55) BASED
C                  ON DEPTHS TYPICAL OF Z. MARINA IN THE LOWER
C                  CHESAPEAKE BAY.
C          9. PARZ:  SUBMARINE IRRADIANCE AT DEPTH Z
C                  (MICROEINSTEINS/M-2/SEC-1)
C
C      TLAG=LDAY*0.842105263
C      Z=MLW
C      IF (TIDE .EQ. 'Y' .OR. TIDE .EQ. 'y')
C      &   Z = MSL + 0.5*TAMP*COS(2.*PI*(IHR-TLAG)/12.)
C
C      EVALUATE PARSEC AND PARZ ONLY DURING DAYLIGHT PERIOD
C
C      DAYLT=DIM(COS((2.*PI*(IHR-12))/(PP*2.)),0.0)
C      IF(DAYLT.GT.0.0) THEN
C          PARHR=(PARD/(0.63662*PP))*DAYLT
C          PARSEC=277.78*PARHR
C          PARZ=EXP((-AKZ*Z)+LOG(PARSEC))
C      ELSE
C          PARHR=0.0
C          PARSEC=0.0
C          PARZ=0.0
C      ENDIF
C
C*****
C      COMPUTE VARIABLE FLUX COEFFICIENTS AS
C      DRIVEN BY THE ABOVE FITTED OR ASSUMED
C      FUNCTIONAL RELATIONSHIPS.
C*****
C
C      VASCULAR PLANT PHOTOSYNTHESIS (T0203) AS A FUNCTION OF
C      TEMP AND PARZ WHICH INCLUDES THE EFFECTS OF WATER COLUMN
C      PAR ATTENUATION AND EPIPHYTE GROWTH
C
C      EPLR=(1.-DIM(1.,(SQRT(DIM((X05/X03),0.1)/(3.0-0.1)))))*0.5
C      PARVP=DIM((PARZ-(PARZ*0.75*EPLR)),0.0)
C      PM03=((0.000162*TEMP)+.0041)*
C      +(1.-DIM(TEMP,25.)/(35.-25.))
C      VPIK=((15220.78*PM03)-31.86)
C      DIFFL=DIM(1.0,(0.3*((DIM((X05/(X03+.1E-15)),.1))/(3.-.1))))
C      T0203=DIM((PM03*(PARVP/(VPIK+PARVP+1.E-15))),0.0)*DIFFL
C
C      PARVP= PAR REACHING THE PLANT CANOPY
C      VPIK=  MICHAELIS-MENTEN HALF SATURATION COEFF.
C            DERIVED AS FUNCTION OF PMAX (PENHALE 1977)
C      PM03=  THE PREDICTED P-MAX AT GIVEN TEMP AND PARZ.
C            INCREASES LINEARLY WITH TEMPERATURE TO 25 C;
C            ABOVE 25C, PMAX DECLINES LINEARLY, SUCH THAT AT
C            30 C, PMAX=.5 (MAXIMUM).
C      EPLR=  PAR ATTENUATION DUE TO EPIPHYTE GROWTH, ESTIMATED
C            AS A FUNCTION OF THE SPATIAL FEEDBACK TERM, FB0505.
C            AS THE RATIO OF EPIPHYTE TO LEAF BIOMASS APPROACHES 3,
C            THE LIGHT REDUCTION DUE TO EPIPHYTE ATTENUATION
C            APPROACHES
C            75%. ABSORBANCE AT ANY TIME IS REDUCED BY 50% TO
C            ACCOUNT FOR THE PRESENCE OF NEW, UNEPIPHYTIZED LEAVES
C            IN THE CANOPY.
C      DIFFL= REDUCTION IN PHOTOSYNTHETIC RATE DUE TO LIMIT TO
C            BICARBONATE DIFFUSION CAUSED BY EPIPHYTE LAYER;
C            APPROACHES 30% AS RATIO OF EPIPHYTE TO LEAF BIOMASS
C            APPROACHES MAXIMUM.

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C*****
C
C      VASCULAR PLANT RESPIRATION (R0302) AS A FUNCTION OF TEMPERATURE
C      AND PHOTOSYNTHESIS.
C      R0302=((0.305*DIM(((0.0104*TEMP)+.3432)*T0203),0.0))+
C      +EXP((.1370*TEMP)-10.09))
C
C      VASCULAR PLANT LEAF MORTALITY AS A FUNCTION OF TEMPERATURE
C      AND DERIVED SUCH THAT AT 0 DEGRESS, LEAF MORTALITY IS 0.5% PER DAY
C      AND AT 32 DEGRESS, LEAF MORTALITY IS 3.0% PER DAY.
C
C      U0307=(0.0175 - 0.0125*COS(2.0*PI*IDAY/365.))/24.0
C*****
C
C      EPIPHYTE PHOTOSYNTHESIS (T0205) USING THE SAME FUNCTIONAL
C      RELATIONSHIPS AS FOR VASCULAR PLANT PHOTOSYNTHESIS.
C
C      EPIK=50.+(100.*(DIM(TEMP,10.)/(30.-10.)))
C      PM05=EUTROF*((.0003801*TEMP)*(1.0-(DIM((TEMP-25.),0.0)/
C      +(45.-25.))))
C      T0205=DIM((PM05*(PARZ/(PARZ+EPIK+.1E-15))),0.0)
C
C      EPIPHYTE RESPIRATION
C
C      NOTE: ASSUMES THE SAME FUNCTIONAL RELATIONSHIPS AS FOR
C      THE VASCULAR PLAN EXCEPT IT IS 10% THE SPECIFIC RATE
C
C      R0502=0.5*((0.5*DIM(((0.0104*TEMP)+.3432)*T0205),0.0))+
C      +EXP((.1370*TEMP)-10.09))
C*****
C      INGESTION AND RESPIRATION OF GRAZERS
C*****
C
C      T0306=GRAZF*(.00325*EXP(.0921*TEMP)/24.)*DIM(TEMP,10.)/(30.-10.)
C      T0506=GRAZF*0.805*((DIM(TEMP,1.0)/(30.-10.))/24.0)
C      R0602= (.0001046*TEMP)+.0008009
C*****
C*****
C      END CALCULATIONS FOR          *****
C      VARIABLE COEFFICIENTS        *****
C*****
C*****
C
C      COMPUTE METABOLIC CORRECTION TERMS FOR
C      SELF-CONTROLLED FEEDBACK TERMS
C
C      C0203=DIM((1.-(R0302/(T0203+.1E-15))),0.0)
C      C0205=DIM((1.-(R0502/(T0205+.1E-15))),0.0)
C      C0306=DIM((1.-(R0602/((T0306*AE0306)+.1E-15))),0.0)
C      C0506=DIM((1.-(R0602/((T0506*AE0506)+.1E-15))),0.0)
C
C      CALCULATE TOTAL MULTIPLICATIVE FEEDBACK TERMS
C
C      TF0203=DIM(1.,(FBP0203*(1.-(FB0303*C0203))))
C      TF0205=DIM(1.,(FBP0205*(1.-(FB0505*C0205))))
C      TF0306=DIM(1.,(FBP0306*(1.-(FB0606*C0306))))
C      TF0506=DIM(1.,(FBP0506*(1.-(FB0606*C0506))))
C
C      CALCULATE HOURLY FLUXES
C
C      VASCULAR PLANT: ROOT/RHIZOME EXCHANGES
C
C      F0203=T0203*X03*(1.-TF0203)
C      F0302=R0302*X03
C      F0304=T0304*DIM(F0203,F0302)*(1.-FB0404)
C      F0402=R0402*X04
C      F0307=U0307*X03*(DIM((TEMP-10.),0.0)/(30.-10.))
C
C      F0307=U0307*X03

```

C
C
C

EPIPHYTE EXCHANGES

F0205=T0205*X05*(1.-TF0205)
F0502=R0502*X05
F0507=(U0507*X05)+(DIM((X05/X03+.1E-15),0.0)*F0307)

C
C
C

GRAZER EXCHANGES

F0306=TP0306*T0306*X06*(1.-TF0306)
F0506=TP0506*T0506*X06*(1.-TF0506)
F0602=R0602*X06
F0607=(U0607*X06)+(F0306*(1.-AE0306))+(F0506*(1.-AE0506))

C
C
C

ENVIRONMENTAL EXCHANGES

F0102=0.0
F0201=0.0

C
C
C
C
C
C

LOSS OF GRAZERS TO FISH PREDATION INCORPORATED USING
PRED=SEASONAL PATTERN OF FISH ABUNDANCE IN ZOSTERA MEADOWS OF
NORTH CAROLINA (R: 0 TO 1 GRAMS CARBON/M2), AND
DR=DAILY RATION (%/DAY) OF PREDATORY FISH BASED ON TEMPERATURE.

PRED=0.5-(0.5*COS((2*PI*IDAY)/365))
DR=0.3*(DIM((-13.1+(2.29*TEMP)-(0.032*(TEMP**2))),0.0)/100./24.)*
+(DIM(1.0,(DIM(0.2,X06)/(0.2-0.1))))
F0600=DR*PRED
F0006=0.0

C
C
C

ASSIGN FLUX VALUES FOR EACH ITERATION

DFLUX(1)=F0203
DFLUX(2)=F0302
DFLUX(3)=F0304
DFLUX(4)=F0402
DFLUX(5)=F0307
DFLUX(6)=F0205
DFLUX(7)=F0502
DFLUX(8)=F0507
DFLUX(9)=F0306
DFLUX(10)=F0506
DFLUX(11)=F0602
DFLUX(12)=F0607
DFLUX(13)=F0600
DFLUX(14)=F0006
DFLUX(15)=F0102
DFLUX(16)=F0201

C
C
C

SUM AND STORE FLUX VALUES IN STFLUX(I)

DO 12 I=1,16
TFLUX(I)=(DFLUX(I)*DT)+STFLUX(I)
12 CONTINUE
DO 13 I=1,16
STFLUX(I)=TFLUX(I)
13 CONTINUE

C
C
C

SUM HOURLY FLUXES IN DAFLUX(I)

DO 51 I=1,9
DAFLUX(I)=DAFLUX(I)+(DFLUX(I)*DT)
51 CONTINUE

C
C
C

SUM HOURLY RATES IN DARATE(I)

DARATE(1)=DARATE(1)+(T0203*(1.-TF0203))
DARATE(2)=DARATE(2)+R0302

C
C
C

COMPUTE ITERATIVE COMPARTMENTAL CHANGES

DX(1)=F0201-F0102
DX(2)=0.0


```

DX(3)=F0203-(F0302+F0304+F0307+F0306)
DX(4)=F0304-F0402
DX(5)=F0205-(F0502+F0507+F0506)
DX(6)=(F0306+F0506+F0006)-(F0602+F0607+F0600)
DX(7)=(F0307+F0507+F0607)
C
C
C   CALCULATE STATE VARIABLE CHANGE
C
C   DO 14 I=1,7
C   X(I)=DIM((X(I)+DX(I)*DT),0.0)
14 CONTINUE
C
C   CALCULATE THE RATIO BETWEEN EPIPHYTE AND PLANT BIOMASS
C
C   RA0503=(X(5)/(X(3)+0.1E-15))
C
C   INCREMENT DAY-HOUR COUNTER (IHR)
C
C   IHR=IHR+DT
C   IF (IHR.GT.24) THEN
C   IHR=1
C   ELSE
C   IHR=IHR
C   ENDIF
C
C 100 CONTINUE
C
C *****
C ***** END DAILY LOOP *****
C *****
C
C   INCREMENT PRINTER AND DAY COUNTERS
C
C   IPRT=IPRT+1
C   IDAY=IDAY+1
C   LDAY=LDAY+1
C   JCDAY=JCDAY+1
C
C   RESET DAY-HOUR COUNTER
C
C   IHR=12
C
C   CHECK LUNAR DAY COUNTER FOR COMPLETION OF LUNAR CYCLE
C
C   IF (LDAY.GT.30) THEN
C   LDAY=0
C   ELSE
C   LDAY=LDAY
C   ENDIF
C
C   CHECK THE PRINT/STORAGE INTERVAL COUNTER
C
C   IF (ISTORE.GT.ISTMAX) THEN
C   ISTORE=ISTMAX
C   GO TO 21
C   ELSE
C   ISTORE=ISTORE
C   ENDIF
C
C   IF (IPRT.LT.NDPRT) GO TO 21
C
C IF IPRT = NDPRT PRINT OUT STATE VARIABLE STANDING STOCKS
C   WRITE(NSST,210) IDAY,X,RA0503
C
C WRITE DAILY INTEGRATED FLUXES TO FILE -- UNITS G C / M2 / D
C   WRITE(NFLX,599) IDAY,DAFLUX
C
C WRITE DAILY INTEGRATED RATES TO FILE -- UNITS G C / G C / DAY
C   WRITE(NRAT,598) IDAY,DARATE
C
C   RESET THE PRINTING INTERVAL TO ZERO
C

```

```

IPRT=0
C
20 CONTINUE
C
C      INCREMENT THE STORAGE COUNTER FOR ALL OTHER OUTPUTS
C
C      ITIME(ISTORE)=IDAY
C
C      STORE THE FEEDBACK VALUES
C
C      DO 16 I=1,12
C      FB(I,ISTORE)=SFB(I)
16 CONTINUE
C
C      STORE THE DAILY FLUX VALUES
C
C      DO 17 I=1,16
C      FLUX(I,ISTORE)=SIFLUX(I)
17 CONTINUE
C
C      STORE THE ACCUMULATED (SUMMED) FLUX VALUES
C
C      DO 18 I=1,16
C      AFLUX(I,ISTORE)=STFLUX(I)
18 CONTINUE
C
C      STORE PHYSICAL DATA FOR TABLUAR OUTPUT
C
C      DO 19 I=1,7
C      PHYSICAL(I,ISTORE)=PHYSDAT(I)
19 CONTINUE
C
C      STORE CALCULATED RATE COEFFICIENTS FOR TABULAR OUTPUT
C
C      DO 199 I=1,14
C      TRATES(I,ISTORE)=SRATES(I)
199 CONTINUE
C
C      ADVANCE THE STORAGE COUNTER AND RESET THE PLOTTING
C      COUNTER
C
C      ISTORE=ISTORE+1
C
C      21 CONTINUE
C
C      RESET THE DAILY FLUX STORAGE TO ZERO
C
C      DO 53 I=1,9
C      DAFLUX(I)=0.0
53 CONTINUE
C
C      RESET THE DAILY RATE STORAGE TO ZERO
C
C      DO 61 I=1,2
C      DARATE(I)=0.0
61 CONTINUE
C
110 CONTINUE
C
C*****
C*****      END ONE YEAR LOOP      *****
C*****
C
C      CALCULATE ANNUAL MASS BALANCE FOR EACH COMPARTMENT
C      WHERE:      SX__I = SUM INPUTS
C                 SX__O = SUM OUTPUTS
C                 SDX__ = MASS BALANCE
C
C      SX01I=STFLUX(16)
C      SX01O=STFLUX(15)
C      SDX01=SX01I-SX01O
C      SX02I=STFLUX(15)+STFLUX(2)+STFLUX(4)+STFLUX(7)+STFLUX(11)

```



```

SX020=STFLUX(16)
SDX02=SX02I-SX020
SX03I=STFLUX(1)
SX030=STFLUX(2)+STFLUX(3)+STFLUX(5)+STFLUX(9)
SDX03=SX03I-SX030
SX04I=STFLUX(3)
SX040=STFLUX(4)
SDX04=SX04I-SX040
SX05I=STFLUX(6)
SX050=STFLUX(7)+STFLUX(8)+STFLUX(10)
SDX05=SX05I-SX050
SX06I=STFLUX(9)+STFLUX(10)+STFLUX(14)
SX060=STFLUX(11)+STFLUX(12)+STFLUX(13)
SDX06=SX06I-SX060
SX07I=STFLUX(5)+STFLUX(8)+STFLUX(12)
SX070=0
SDX07=SX07I-SX070

```

C
C
C
C

```

      BEGIN NEXT YEAR SIMULATION AND INCREMENT YEAR COUNTER
      AND RESET JULIAN DAY COUNTER (JCDAY)

```

```

IYEAR=IYEAR+1
JCDAY=1

```

C

```

120 CONTINUE
22  CONTINUE
23  CONTINUE

```

C
C
C
C

```

      OUTPUT ANNUAL FLOWS, FEEDBACK AND COMPARTMENTAL DYNAMICS
      AND PHYSICAL DATA IN TABULAR FORMATS

```

C

```

OUTPUT TABLE:  PHYSICAL DATA

```

```

      WRITE(NPHY,299)
      WRITE(NPHY,300)
      WRITE(NPHY,301)
      WRITE(NPHY,302) (ITIME(J), (PHYSICAL(I,J), I=1,7), J=1, ISTORE)

```

C

```

OUTPUT TABLE:  CALCULATED RATE COEFFICIENTS

```

```

      WRITE(NMBL,399)
      WRITE(NMBL,400)
      WRITE(NMBL,401)
      WRITE(NMBL,402) (ITIME(J), (TRATES(I,J), I=1,11), J=1, ISTORE)
      WRITE(NMBL,403)
      WRITE(NMBL,404)
      WRITE(NMBL,405)
      WRITE(NMBL,406) (ITIME(J), (TRATES(I,J), I=12,14), J=1, ISTORE)

```

C

```

OUTPUT TABLE:  FLUXES INCREMENTED BY DAY

```

```

      WRITE(NMBL,229)
      WRITE(NMBL,230)
      WRITE(NMBL,232) (ITIME(J), (FLUX(I,J), I=1,8), J=1, ISTORE)
      WRITE(NMBL,229)
      WRITE(NMBL,231)
      WRITE(NMBL,232) (ITIME(J), (FLUX(I,J), I=9,16), J=1, ISTORE)

```

C

```

OUTPUT TABLE:  FLUXES SUMMED OVER THE SIMULATION PERIOD

```

```

      WRITE(NMBL,233)
      WRITE(NMBL,230)
      WRITE(NMBL,232) (ITIME(J), (AFLUX(I,J), I=1,8), J=1, ISTORE)
      WRITE(NMBL,233)
      WRITE(NMBL,231)
      WRITE(NMBL,232) (ITIME(J), (AFLUX(I,J), I=9,16), J=1, ISTORE)

```

C

```

OUTPUT TABLE:  FEEDBACK TERMS

```

```

      WRITE(NMBL,239)
      WRITE(NMBL,240)
      WRITE(NMBL,241) (ITIME(J), (FB(I,J), I=1,6), J=1, ISTORE)
      WRITE(NMBL,239)
      WRITE(NMBL,242)
      WRITE(NMBL,241) (ITIME(J), (FB(I,J), I=7,12), J=1, ISTORE)

```

C

```

OUTPUT TABLE:  COMPARTMENTAL ANNUAL MASS BALANCE

```

```

      WRITE(NMBL,249)

```

```
WRITE (NMBL, 250)
WRITE (NMBL, 251) SX01I, SX01O, SDX01
WRITE (NMBL, 252) SX02I, SX02O, SDX02
WRITE (NMBL, 253) SX03I, SX03O, SDX03
WRITE (NMBL, 254) SX04I, SX04O, SDX04
WRITE (NMBL, 255) SX05I, SX05O, SDX05
WRITE (NMBL, 256) SX06I, SX06O, SDX06
WRITE (NMBL, 257) SX07I, SX07O, SDX07
```

C

```
STOP
END
```


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