# ECOLOGY of the ANTARCTIC SEA ICE ZONE EASIZ



# WATER COLUMN PROCESSES

**Comparative Plankton Ecology** 

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## ANTARCTIC SEA ICE ZONE Productivity long-recognized



Sir Alastir Hardy, Great Waters



- 1. HOW DO VARIATIONS IN SEA ICE EXTENT & DURATION INFLUENCE WATER COLUMN ECOLOGY & BIOGEOCHEMISTRY?
  - -- primary production & sedimentation in LTER study region along West Antarctic Peninsula
- 2. HOW DOES PHYTOPLANKTON COMMUNITY STRUCTURE INFLUENCE BIOGEOCHEMICAL PROCESSES?
  - -- structure of carbon exchanges in Ross Sea and WAP foodwebs.
- 3. EMPHASIS ON LARGE-SCALE PROCESSES AND INTERANNUAL VARIATIONS

#### PALMER, ANTARCTICA Long Term Ecological Research Program (LTER)

- Part of US LTER Network of 26 sites
- 1991 present
- Regional (200,000 km<sup>2</sup>) and local sampling
- Focus on sea ice dynamics, water column processes and apex predators
- Physics, remote sensing and ice, microbial biogeochemistry, sedimentation, primary production, krill, penguins



#### **1. SEA ICE VARIABILITY AND WATER COLUM PROCESSES**

- Sea ice extent, duration and retreat
- Primary production
- Sedimentation integrator of water column processes and link to benthos
- Relationships among these processes

#### LTER Sampling Grid



~100 stations surface to bottom grid sampled every January, 1991-2004

LTER sample grid Palmer Station

Sediment Trap

## Measuring and characterizing sea ice:



Solid lines: Observations Dotted lines: 1991-01 mean <u>Extent</u> (black lines): the area enclosed by the 15% sea ice concentration contour – includes open water in the SIZ.

<u>Area</u> (red): the area covered by sea ice concentrations > 15% within the total extent.

**Open water** (blue): Extent minus area.

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Data from NASA-GSFC SSMI-SSMR Time Series processed by Sharon Stammerjohn and Ray Smith for Palmer LTER

#### Palmer LTER Sea Ice Indices, 1991 - 2002



#### Sea Ice Advance and retreat in LTER grid



#### Interannual anomalies in date of ice retreat (WAP)



Symbol in upper center of contour plots is location of sediment trap mooring at 64.5S, 66W

To describe interannual variability over the decade of sampling:

Standardized anomaly =  $(X_i - \overline{X})/\overline{X}$ 

where X is annual mean over the decade.

range -1 to +1 (0 = average)

## INTERANNUAL ANOMALIES



## Sediment trap variability, 1993-2003







#### Palmer annual sedimentation rate



# CONCEPTUAL MODEL OF SIZ DYNAMICS Ice extent & retreat $\rightarrow$ bloom $\rightarrow$ sedimentation

#### PALMER STATION 1995 - 96



**250** 10 -Chl Ice Extent -0.08 60 Flux -0.82 200 8 Mean Ice 94186 150 -6 N 40 4 R guiter 100 -4 1000 km mgC m 20 50 2 0 0 -0 1996 1997 19

PALMER STATION 1996-1997

#### ICE RETREAT, PRIMARY PRODUCTION AND SEDIMENTATION

-0.08

-0.04

**ICE RETREAT** 

0.00

0.04

**ICE RETREAR & PP** 

0.12

0.08





#### SEA ICE VARIABILITY AND WATER COLUM PROCESSES

#### SUMMARY

- +/- 10-25% variations in ice translate to order of magnitude interannual variability in PP and sedimentation
- Sea ice coverage and duration (day of retreat) appear to influence annual variations in production and sedimentation
- 1999: "average" sedimentation but anomalous high relative to PP and ice retreat...a year dominated by salps not krill.
- Need more years!!

#### 2. COMPARATIVE FOOD WEB STRUCTURE EFFECTS OF PHYTOPLANKTON COMPOSITION

#### HOW DOES PHYTOPLANKTON COMMUNITY STRUCTURE INFLUENCE BIOGEOCHEMICAL PROCESSES?

#### COMPARISON OF FOODWEBS IN ROSS SEA POLYNYA AND WEST ANTARCTIC PENINSULA IN JANUARY USING AN INVERSE APPROACH

- ✓ Fates of primary production
- ✓ POC and DOC fluxes
  - **Foodweb characterization (Legendre & Rassoulzedegan)**

#### FOODWEB RECONSTRUCTION Chesapeake Bay



#### **ROSS SEA POLYNYA STUDY REGION**



ROSS SEA POLYNYA STUDIED INTENSIVELY 1994-1999 US JGOFS AESOPS PROGRAM ROAVERRS PROGRAM ITALIAN RESEARCH IN TERRA NOVA BAY PHYTOPLANKTON COMMUNITIES WELL CHARACTERIZED

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#### **ROSS SEA vs WAP**

- 1. ROSS: dominated by Phaeocystis. WAP: diatoms
- 2. ROSS: grazer biomass low (but few data for polynya) WAP: dominated by krill
- 3. BACTERIAL PRODUCTION is ~10% of annual NPP and lower during bloom.
- 4. ROSS: dominated by POC accumulation WAP: ???

#### **RECONSTRUCTING FOODWEBS: THE PROBLEM**

1. MEASUREMENTS ARE NOT ADEQUATE TO FULLY SPECIFY FLOW STRUCTURES IN FOODWEBS. MANY PATHWAYS OR COMPARTMENTS CANNOT BE MEASURED EASILY OR ROUTINELY.

2. WHAT LEVEL OF DETAIL IS NECESSARY? FUNCTIONAL GROUPS? INDIVIDUAL SPECIES?



#### AGGREGATED FOOD WEB STRUCTURE



Small phytoplankton\* Phaeocystis colonies Diatoms Protozoan grazers Microzooplankton Krill +Mesozooplankton Myctophids Penguins Bacteria Detritus DOC

\*mostly Phaeocystis unicells < 5 um.

Flows proportional to width of arrows

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#### THE INVERSE METHOD

- 1. APPROACH BORROWED FROM GEOPHYSICS (EG, SEISMOLOGY). APPLIED TO FOODWEB RECONSTRUCTION BY VEZINA AND PLATT (1986) – GROWING NUMBER OF STUDIES.
- 2. USES OBSERVATIONS AND KNOWN BIOLOGICAL CONSTRAINTS (EG, GROWTH & ASSIMILATION EFFICIENCY, BIOMASS-SPECIFIC RESPIRATION) TO GIVE BEST-FIT PATTERNS OF FLOWS IN SPECIFIED FOODWEBS.
- 3. IN ESSENCE, SOLVING SYSTEMS OF LINEAR BALANCE EQUATIONS. NONSTEADY STATES ARE ADDRESSED.
- 4. PROVIDES AND OBJECTIVE AND CONSISTENT METHOD FOR FOODWEB RECONSTRUCTION. DIFFERENT ASSUMPTIONS CAN BE TESTED.

#### INPUT DATA: ROSS OBSERVATIONS (Primary & Bacterial Production)



NPP data courtesy Walker<sup>30</sup> Smith

#### INPUT DATA: ROSS OBSERVATIONS (POC and DOC accumulation)



#### INPUT DATA: WAP OBSERVATIONS (example)

Observations taken from grid stations within Adelie penguin foraging region determined with remote transmitters on birds.



INPUT DATA: WAP OBSERVATIONS (example)



### WAP 1996 GPP = 1200 ROSS SEA 96 GPP = 550





#### **RESULTS: DOC and POC PRODUCTION**



**RESULTS: FOODWEB CLASSIFICATION** *after Legendre and Rassoulzedegan, 1996* 





Key Structuring Flows In the foodweb:

- 1. Large Phytoplankton Production
- 2. Consumption and export via large grazers & predators (F<sub>T</sub>)

Flows normalized to total NPP





Key Structuring Flows In the foodweb:

- 1. Large Phytoplankton Production (P<sub>L</sub>)
- 2. Consumption and export via large grazers & predators (F<sub>T</sub>)
- Ungrazed, sinking phytoplankton (D<sub>T</sub>)
- 4. Recycled flows via detritus and DOC =  $1-(F_T + D_T)$

Flows normalized to total NPP

Legendre & Rassoulzedegan Foodweb models	$P_L/P_T$	R <sub>T</sub> /P <sub>T</sub>	F <sub>T</sub> /Ρ <sub>T</sub>	D <sub>T</sub> /P <sub>T</sub>
Sinking of ungrazed cells	1.00	0	0	1.00
Herbivorous foodweb	0.80	0.30	0.60	0.10
Multivorous foodweb	0.35	0.60	0.30	0.10
Microbial foodweb	0.10	0.80	0.20	0
Microbial Loop	0	1.00	0	0
INVERSE RESULTS				1
North Atlantic Bloom	0.50	0.78	0.04	0.18
WAP 1996	0.67	0.69	0.19	0.11
ROSS SEA 1996	0.54	0.47	0.02	0.50

#### COMPARATIVE FOOD WEB STRUCTURE SUMMARY

- 1. PLAUSIBLE FOODWEBS, CONSISTENT WITH OBSERVATIONS, CAN BE RECOVERED FOR BOTH AREAS.
- 2. BOTH REGIONS DOMINATED BY LARGE PHYTOPLANKTON BUT TAXON DIFFERENCES LEAD TO CONTRASTS IN GRAZER ABUNDANCE AND IMPACTS
- 3. WAP: GRAZER-DOMINATED SYSTEM ROSS: DETRITAL-DOMINATED
- 4. MOST DETRITAL FLOW ORIGINATES FROM UNGRAZED PHYTOPLANKTON
- 5. FOODWEB STRUCTURE DOMINATED BY LARGE PHYTOPLANKTON BUT TEND TOWARDS MICROBIAL FOODWEB STRUCTURE – EVEN WITH LOW BP AND DOC.

#### **FINAL THOUGHTS**

- 1. IMPORTANCE OF SEA ICE IN STRUCTURING SYSTEMS
- 2. NEED TO WORK LONG TERM AT REGIONAL SCALES
- 5. NEED FOR MODELS TO INTEGRATE SPARSE OBSERVATIONS

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PALMER MEAN SEA ICE EXTENT 1979-2003



#### Open water in ice margin and primary production More open ice margin, less PP



## Sea Ice Area and Day of Retreat larger area – later retreat

