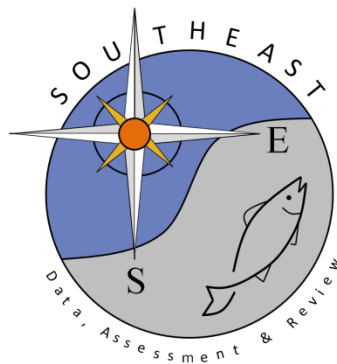


Zooplankton estimates for 2025 ERP Benchmark

M. Celestino, D. Chagaris and A. Buchheister

SEDAR102-RW-06

25 July 2025



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SEDAR 102 WP-06: Zooplankton estimates for 2025 ERP Benchmark

M Celestino, D Chagaris, A Buchheister

2025-05-16

Zooplankton data background & early discussions

For the present assessment we sought to develop a time series of zooplankton abundance to which the NWACS models could be fit. A subgroup of the ERP-WG formed to review existing ecological indicator datasets (e.g., State of the Ecosystem reports, regional surveys, etc.) to determine whether indicators could be useful to this end. The outcome of the subgroup's work suggested that indicator data from existing sources could be fed directly into one or more of our ERP models. Review of NMFS (2023) by the subgroup suggested that the NMFS's EcoMon program could be a rich source of data for the ERP assessment.

While NMFS (2023) does generate specific zooplankton indices of abundance, early discussions with NMFS (H Walsh; harvey.walsh@noaa.gov) indicated they routinely generate unique custom zooplankton indices of abundance for stock assessments and that they would be happy to do the same for us. Indices of abundance can be provided as either number per 100 m³ or number per 10 m², for any number of zooplankton categories we would like (only number per 10 m² were of interest to us). H Walsh sent us a spreadsheet with the species composition of their current categories and indicated that we can, for example, rearrange species membership, or change the number of categories from however many they have to just the ones in NWACS etc. Indices can be generated by ecological production unit (EPU) or coastwide (or both).

H Walsh sent us a spreadsheet with nearly all (see next bullet) taxa collected in the EcoMon survey:

- All of the taxa in the spreadsheet account for ~95% of all plankton NMFS captures in the EcoMon survey (i.e., omits the rarest 5% of taxa).
- Dave, Andre, and I met via conference call a number of times to discuss the data needs for NWACS-Full and NWACS-MICE. From Link et al. (2006), and used in SEDAR (2020), the 5 zooplankton groups we were interested in:
 - Large copepods: the V and VI copepodites stages of *Calanus finmarchicus*, *Metridia lucens*, and *Centropages typicus*. a,b
 - Small copeopods: stage I-IV copepodites of the large copepod species (i.e., stage I-IV copepodites of *Calanus finmarchicus*, *Metridia lucens*, and *Centropages typicus*) and the I-VI copepodites stages of *Centropages hamatus*, *Pseudocalanus* spp., *Temora longicornis*, *Paracalanus parvus*, *Nannocalanus minor*, and *Clausocalanus arcuicornis*. a,b

- Gelatinous zooplankton: Cnidaria (both the medusae and hydrozoans); the Ctenophores (comb jellies); the colonial Siphonophores; and the colonial Salpidae. c
- Micronekton: the largest size animals taken in plankton nets, typically having body lengths of 5-10mm or more. For this study, the micronekton group is considered inclusive of the crustacean groups: amphipoda, euphausiacea, mysidacea, and similar decapoda captured in plankton nets. Chaetognatha are also included in this group. d
- Microzooplankton = 13% of phytoplankton biomass estimate.

a Methods described in Link et al (2006): Plankton samples were collected seasonally on two types of cruises: 1) broad scale surveys dedicated to plankton where sampling was done at standard or randomly selected stations spaced approximately 8-35 km apart; and 2) trawl and dredge surveys where plankton stations were selected from a stratified random plan at locations uniformly distributed over the region. Samples were all collected with a 61 cm bongo frame fitted with a 0.333 mm mesh net towed obliquely to a maximum depth of 200 m or 5 m from the bottom and back to the surface. A digital flowmeter was suspended in the center of the bongo frame to measure volume of water filtered during the tow. Samples were then reduced to approximately 500 organisms by subsampling with a modified box splitter.

b Copepod abundance by size group was converted to biomass using the length to wet weight (W) equation given by Pearre (1980): $W = 0.08810L^{2.8514}$ (where L is from Pearre 1980).

c Gelatinous zooplankton biomass was estimated from 60 cm bongo tows with 333 mm mesh nets taken on NEFSC monitoring cruises. Mean abundance per m^3 for each station are the calculated mean of the abundance for each stratum sampled. Mean station abundance was multiplied by the sampling depth to calculate number per m^2 . Individual group biomasses were calculated using the following relationship (Reeve and Walter 1976): $\log DW = 2.65 \log L$ Where DW = dry weight (g) and L is length (mm). This relationship was established for ctenophores and is assumed to be similar enough for all other gelatinous zooplankton groups such that we use it for all these zooplankton taxa. A mean length of 1.3 mm was assumed for this calculation. Total biomass for all groups was then integrated into an annual average, summed across all gelatinous zooplankton taxa, and then converted to g wet weight per m^2 . Conversion to wet weight from dry weight was approximated from Pages (1997), with DW = 4.48% of WW.

d The mean abundance (no / 10 m^2) of micronekton were calculated distinctly for each of the groups listed above. These estimates of mean abundance were then converted to biomass based upon established size-biomass relationships. Here we assume that most micronekton were roughly equivalent to a common micronekton taxa, the amphipod Gamarrus sp. Mean abundance was converted to dry weight from the relationship established in Avery et al (1996), using a mean length of 6 mm which produced a dry weight estimate of 1.2 mg. This average weight was then multiplied by the abundance estimates to obtain a total biomass for each group. However, for the chaetognaths group, we used a dry weight of 0.026 mg. After conversions to biomass were done for each

micronekton group, the values were then converted to biomass and integrated into an annual estimate. This was then converted to g per weight per m².

- (1) Pearre, S Jr. 1980. The copepod width-weight relation and its utility in food chain research. *Can J Zool.* 58:1884-1891.

Into the weeds:

As part of our (Dave, Andre, and I) discussions, we agreed to, rather than classifying copepods as large or small based on observed size (or stage), as a simplifying measure, to instead classify copepods as large or small based on an average size of the adult animal, reasoning that diet habits are likely similar among similarly sized copepods (e.g., small copepods likely eat bacteria, while larger copepods likely eat larger algae ([link](#); e.g., herbivorous copepods vs predatory copepods). A series of Google and ChatGPT searches aided in assigning an average size to the various copepod species collected in the EcoMon survey; additionally, the spreadsheet from H Walsh had some initial large or small copepod designations, which were also used as a guide.

As was done in Link et al. (2006), chaetognaths were grouped as a separate category within the micronekton zooplankton group – they are included in the NWACS ‘micronekton’ category but have such a different assumed average weight than the other micronekton (0.026 mg vs 1.2 mg) that we thought it was appropriate to break them out (as was done for EMAX), then recombine for total micronekton.

We assigned all taxa in ZooTaxaUsed_NWACS_submitted.xlsx to one of our various NWACS categories (see above, Appendix 2 of SEDAR 2020, and Buchheister et al. 2017). There were a number of taxa that were binned into an ‘omit’ category - that either didn’t quite fit into the descriptions of NWACS categories or are classified as something other than zooplankton in our categories. To get a handle on the magnitude of this ‘omit’ category, we requested an index of abundance on the ‘omit’ category (i.e., all ‘omit’ taxa combined).

We talked about attempting to convert zooplankton biomass per m² to biovolume, and did some Google searches on conversions to that end, but in the end, decided not to use this approach (at least partly due to uncertainties in conversion factors). We also considered obtaining biovolume for each of our NWACS zooplankton groups, but learned that this is not physically possible; biovolume = laboratory displacement volume = all plankton + preservative in graduated cylinder poured through mesh cone into second cylinder. Displacement volume = the difference in volume between 1st and 2nd cylinder. That is, NMFS biovolume is calculated for all zooplankton all at once.

For the MICE model, we agreed that seasonal resolution of indices of abundance would be helpful for exploration, but this level of resolution was not necessary for NWACS-Full. Additionally, since the MICE model pools all zooplankton species into a single group (vs NWACS-Full that keeps all zooplankton groups separate), we also agreed to request biovolume of all zooplankton (seasonally, and by EPU) from NMFS.

Pteropods have been identified as an increasingly important component of the zooplankton (NMFS 2023). Pteropods were not mentioned in the EMAX (Link et al. 2006) or NWACS

model documentation (Buchheister et al. 2017) and so we believe they were omitted / not included, and so we classified them as a separate zooplankton group that could be treated separately if need be. We also suspected that the energetic value of pteropods could differ from other plankton and so keeping as a separate group seemed sensible. With some Google searches we were able to find some average size and weight information for pteropods. We thought that micronekton could reasonably be grouped with micronkton or omitted, and this is a decision point we discussed with the full ERP-WG at their Nov 2004 methods workshop and March 2025 model workshop.

Deeper into the weeds - pteropods:

From Googling around, we initially assumed pteropods that EcoMon collects hover around 5-6mm (see <outputs (2025) Feb 6th 2025> in ERP_zooplankton_data_1977to2022.xlsx); we found a citation for the average size of a pteropod in the Scotia Sea = 2.74 mm with dry weight = 0.62 mg; and we found a length weight relationship for Scotia Sea pteropods - if the relationship holds in the north western Atlantic, then a ~5.5mm pteropod might weigh around ~1.8 mg, which is about what we are assuming for micronekton (minus chaetognaths; see table below). We considered bundling pteropds with micronekton. But also wondered whether the assumptions might be a little too shaky (average size of a pteropod in NWACS, applying an Antarctic pteropod len-wt relationship to NWACS), bolstered by the omission in EMAX, as to support their continued omission from NWACS? We also wondered whether if by including pteropods, while EMAX didn't, would we need to re-configure some of the EwE inputs (P:B, Q:B, etc)? We also learned later that most plankton in EcoMon are < 1.5 mm.

Pteropod information.

Species or group	Assumed average length (mm)	Assumed average dry weight (mg)	Source of assumption
Micronekton (without chateognaths; no pteropds)	6.00	1.200	EMAX
Micronekton (chateognaths only)	4.00	0.026	EMAX
<i>Limacina antartica</i> (Scotia Sea pteropod)	2.74	0.620	Literature(a)
Generic NWACS pteropod	5.50	1.800	Literature(b)

Footnotes

Footnote	Source
(a)	Bednarsek et al. 2017
(b)	Bednarsek et al. 2012 (len-wt rel) & Google (mean size)

Size of EPU:

Units = m²; values from Brandon Beltz, brandon.beltz@noaa.gov, pers. comm. 28 October 2024.

EPU	area
SS	30,209,438,176
GOM	68,311,878,130
GB	57,307,696,043
MAB	126,862,117,858

Final 2025 benchmark methods

Small copepods:

We converted numeric abundance of small copepods to weight (mg to g) via: (sm copepod index) x $((0.0881 \cdot 2.0^{2.8514})/1000)$, assuming mean size = 2.0 mm to get final densities in the ballpark of the SEDAR (2020) estimates. Finally, we converted to g per m² via $\div 10$. Length-wt conversion from Table 4 of Pearre (1980) (wet weight predicted from prosome length).

Large copepods:

We converted numeric abundance of large copepods to weight (mg to g) via: (lg copepod index) x $((0.0881 \cdot 2.2^{2.8514})/1000)$, assuming mean size = 2.2 mm to get final densities in the ballpark of the SEDAR (2020) estimates. Finally, we converted to g per m² via $\div 10$. Length-wt conversion from Table 4 of Pearre (1980) (wet weight predicted from prosome length).

Gelatinous zooplankton

We converted numeric abundance of gelatinous zooplankton to weight (mg to g) via: (gel zoop index) x $((\exp(-1.54 + 2.65 \cdot \log(0.55)))/0.0448)/1000$, assuming mean size = 0.55 mm to get final densities in the ballpark of the SEDAR (2020) estimates. Finally, we converted to g per m² via $\div 10$. The conversion from length to dry weight was from Reeve and Walter (1976), and conversion to WW ($\div 0.0448$) was from Link et al. (2006).

Chateognaths

We converted numeric abundance of chateognaths to weight (mg to g) via: (chaeto index) x $((0.00097 \cdot 4^{2.365})/1000)$, assuming mean size = 4 mm to get final densities in the ballpark of the SEDAR (2020) estimates. We needed to convert from DW to WW via $\div 0.083$ (Table 7.1 in Link et al. 2006), then finally, converted to g per m² via $\div 10$. As a reminder, chaetognaths are micronekton, but literature suggests that they have a very different average weight than the other micronekton, so total mass was calculated separately, and subsequently combined with other micronekton.

Micronekton

We converted numeric abundance of micronekton (excluding pteropods) to weight (mg to g) via: (micronekton index) $\times$ (0.28/1000); avg weight came from the literature (0.28/1e3), converted to WW via $DW \times 9.30$ (Table 7.1 in Link et al. 2006), and finally, converted to g per m^2 via $\div 10$.

Pteropods (not included in 2025 benchmark)

Though not used, we are documenting for future explorations. We converted numeric abundance of pteropods via: (ptero index) $\times$ $(0.137 \times 1.5^{1.5005})/1000$, assuming mean size = 1.5 mm. Given the available length weight equation, this size approximated the average weight of 'other micronekton.' Finally, we converted to weight per m^2 via $\div 10$. Note: converted DW to WW assuming the same relationship as with other micronekton (i.e., $DW \times 9.30$).

Microzooplankton

We made the same assumptions as reported in Link et al. (2006):

- "...assumed that the microzooplankton were primarily composed of protozoans..."
- "Unfortunately traditional zooplankton sampling nets destroy the fragile protozoans, so we lack a monitoring database..."
- "Since we didn't have any independent data on protozoan biomass and rates on the Northeast Continental Shelf, we decided to relate the MZ biomass (in carbon units) to that of phytoplankton (in carbon units) based on Figure 3 in Caron et al. (1990) which showed a relationship (log-log) between ciliate and phytoplankton biomass. *We assumed that MZ (microzooplankton) biomass was 0.13 of the phytoplankton biomass* (emphasis added), similar to values for unfertilized North Sea mesocosms (Baretta-Bekker, 1994) and Narragansett Bay (Monaco, 1997). Given the boom and bust life history strategy of protozoans, we assumed that their annual biomass would be a relatively small fraction of the annual phytoplankton biomass."

Consequently, we are unaware of direct data sources for microzooplankton. Therefore, to estimate the biomass of microzooplankton, we looked to SEDAR (2020) and found that a density of 7 g/ m^2 was assumed. Discussions with A Buchheister suggested that the value likely came from Link et al. (2006) (which summed to 3 g/ m^2); SEDAR (2020) provides some additional information on how the EMAX value was modified for SEDAR (2020):

- "Section 3.3.1 Parameters for lower and higher trophic levels that leverage the EMAX models... We relied heavily on ecosystem models developed by the NEFSC for the Energy Modeling and Analysis exercise (EMAX) project (Link et al. 2006, 2008)... We preferentially used the original EMAX input values as our starting point, but some values were changed during the model balancing process."
- "Section 3.4. Description of Ecopath balancing procedure... Link et al (2006) had recognized that the EMAX model values for unassimilated fractions were low for several low-trophic-level groups (e.g., bacteria, *microzooplankton* (emphasis added), copepods, and micronekton); therefore, we increased the unassimilated fractions

for those groups to balance the detritus group and to bring the values more in line with general recommendations (Link et al. 2006; Christensen et al. 2008).” [from pdf page 516 of 560]

So as a starting point, combined with the details in Link et al. (2006), we used the ratio of microzooplankton to phytoplankton from SEDAR (2020; Appendix 2 parameters for the balanced NWACS ecosystem model): $\frac{7}{30} = 0.23$. In the absence of any direct data sources, the percentage of phytoplankton approach was retained.¹ Further, we used this ratio, multiplied by estimates of phytoplankton from the [GLORYS](#) data set (provided by D Chagaris) to develop the final microzooplankton time series.

Recollections from Andre via email: As best we can recall, the percentage approach wasn’t used in SEDAR (2020), but rather the EMAX value would have been used, tweaked as part of balancing and calibration, and finalized for use in SEDAR (2020). Since this value equated with 23% of phytoplankton, we are using that as a starting value for the present benchmark.

A note on the area sampled

The EcoMon samples are collected over a spatial area of 282,691.1 km², but the NWACS spatial domain is 441,000 km², which is calculated from spatial grids used in Ecospace. The 2025 NWACS spatial domain is a larger area than the survey domain used in SEDAR (2020). Since the 2025 model domain is larger than the area sampled by EcoMon, rather than use absolute biomass estimates from EcoMon as inputs to NWACS, we used the **density** estimates as the inputs to NWACS-MICE and NWACS-Full (g/m² = mt/km²). We felt comfortable applying the EcoMon densities to the full updated NWACS model domain since Ecomon samples across a broad representative spatial domain.

Research recommendations

Things that we weren’t able to accomplish for the 2025 benchmark include the items below and might be explored as part of the next benchmark:

- Estimates of uncertainty: communication with H Walsh suggested that a new pc and some associated lost files conspired to complicate generation of estimates of uncertainty.
- Pteropods: NMFS was able to generate a time series of pteropod abundance indices, but we felt that additional vetting and decision points could be explored. For example, are the energetics and biology (e.g., P:B, Q:B) of pteropods similar enough to one of our existing zooplankton groups (e.g., micronekton) to combine, or are they different enough to warrant a separate group. Additionally, more localized information on mean size and weight would be helpful (current weight is estimated

¹ We also discussed the possibility of having EwE estimate the micronekton biomass values, but due to time limitations we do not believe this was explored, but could be explored as part of the next benchmark.

using an assumed mean size and a length-weight equation from the Scotian Sea (Bednarsek et al. 2012)).

Literature cited

Bednarsek, N, GA Tarling, S Fielding, and DCE Bakker. 2012. Population dynamics and biogeochemical significance of *Limacina helicina antarctica* in the Scotia Sea (Southern Ocean). *Deep-Sea Research II* 5-60, 105-116.

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http://hjord.cbl.umces.edu/NWACS/TS_694_17_NWACS_Model_Documentation.pdf

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Reeve R and M Walter. 1976. A large scale experiment on the growth and predation potential of ctenophore populations. In: Mackie, G., ed. *Coelenterate ecology and behavior*. New York, NY: Plenum Pubcorp; pp 187-199.

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<http://sedarweb.org/sedar-69>

Appendix (code)

```
# Pared down from zoop_summary_03c-Feb6.r
# a = all rows, all columns
require(readxl)
require(reshape)

## Loading required package: reshape

require(plotrix)

## Loading required package: plotrix

require(latticeExtra)

## Loading required package: latticeExtra

## Loading required package: lattice

##
## Attaching package: 'latticeExtra'

## The following object is masked from 'package:ggplot2':
##
##      layer

require(plyr)

## Loading required package: plyr

##
## Attaching package: 'plyr'

## The following objects are masked from 'package:reshape':
##
##      rename, round_any

#new0 <- options(scipen=2)

a <- as.data.frame(read_xlsx("X:\\zooplankton\\zoopIndices\\ERP_zooplankton_data_1977to20
23_5Feb2025_v1b.xlsx",
  sheet="Data",na="NaN")) # Winter = Jan, Feb, March

b <- as.data.frame(read_xlsx("X:\\zooplankton\\zoopIndices\\ERP_zooplankton_data_1977to20
23_5Feb2025_v2b.xlsx",
  sheet="Data",na="NaN")) # winter = Dec, Jan, Feb

# compare results with winter defined as months 1:3 vs 12:2 (generally pretty similar):
cbind(
  cbind(
    meanJan=round(colMeans(a[a$Year<2023,-c(1:4)],na.rm=TRUE)),
    meanDec=round(colMeans(b[b$Year<2023,-c(1:4)],na.rm=TRUE))),

  cbind(
    minJan=round(apply(a[a$Year<2023,-c(1:4)],2,min,na.rm=TRUE)),
    minDec=round(apply(b[b$Year<2023,-c(1:4)],2,min,na.rm=TRUE))),
```

```

cbind(
maxJan=round(apply(a[a$Year<2023,-c(1:4)],2,max,na.rm=TRUE)),
maxDec=round(apply(b[b$Year<2023,-c(1:4)],2,max,na.rm=TRUE)))
)

##                meanJan meanDec minJan minDec  maxJan  maxDec
## Small copepods      236492  246568      0      0  2647596  2647596
## Large copepods      214203  211140      0      0  1587002  1587002
## Gelatinous zooplankton  68662   67595      0      0   894248   731699
## Chaetognatha         8529    8172      0      0   297363   194073
## Micronekton          29371   27809      0      0  4166377  4166377
## Pteropod            11862    9825      0      0   388950   163670
## Omit                 74285   68771      0      0  2020761  1935041

norm <- function(x) {
  x/mean(x)
}

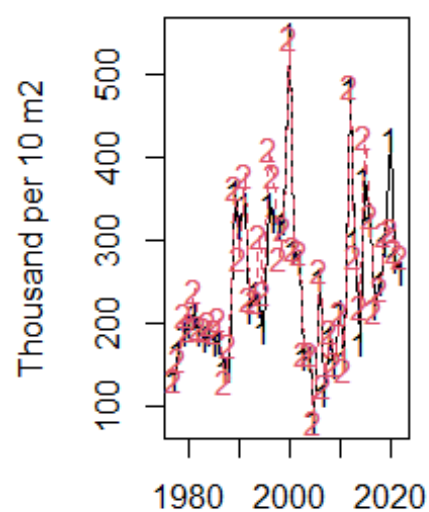
# Plot indices over time, comparing index that defines Winter starting in Jan vs Dec
# (Very similar)
op <- par(mfrow=c(1,2))
for(i in c("Small copepods","Large copepods","Gelatinous zooplankton","Chaetognatha","Micronekton","Pteropod","Omit")) {

  matplot(1977:2022,
    data.frame(winterJan= tapply(a[a$Year<2023,i],a[a$Year<2023,"Year"],mean,na.rm=TRUE),
      winterDec=tapply(b[b$Year<2023,i],b[b$Year<2023,"Year"],mean,na.rm=TRUE))/1e3,type=
"o",ylab="Thousand per 10 m2",xlab="",main=i)

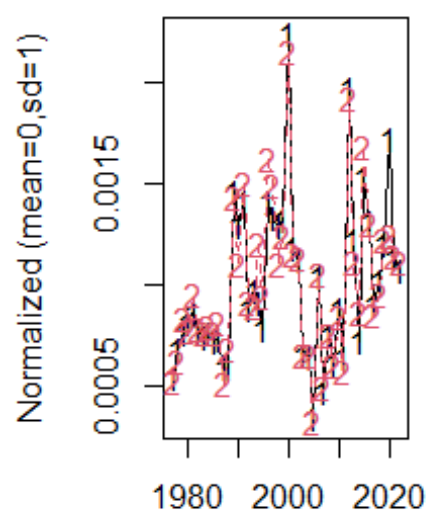
  matplot(1977:2022,
    data.frame(winterJan= norm(tapply(a[a$Year<2023,i],a[a$Year<2023,"Year"],mean,na.rm=TRUE)),
      winterDec=norm(tapply(b[b$Year<2023,i],b[b$Year<2023,"Year"],mean,na.rm=TRUE)))/1e3
,type="o",ylab="Normalized (mean=0,sd=1)",xlab="",main=i)
}

```

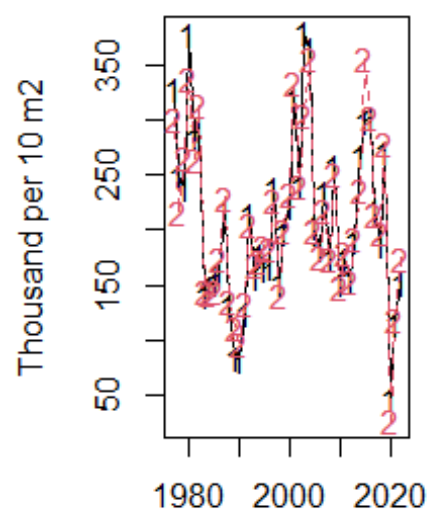
Small copepods



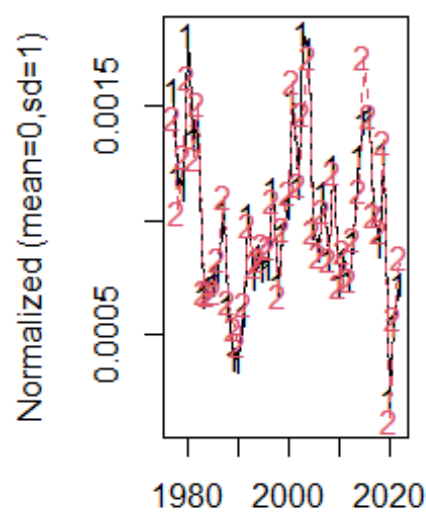
Small copepods



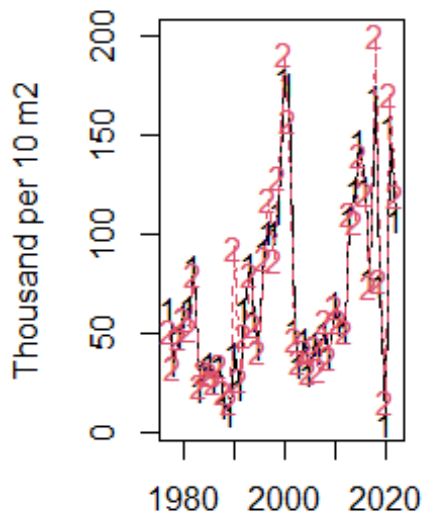
Large copepods



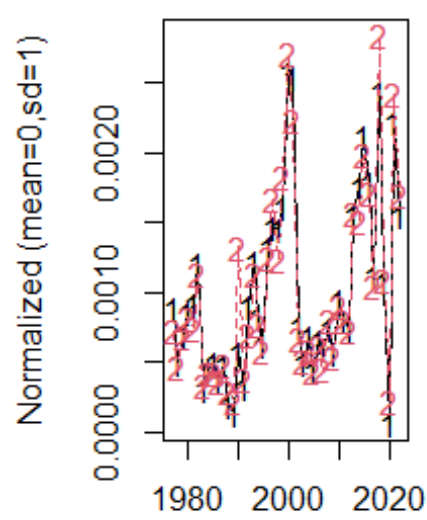
Large copepods



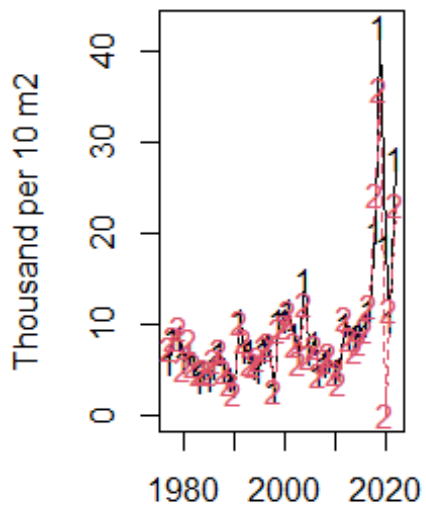
Gelatinous zooplankton



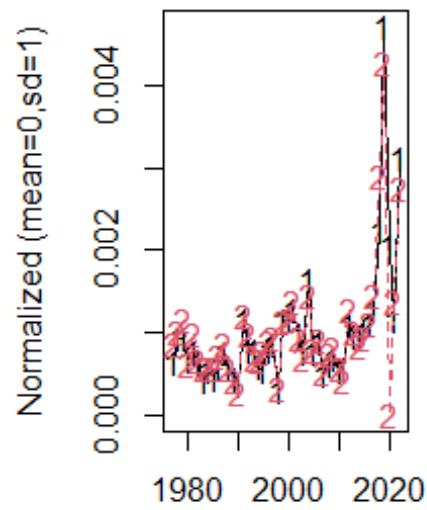
Gelatinous zooplankton



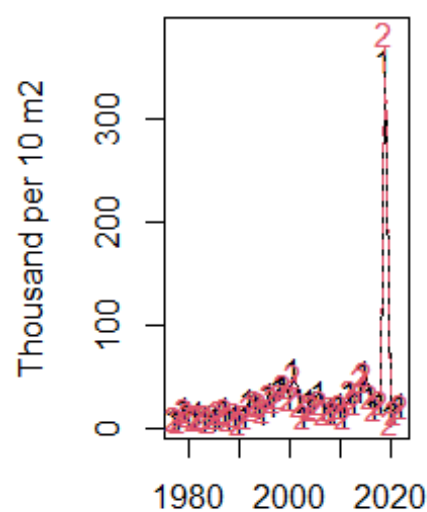
Chaetognatha



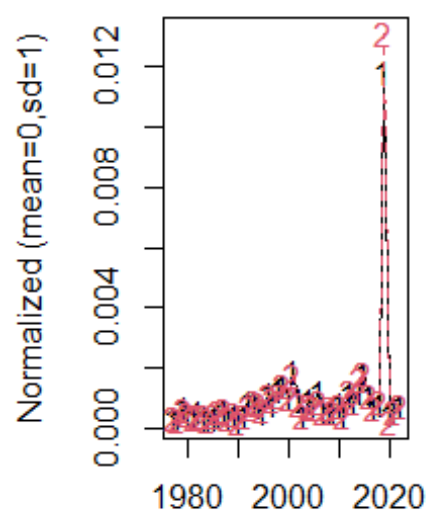
Chaetognatha



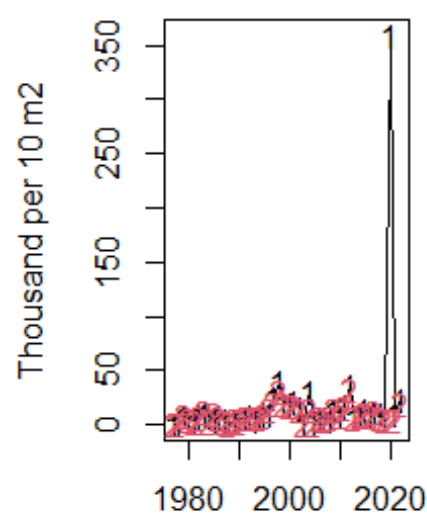
Micronekton



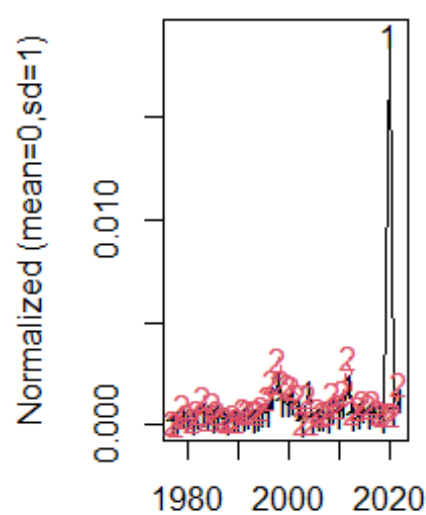
Micronekton

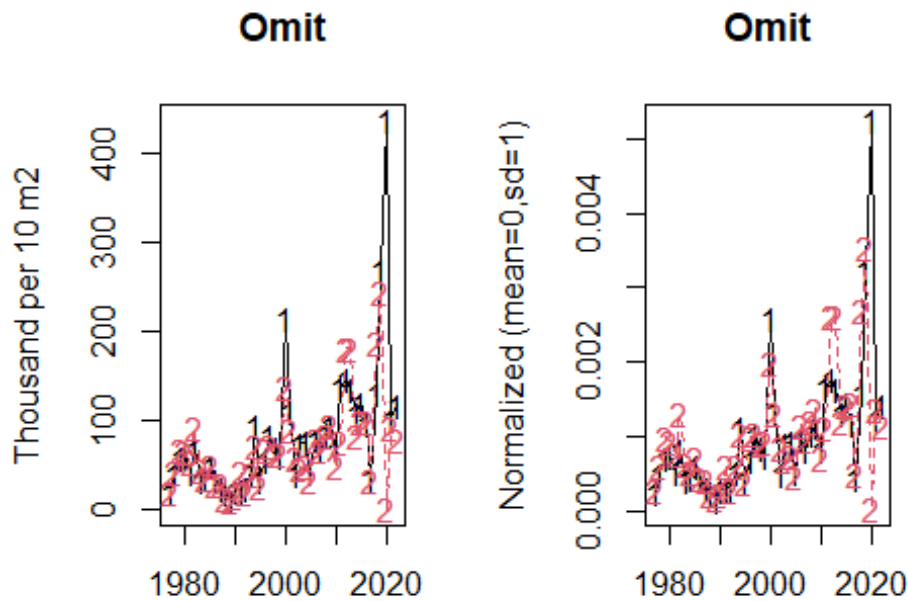


Pteropod



Pteropod





```
# Look at correlation in mean abundance as a function of how winter is defined:
# (again, very similar)
for(i in c("Small copepods", "Large copepods", "Gelatinous zooplankton", "Chaetognatha", "Micronekton", "Pteropod", "Omit")) {
  cat("\n\n\n")

  print(i)

  print(round(cor(data.frame(winterJan= tapply(a[a$Year<2023,i],a[a$Year<2023,"Year"],mean,na.rm=TRUE),
    winterDec=tapply(b[b$Year<2023,i],b[b$Year<2023,"Year"],mean,na.rm=TRUE)),method="spearman"),2))
}

##
##
##
## [1] "Small copepods"
##      winterJan winterDec
## winterJan    1.00    0.96
## winterDec    0.96    1.00
##
##
##
## [1] "Large copepods"
##      winterJan winterDec
## winterJan    1.00    0.98
## winterDec    0.98    1.00
##
##
##
## [1] "Gelatinous zooplankton"
```

```

##           winterJan winterDec
## winterJan      1.00      0.96
## winterDec      0.96      1.00
##
##
## [1] "Chaetognatha"
##           winterJan winterDec
## winterJan      1.00      0.85
## winterDec      0.85      1.00
##
##
## [1] "Micronekton"
##           winterJan winterDec
## winterJan      1.00      0.96
## winterDec      0.96      1.00
##
##
## [1] "Pteropod"
##           winterJan winterDec
## winterJan      1.00      0.84
## winterDec      0.84      1.00
##
##
## [1] "Omit"
##           winterJan winterDec
## winterJan      1.0      0.8
## winterDec      0.8      1.0

# Punchline: defining winter as 1:3, or 12:2 has little impact, but we went with
# Winter = Jan-Mar for consistency with other data source
rm(b) # delete this before it gets in the way...

# Interpolate missing samples? No!

#####
# Small copepods
# Multiply by avg wt of sm copepod (originally assumed mean size of sm copepod = 1.2 mm (
EMAX), but after some tweaks to get closer to SEDAR(2020) increased to 2.0mm; and note th
at when I went to the primary literature, I was able to confirm the conversion is for wet
weight! (not dry wt as reported in EMAX)):
a$`Small copepods` <- a$`Small copepods` * ((0.0881*2.0^2.8514)/1000)

# See how close we can get to EMAX:
tmp <- ddply(a[is.element(a$Year,1996:2000)],.(EPU,Season),summarize, .mean=mean(`Small
copepods`,na.rm=TRUE)) #

a$`Small copepods` <- a$`Small copepods` * 1 # already WW so no need to convert
mean(a$`Small copepods`,na.rm=TRUE)

## [1] 151.4111

```

```

# Now convert to wt per m2
a$`Small copepods` <- a$`Small copepods`/10
mean(a$`Small copepods`,na.rm=TRUE)

## [1] 15.14111

#####
# Large copepods
# Multiply by avg wt of lg copepod (originally assumed avg size of lg copepod to be 2.5mm
, but after tuning to SEDAR (2020) using 2.2 mm; and as above, the primary literature con
firms the equation below converts length to wet weight!
a$`Large copepods` <- a$`Large copepods` * ((0.0881*2.2^2.8514)/1000)

# See how close we can get to EMAX:
tmp <- ddply(a[is.element(a$Year,1996:2000),c("EPU","Season","Large copepods")],.(EPU,Sea
son),summarize, .mean=mean(`Large copepods`,na.rm=TRUE))

a$`Large copepods` <- a$`Large copepods` * 1 # Already wet wt (WW), so no need to convert
mean(a$`Large copepods`,na.rm=TRUE)

## [1] 180.173

# Now convert to wt per m2
a$`Large copepods` <- a$`Large copepods`/10
mean(a$`Large copepods`,na.rm=TRUE)

## [1] 18.0173

#####
# Gel zooplankton
# See how close we are to EMAX (numbers per 10m2; Table 6.1):
ddply(a[is.element(a$Year,1996:2000),],.(EPU), summarize, .mean=mean(`Gelatinous zooplan
kton`,na.rm=TRUE)) # We're in the right ballpark with abundance!

##      EPU      .mean
## 1  GB  93973.4
## 2  GOM 132282.7
## 3  MAB 108197.7
## 4   SS 132206.1

# Final
a$`Gelatinous zooplankton` <- a$`Gelatinous zooplankton` * ((exp(-1.54+2.65*log(0.55))/0.
0448)/1000)# EMAX w/ typo corrected, and have tweaked the mean size of gelzooop to tune to
SEDAR (2019), and converted to WW
# Then convert to abundance per m2
a$`Gelatinous zooplankton` <- a$`Gelatinous zooplankton`/10
mean(a$`Gelatinous zooplankton`,na.rm=TRUE) # 6.7 g kg per m2

## [1] 6.709856

#####
# Chaetognatha
# Multiply by avg wt of Chaetognath (g): DW

a$`Chaetognatha` <- a$`Chaetognatha` * ((0.00097*4^2.365)/1e3) # DW g/10m2

# How close to EMAX can we get:

```

```

tmp <- ddply(a[is.element(a$Year,1996:2000)],,.(Season,EPU), summarize, .mean=mean(`Chaetognatha`,na.rm=TRUE)*1000) # convert back to mg for comparison with EMAX tables
apply(reshape(tmp,direction="wide",idvar="Season",timevar="EPU"),[-1],2,mean) # Avg DW (mg) Pretty good!

## .mean.GB .mean.GOM .mean.MAB .mean.SS
## 225.19257 103.57818 389.55422 56.49015

((apply(reshape(tmp,direction="wide",idvar="Season",timevar="EPU"),[-1],2,mean)/1000)/0.083)/10 # WW g/m2; very good!

## .mean.GB .mean.GOM .mean.MAB .mean.SS
## 0.27131635 0.12479299 0.46934244 0.06806042

# Convert to wet weight:
a$`Chaetognatha` <- a$`Chaetognatha`/0.083 # DW = 8.3% (+/- 0.2%) of the wet weight

# Convert per 10m2 to per m2
a$`Chaetognatha` <- a$`Chaetognatha`/10
mean(a$`Chaetognatha`,na.rm=TRUE) # mean g/m2; very close to EMAX Table 7.1

## [1] 0.2662079

#####
# Other micronekton
# How close can we get to EMAX:
tmp <- ddply(a[is.element(a$Year,1996:2000)],,.(Season,EPU), summarize, .mean=mean(`Micronekton`,na.rm=TRUE))
apply(reshape(tmp,direction="wide",idvar="Season",timevar="EPU"),[-1],2,mean) # quite a bit higher than EMAX abundance per 10 m2

## .mean.GB .mean.GOM .mean.MAB .mean.SS
## 33046.69 14592.73 76284.59 32307.15

# Multiply by avg wt of a micronekton (g): DW
a$`Micronekton` <- a$`Micronekton` * (0.28/1e3) # DW g/10m2 # Assume
# dry wt = 1.2 mg, or 0.0012 g each

# Convert to wet weight:
a$`Micronekton` <- (9.299855693452*a$`Micronekton`) # From EMAX Table 7.1, assume DW = 0.1075286 x WW; therefore 1/0.1075286

# How close to EMAX did we get:
ddply(a,.(EPU), summarize, .mean=mean(`Micronekton`,na.rm=TRUE))

## EPU .mean
## 1 GB 58.05299
## 2 GOM 38.79491
## 3 MAB 113.95131
## 4 SS 99.00890

# Convert per 10m2 to per m2
a$`Micronekton` <- a$`Micronekton`/10
mean(a$`Micronekton`,na.rm=TRUE)

## [1] 7.769656

```

```
#####
# Pteropods
tmp <- dply(a, .(Season,EPU), summarize, .mean=mean(`Pteropod`,na.rm=TRUE))
apply(
  reshape(tmp,direction="wide",idvar="Season",timevar="EPU")[,-1],
  2,mean)/10 # number per m2

## .mean.GB .mean.GOM .mean.MAB .mean.SS
## 1475.2357 606.4824 2133.6823 534.7312

# Multiply by avg wt of a Pteropod (g): DW
a$`Pteropod` <- a$`Pteropod` * ((0.137*1.5^1.5005)/1e3) # DW g/10m2 (assume avg size of
Pteropod is 1.5mm; see spreadsheet for details, but essentially, given the len-wt equatio
n I have, this is the size that approximates the avg wt of 'Other micronekton' and makes
sense (Harvey noted that most things they see in nets are < about 1.5mm).

# Convert to wet weight (assuming same relationship as Other micronekton...):
a$`Pteropod` <- (9.299856*a$`Pteropod`)

# Convert per 10m2 to per m2
a$`Pteropod` <- a$`Pteropod`/10
mean(a$`Pteropod`,na.rm=TRUE)

## [1] 2.863368

#####
# Area of each EPU:
epu <- data.frame("EPU"=c("MAB","GOM","SS","GB"), "areaKm2"=c(126862117858, 68311878130,
30209438176, 57307696043)/1e6)

# units = square km; note that
by having units = km2, multiplying grams per m2 by km2, results in MT!

A <- merge(a,epu,by.x="EPU",by.y="EPU",all.x=TRUE)
B <- A[A$Year>1984,] # Note th
at for final run, including Scotian Shelf; in reality, should probably only use ~1/2 of S
S, but to keep me sane, going with the entire area.
B <- B[order(B$Year,B$EPU,B$Season),]

out <- dply(B, .(Year,EPU,Season), summarize,
  smCopepods=`Small copepods`*areaKm2,
  lgCopepods=`Large copepods`*areaKm2,
  gelZoop=`Gelatinous zooplankton`*areaKm2,
  chateo=`Chaetognatha`*areaKm2,
  micronek=`Micronekton`*areaKm2,
  pteropod=`Pteropod`*areaKm2
)

# Note th
at for final run, excluding Pteropods (not clear which, if any, group they should be comb
ined with, plus I had to make a number of assumptions about which I am not super clear on
(e.g., assumed length etc; see code above for all assumptions).
out2 <- data.frame(out[,c("Year","EPU","Season","smCopepods","lgCopepods","gelZoop")], "mi
cronek"=rowSums(out[,c("chateo","micronek")],na.rm=TRUE))#,pteropod=out[, "pteropod"])

# Average over seasons within a year (and EPU) # Skip do
```

```

wn to line approx XXXX for calcs with calcs by season (rather than avg over seasons)
out3 <- ddply(out2, .(Year,EPU), summarize, # I think
regular means (not weighted) are ok here, because I have already expanded the
  smCopepods=mean(smCopepods,na.rm=TRUE), # bioma
ss by the area of the EPU above!
  lgCopepods=mean(lgCopepods,na.rm=TRUE),
  gelZoop=mean(gelZoop,na.rm=TRUE),
  micronek=mean(micronek,na.rm=TRUE)#,
# pteropod=mean(pteropod,na.rm=TRUE)
)

```

```

G <- reshape::melt(out3,id=c("Year","EPU"))
colnames(G)[3:4] <- c("spp","totalWt")

```

```

H <- reshape(G,direction="wide", idvar=c("Year","spp"),timevar="EPU")

```

```

J <- data.frame(H[,c(1:2)],"totalWt"=rowSums(H[, -c(1,2)],na.rm=TRUE))

```

```

K <- reshape(J,direction="wide",idvar="Year",timevar="spp") # These w
ere the main outputs for NWACS-FULL (non-seasonal estimates) in total biomass
print(K)

```

##	Year	totalWt.smCopepods	totalWt.lgCopepods	totalWt.gelZoop	totalWt.micronek
## 1	1985	3431554	3425055	1268670.55	1112159.9
## 5	1986	3440344	3639841	800741.25	1370655.8
## 9	1987	2644889	4744488	1033159.61	2045327.5
## 13	1988	3588997	2888617	412271.91	776389.9
## 17	1989	7195068	1806078	274026.57	421091.0
## 21	1990	6789591	1405306	1075466.97	542611.8
## 25	1991	6417473	2577277	602044.93	905297.4
## 29	1992	4542449	4547651	1886828.09	2991831.8
## 33	1993	4349079	3720062	2503117.60	2363433.1
## 37	1994	4057677	3900582	1537559.39	2134298.4
## 41	1995	3548378	3543290	1248087.65	1386493.7
## 45	1996	6258443	3428739	2749291.72	3543381.0
## 49	1997	6126498	4336470	2760482.99	1809456.5
## 53	1998	6258336	3541391	3264596.80	3592821.6
## 57	1999	5644321	3943443	2367019.98	2445039.3
## 61	2000	8814913	5068131	4873471.86	2462884.7
## 65	2001	5307195	6967732	5461644.81	4754488.9
## 69	2002	5208322	5179505	1880889.14	3153318.9
## 73	2003	3146642	9658619	1175894.95	1137838.2
## 77	2004	3190861	7758209	1581424.72	3057422.3
## 81	2005	1677783	4762070	1026346.53	2826247.1
## 85	2006	4556767	4079822	1638105.96	4068321.2
## 89	2007	2564104	4951487	1541922.26	2002032.9
## 93	2008	3830498	3789274	1780934.58	1449205.0
## 97	2009	2944705	5356916	1375016.81	1549235.4
## 101	2010	3732773	3319225	1785077.23	1956917.6
## 105	2011	3137070	3553507	1577343.09	1346850.0
## 109	2012	7665340	3179961	1246596.34	2778440.6
## 113	2013	5504799	4176236	2766185.10	2665814.0
## 117	2014	3412276	5361754	3201763.47	3040033.3
## 121	2015	6770999	7423748	4309090.36	3672034.8
## 125	2016	5004780	6365594	3457567.62	2104169.9

## 129 2017	4174560	4289158	2206050.91	1454232.5
## 133 2018	4716700	4953572	4884686.04	1659904.4
## 137 2019	5160204	6022847	2360542.80	11621742.7
## 141 2020	3412234	493223	48392.79	140113.0
## 145 2021	4793319	2732574	3783866.95	1496381.3
## 149 2022	4690930	3560822	2700313.60	1642634.7
## 153 2023	5863249	7071742	1765079.98	3411208.5

Look at g/m2 over time across (and mean) # Here are the density estimates that were actually input into NWACS (note that for final input, 2020 est was omitted):

```
K[, -1]/sum(epu$areaKm2) #[epu$EPU!="SS", "areaKm2"])
```

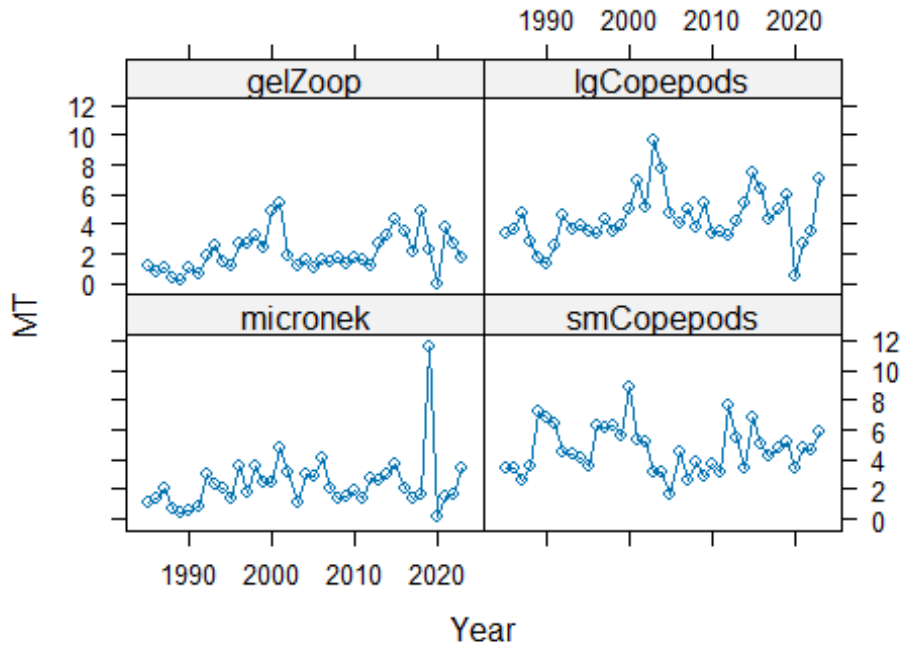
##	totalWt.smCopepods	totalWt.lgCopepods	totalWt.gelZoop	totalWt.micronek
## 1	12.138881	12.115890	4.4878329	3.9341876
## 5	12.169975	12.875682	2.8325659	4.8485984
## 9	9.356109	16.783292	3.6547295	7.2352024
## 13	12.695825	10.218281	1.4583829	2.7464247
## 17	25.452046	6.388873	0.9693497	1.4895798
## 21	24.017702	4.971171	3.8043888	1.9194511
## 25	22.701359	9.116937	2.1296916	3.2024259
## 29	16.068592	16.086997	6.6745217	10.5833948
## 33	15.384560	13.159457	8.8546025	8.3604783
## 37	14.353746	13.798035	5.4390083	7.5499307
## 41	12.552137	12.534138	4.4150223	4.9046240
## 45	22.138802	12.128924	9.7254262	12.5344611
## 49	21.672057	15.339957	9.7650145	6.4008253
## 53	22.138424	12.527420	11.5482817	12.7093537
## 57	19.966388	13.949651	8.3731668	8.6491547
## 61	31.182135	17.928157	17.2395641	8.7122814
## 65	18.773828	24.647862	19.3201846	16.8186701
## 69	18.424073	18.322135	6.6535131	11.1546440
## 73	11.131024	34.166685	4.1596457	4.0250227
## 77	11.287446	27.444118	5.5941788	10.8154164
## 81	5.935038	16.845488	3.6306287	9.9976504
## 85	16.119243	14.432085	5.7946847	14.3914003
## 89	9.070339	17.515537	5.4544416	7.0820505
## 93	13.550118	13.404292	6.2999309	5.1264608
## 97	10.416686	18.949713	4.8640253	5.4803114
## 101	13.204420	11.741526	6.3145852	6.9224583
## 105	11.097166	12.570281	5.5797403	4.7643872
## 109	27.115601	11.248887	4.4097469	9.8285384
## 113	19.472840	14.773141	9.7851853	9.4301295
## 117	12.070687	18.966829	11.3260132	10.7539039
## 121	23.951934	26.260988	15.2431042	12.9895650
## 125	17.704056	22.517840	12.2309024	7.4433529
## 129	14.767212	15.172594	7.8037500	5.1442452
## 133	16.684994	17.522913	17.2792335	5.8717950
## 137	18.253858	21.305398	8.3502542	41.1110977
## 141	12.070537	1.744742	0.1711861	0.4956398
## 145	16.956029	9.666290	13.3851633	5.2933436
## 149	16.593836	12.596157	9.5521695	5.8107047
## 153	20.740831	25.015790	6.2438463	12.0669101

```
apply(K[, -1]/sum(epu$areaKm2), 2, mean) # (sum(epu[epu$EPU!="SS", "areaKm2"])), 2, mean)
```

```
## totalWt.smCopepods totalWt.lgCopepods totalWt.gelZoop totalWt.micronek
## 16.650783 15.557799 7.456863 8.425592
```

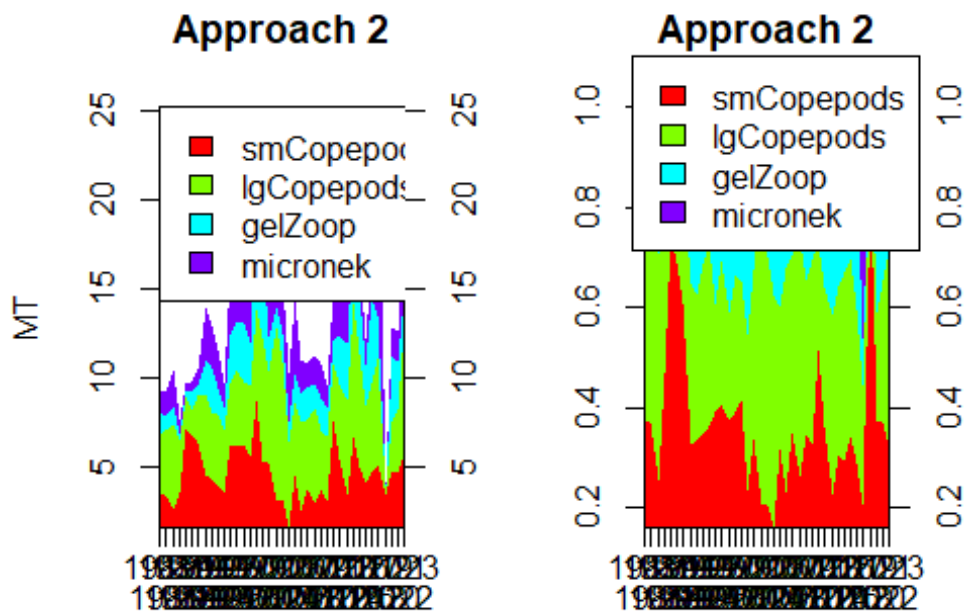
```
xyplot((value/1e6)~Year | as.factor(gsub("totalWt.", "", variable)), data=melt(K,id="Year")
,as.table=TRUE,type="o",ylab="MT",main="Approach 2")
```

Approach 2



```
stackpoly(x=K[,1], y=K[, -1]/1e6, staxx=TRUE,stack=TRUE,col=rainbow(4),main="Approach 2",y
lab="MT")
legend("topleft",legend=gsub("totalWt.", "", c(colnames(K)[-1])),fill=rainbow(4))
```

```
stackpoly(x=K[,1], y=K[, -1]/rowSums(K[, -1]), staxx=TRUE,stack=TRUE,col=rainbow(4),main="A
pproach 2")
op <- par(xpd=TRUE)
legend(x=1983,y=1.1,legend=gsub("totalWt.", "", c(colnames(K)[-1])),fill=rainbow(4),bg="whi
te")
```



```
# Any relationships among copepods and pteropods? Not run since we have lopped pteropods
#plot(K[-36,c("totalWt.smCopepods", "totalWt.pteropod")])
#plot(K[-36,c("totalWt.lgCopepods", "totalWt.pteropod")])
#plot(data.frame("copepods"=rowSums(K[-36,c("totalWt.smCopepods", "totalWt.lgCopepods")]),
"pteropods"=K[-36, "totalWt.pteropod"])))
```

```
# Do *NOT* average over season:
y 2025
```

```
# Februar
```

```
if(FALSE) {
out3 <-
out2 |>
# filter(Year==2022) |> # do a test year to make sure things are working the way I think they should
group_by(Year,Season) |>
summarize(smCopepods=sum(smCopepods,na.rm=TRUE),
lgCopepods=sum(lgCopepods,na.rm=TRUE),
gelZoop=sum(gelZoop,na.rm=TRUE),
micronek=sum(micronek,na.rm=TRUE)) |>
print(n=Inf,width=Inf)
}
```

```
# Equivalent, and much more straightforward than code immediately above:
```

```
out3 <-
ddply(out2, .(Year,Season), summarize,
```

```
# summing
```

over EPU

```
smCopepods=sum(smCopepods,na.rm=TRUE),  
lgCopepods=sum(lgCopepods,na.rm=TRUE),  
gelZoop=sum(gelZoop,na.rm=TRUE),  
micronek=sum(micronek,na.rm=TRUE)#,  
# pteropod=sum(pteropod,na.rm=TRUE)  
)
```

```
reshape(ddply(out3,.(Year,Season), summarize,  
  sumZoop=sum(smCopepods,lgCopepods,gelZoop,micronek)),  
  direction="wide",idvar="Year",timevar="Season")
```

Note th

at 0s should actually be NAs

##	Year	sumZoop.Fall	sumZoop.Spring	sumZoop.Summer	sumZoop.Winter
## 1	1985	11866997	10583094	10830011	3669655
## 5	1986	7072820	11167990	13981895	4783624
## 9	1987	9334159	16436986	12817552	3282758
## 13	1988	9263438	2339411	10212986	5162853
## 17	1989	12703981	0	0	7530727
## 21	1990	13823029	3316883	7070015	7342413
## 25	1991	15427760	0	13022019	3961795
## 29	1992	11415652	22037082	16922278	5500028
## 33	1993	14252968	19410488	8675370	4910128
## 37	1994	13643757	18013068	3439447	4738713
## 41	1995	8921034	12194187	12709408	5080367
## 45	1996	13924069	29349385	16207159	3321683
## 49	1997	16218188	9908675	23972698	5246246
## 53	1998	16363006	27770950	7994368	8077239
## 57	1999	17636641	11116963	17800414	7802163
## 61	2000	21125320	20838111	24911313	6876127
## 65	2001	12160602	46150646	16497493	7032286
## 69	2002	12205263	22928892	16386076	10167909
## 73	2003	11840605	26138413	15470687	7026271
## 77	2004	10016974	21947399	19250947	11136346
## 81	2005	5787924	16182318	11572917	7626626
## 85	2006	13753976	18670034	17394599	7553457
## 89	2007	9048197	16898982	10624785	7666222
## 93	2008	11197535	7639341	16783860	7778912
## 97	2009	8930440	15168377	15746996	5057678
## 101	2010	9041848	13863149	15664961	4606013
## 105	2011	12322879	13101685	3209166	4382965
## 109	2012	12679288	11810192	28189457	6802414
## 113	2013	14175211	23036242	16010903	5655751
## 117	2014	12501512	28063529	4837925	2684549
## 121	2015	19404099	29098999	5260569	7141123
## 125	2016	12206475	24948716	15745312	0
## 129	2017	13197754	19483093	0	5145389
## 133	2018	13337477	19219627	10569852	4454481
## 137	2019	55576794	10211181	22871891	2832654
## 141	2020	0	0	0	4514301
## 145	2021	15641733	10056105	14929678	2171441
## 149	2022	14261808	15312413	3032964	1755834
## 153	2023	8524460	20006440	29214146	0

Example

: sum(c(NA,NA,NA),na.rm=TRUE) sum is actually NA, not 0!

fin!

```
reshape(out3, direction="wide",idvar=c("Year"),timevar="Season")
```

##	Year	smCopepods.Fall	lgCopepods.Fall	gelZoop.Fall	micronek.Fall
## 1	1985	5621503	4376674	455355.3	1413464.2
## 5	1986	3301310	2452584	358975.5	959950.1
## 9	1987	3595523	2494079	103732.2	3140825.7
## 13	1988	5376774	2595919	101248.5	1189496.3
## 17	1989	8976746	2162722	319553.3	1244960.0
## 21	1990	11528095	1083204	420287.2	791442.1
## 25	1991	10629353	1929672	763738.8	2104996.3
## 29	1992	6395641	2235428	395506.7	2389075.1
## 33	1993	9878515	1980666	595296.4	1798490.5
## 37	1994	8884102	2502210	621298.1	1636145.9
## 41	1995	4589340	2358530	755111.3	1218052.7
## 45	1996	9842861	1989404	1015246.2	1076557.5
## 49	1997	9267882	2481839	1796712.1	2671754.2
## 53	1998	9959304	2943445	1510750.7	1949507.4
## 57	1999	8820127	3592861	1040202.6	4183450.0
## 61	2000	14349206	2097658	2150981.0	2527475.4
## 65	2001	3936137	5373462	752182.2	2098820.9
## 69	2002	6711774	3028381	563399.7	1901707.7
## 73	2003	3994491	4843060	912567.4	2090486.9
## 77	2004	3167717	3714475	355883.2	2778897.9
## 81	2005	1321187	2446586	346818.0	1673332.6
## 85	2006	6364806	3181383	1527853.9	2679933.4
## 89	2007	2656704	3873597	920312.0	1597583.1
## 93	2008	4473059	3435091	890056.1	2399329.4
## 97	2009	3872856	2948010	419687.0	1689886.6
## 101	2010	4609861	1762801	528706.9	2140478.5
## 105	2011	7282673	2411735	495579.9	2132891.1
## 109	2012	6322967	1457182	1599668.5	3299470.6
## 113	2013	9092327	1895519	1240664.1	1946700.6
## 117	2014	6325173	2051316	927106.8	3197916.8
## 121	2015	11746887	2492056	2647050.3	2518106.3
## 125	2016	4970044	2114298	2622446.0	2499687.3
## 129	2017	7861816	1867047	1367351.8	2101538.8
## 133	2018	5758145	2337609	2170168.6	3071554.5
## 137	2019	6695993	4573240	3817228.6	40490332.1
## 141	2020	0	0	0.0	0.0
## 145	2021	8363790	1126134	4227001.4	1924808.7
## 149	2022	8670437	1495086	1768366.0	2327918.1
## 153	2023	3594486	3224390	596598.4	1108986.2
##		smCopepods.Spring	lgCopepods.Spring	gelZoop.Spring	micronek.Spring
## 1		1541976.87	4662643	3142056.6	1236417.78
## 5		1300087.61	5875113	2023519.4	1969270.01
## 9		717818.47	9865187	2915015.5	2938965.06
## 13		53466.69	1668514	582171.2	35259.06
## 17		0.00	0	0.0	0.00
## 21		67376.75	2274950	974556.2	0.00
## 25		0.00	0	0.0	0.00

## 29	323263.47	8334016	6133949.2	7245852.82
## 33	479016.10	8241651	4913743.5	5776077.50
## 37	456116.87	8683843	3455742.6	5417365.24
## 41	975400.14	5741663	3030407.9	2446715.89
## 45	756765.26	9367878	7378321.2	11846421.31
## 49	85611.52	5133252	3216757.1	1473053.75
## 53	2955177.27	6267363	6752357.6	11796052.31
## 57	179184.47	5798914	4245462.4	893401.98
## 61	1901646.06	9245905	5207867.6	4482692.54
## 65	4549640.78	9976344	18292041.6	13332619.59
## 69	1248308.71	9143868	5067795.2	7468919.43
## 73	653234.68	23151976	2169522.8	163678.87
## 77	658732.43	11522486	4218758.7	5547421.78
## 81	69527.66	7737234	1214846.9	7160709.64
## 85	245265.63	6157693	1789232.0	10477843.97
## 89	1515370.48	8292069	3530016.3	3561526.20
## 93	458269.07	3744001	2969605.7	467465.20
## 97	294155.91	10322958	2632619.5	1918643.39
## 101	87019.65	6427073	4193349.3	3155706.17
## 105	370142.40	7169647	3558455.3	2003440.18
## 109	1630409.80	6931089	2013936.9	1234756.05
## 113	1363657.25	9046559	6005879.9	6620145.47
## 117	2030980.54	12386230	6542267.8	7104050.82
## 121	1098212.45	13070394	5970654.0	8959738.96
## 125	2356420.78	12088228	6117317.7	4386749.31
## 129	1466454.72	9780896	4954424.0	3281318.69
## 133	1443448.50	8765503	6956546.1	2054128.94
## 137	368744.55	5865879	2056461.3	1920096.36
## 141	0.00	0	0.0	0.00
## 145	1834812.92	5379498	1605314.8	1236479.19
## 149	1600206.14	6681764	4199220.0	2831222.67
## 153	2193581.05	9684622	3650999.7	4477237.03
##	smCopepods.Summer	lgCopepods.Summer	gelZoop.Summer	micronek.Summer
## 1	4770690.0	3575147.0	1105641.3	1378532.5
## 5	6859422.7	4857373.6	773376.1	1491722.2
## 9	4476390.4	5611511.7	974621.0	1755029.2
## 13	4234480.9	4071853.5	456415.9	1450236.0
## 17	0.0	0.0	0.0	0.0
## 21	5223545.7	130224.4	730566.3	985678.4
## 25	5375246.1	5333215.1	960073.2	1353484.6
## 29	7651072.2	6605512.4	579219.9	2086473.1
## 33	3320963.3	928036.2	3028339.5	1398030.9
## 37	1090095.4	482441.1	889869.6	977040.6
## 41	5376662.4	4922571.9	994805.9	1415367.5
## 45	11512403.6	1454158.3	2089910.8	1150686.6
## 49	8761497.4	8533924.2	4139568.6	2537708.2
## 53	4289270.5	1767346.1	1484272.7	453478.6
## 57	6205203.9	4021180.5	3362821.0	4211208.6
## 61	9347155.1	4347447.1	9216593.2	2000117.7
## 65	5854317.5	6353505.9	1422580.6	2867089.5
## 69	6709206.7	5365103.6	1757796.6	2553969.3
## 73	3609244.6	8158309.6	1576329.8	2126802.8
## 77	4816812.5	9691842.6	1383457.4	3358834.4
## 81	2897362.8	5384763.2	1023103.4	2267687.8
## 85	7664489.8	4395608.8	2815595.6	2518905.0

## 89	2687738.8	4845914.6	1041929.4	2049201.9
## 93	5910713.3	5265246.0	3147579.0	2460321.4
## 97	4703213.8	6732203.0	2289519.6	2022059.2
## 101	8532359.8	3316669.0	1941624.1	1874308.3
## 105	613654.1	692455.9	859179.4	1043876.5
## 109	17896283.1	3248194.1	1257496.7	5787482.9
## 113	7093095.4	4218738.3	3074069.6	1624999.8
## 117	1058762.5	430911.2	1598845.5	1749405.5
## 121	1774333.5	381969.7	1658126.2	1446139.9
## 125	7687874.8	4894254.9	1632939.2	1530242.8
## 129	0.0	0.0	0.0	0.0
## 133	3476359.7	1079127.5	4603758.3	1410606.3
## 137	8329318.5	8046493.0	2704177.8	3791901.6
## 141	0.0	0.0	0.0	0.0
## 145	4173017.4	2291133.6	6506260.2	1959266.6
## 149	886535.5	202388.7	977117.3	966922.7
## 153	11801680.0	8306213.8	1047641.9	8058610.6
##	smCopepods.Winter	lgCopepods.Winter	gelZoop.Winter	micronek.Winter
## 1	1792046	1085754.98	371628.974	420225.28
## 5	2300555	1374293.46	47093.963	1061680.68
## 9	1789825	1007173.18	139269.784	346490.27
## 13	2250624	2189827.40	291832.667	430568.20
## 17	5413389	1449433.40	228499.892	439404.05
## 21	4685648	1027765.26	1235672.669	393326.68
## 25	3247819	468944.55	82322.839	162708.72
## 29	3799817	1015648.61	438636.539	245926.22
## 33	2346655	1300720.99	781619.045	481133.33
## 37	3307254	674105.77	250711.903	506642.00
## 41	3252108	1150394.65	212025.578	465838.73
## 45	2705113	318386.37	198324.482	99858.49
## 49	3479846	767274.36	443815.767	555309.96
## 53	4516016	1853713.60	1535261.110	172247.91
## 57	5127682	1744333.77	438050.861	492096.63
## 61	5149091	729772.34	156009.708	841253.13
## 65	5238857	740540.17	333463.460	719425.41
## 69	6163999	3180666.54	134565.081	688679.16
## 73	4329597	2481129.83	45159.749	170384.31
## 77	4120181	6104030.45	367599.551	544535.04
## 81	2423053	3479696.56	1520617.880	203258.32
## 85	3952507	2584605.08	419742.355	596602.47
## 89	3396604	2794366.69	675431.339	799820.31
## 93	4479951	2712759.41	116497.576	469703.98
## 97	2908593	1424491.91	158241.169	566352.59
## 101	1701850	1770357.63	476628.635	657177.30
## 105	2720924	978918.04	475931.112	207192.18
## 109	4811700	1083377.91	115283.175	792053.03
## 113	3930439	1023197.48	230704.765	471410.10
## 117	821913	1216806.02	537070.377	108759.84
## 121	3282847	385338.24	1708783.782	1764154.07
## 125	0	0.00	0.000	0.00
## 129	3195409	1219529.93	296376.939	434072.47
## 133	4062941	283742.53	4469.333	103327.69
## 137	1448152	1048431.63	51430.116	284640.63
## 141	3412234	493222.97	48392.792	560451.91
## 145	1130863	126920.76	48686.357	864970.77

```
## 149      1203892      63270.92      44196.003      444475.30
## 153           0           0.00           0.000           0.00
```

```
H <- reshape(G,direction="wide", idvar=c("Year","spp"),timevar="EPU")
```

```
J <- data.frame(H[,c(1:2)],"totalWt"=rowSums(H[, -c(1,2)],na.rm=TRUE))
```

```
K <- reshape(J,direction="wide",idvar="Year",timevar="spp")
```

```
# Look at g/m2 over time across (and mean)
K[, -1]/sum(epu$areaKm2)#[epu$EPU!="SS", "areaKm2"])
```

##	totalWt.smCopepods	totalWt.lgCopepods	totalWt.gelZoop	totalWt.micronek
## 1	12.138881	12.115890	4.4878329	3.9341876
## 5	12.169975	12.875682	2.8325659	4.8485984
## 9	9.356109	16.783292	3.6547295	7.2352024
## 13	12.695825	10.218281	1.4583829	2.7464247
## 17	25.452046	6.388873	0.9693497	1.4895798
## 21	24.017702	4.971171	3.8043888	1.9194511
## 25	22.701359	9.116937	2.1296916	3.2024259
## 29	16.068592	16.086997	6.6745217	10.5833948
## 33	15.384560	13.159457	8.8546025	8.3604783
## 37	14.353746	13.798035	5.4390083	7.5499307
## 41	12.552137	12.534138	4.4150223	4.9046240
## 45	22.138802	12.128924	9.7254262	12.5344611
## 49	21.672057	15.339957	9.7650145	6.4008253
## 53	22.138424	12.527420	11.5482817	12.7093537
## 57	19.966388	13.949651	8.3731668	8.6491547
## 61	31.182135	17.928157	17.2395641	8.7122814
## 65	18.773828	24.647862	19.3201846	16.8186701
## 69	18.424073	18.322135	6.6535131	11.1546440
## 73	11.131024	34.166685	4.1596457	4.0250227
## 77	11.287446	27.444118	5.5941788	10.8154164
## 81	5.935038	16.845488	3.6306287	9.9976504
## 85	16.119243	14.432085	5.7946847	14.3914003
## 89	9.070339	17.515537	5.4544416	7.0820505
## 93	13.550118	13.404292	6.2999309	5.1264608
## 97	10.416686	18.949713	4.8640253	5.4803114
## 101	13.204420	11.741526	6.3145852	6.9224583
## 105	11.097166	12.570281	5.5797403	4.7643872
## 109	27.115601	11.248887	4.4097469	9.8285384
## 113	19.472840	14.773141	9.7851853	9.4301295
## 117	12.070687	18.966829	11.3260132	10.7539039
## 121	23.951934	26.260988	15.2431042	12.9895650
## 125	17.704056	22.517840	12.2309024	7.4433529
## 129	14.767212	15.172594	7.8037500	5.1442452
## 133	16.684994	17.522913	17.2792335	5.8717950
## 137	18.253858	21.305398	8.3502542	41.1110977
## 141	12.070537	1.744742	0.1711861	0.4956398
## 145	16.956029	9.666290	13.3851633	5.2933436
## 149	16.593836	12.596157	9.5521695	5.8107047
## 153	20.740831	25.015790	6.2438463	12.0669101

```
apply(K[, -1]/(sum(epu$areaKm2)),2,mean)#[epu$EPU!="SS", "areaKm2"])),2,mean)
```

```

## totalWt.smCopepods totalWt.lgCopepods totalWt.gelZoop totalWt.micronek
## 16.650783 15.557799 7.456863 8.425592

#####
# Biovolume:
.seasons <- data.frame("Season"=c("Winter", "Spring", "Summer", "Fall"), "Qs"=1:4)
aa <- merge(a, .seasons, all.x=TRUE)

bv <- merge(aa[,c("Year", "Season", "Qs", "EPU", "Biovolume")], epu, all.x=TRUE)

# annual biovolume, averaged over all seasons and regions
r1 <- ddply(bv, .(Year), summarize,
  meanBv=mean(Biovolume, na.rm=TRUE))

# mean weighted by size of EPU:
r2 <- ddply(bv, .(Year), summarize,
  meanBv=weighted.mean(Biovolume, areaKm2, na.rm=TRUE))

matplot(1:47, cbind(r1[,2], r2[,2]), type="o")
cor(cbind(r1[,2], r2[,2]), method="spearman")

##      [,1]      [,2]
## [1,] 1.000000 0.970629
## [2,] 0.970629 1.000000

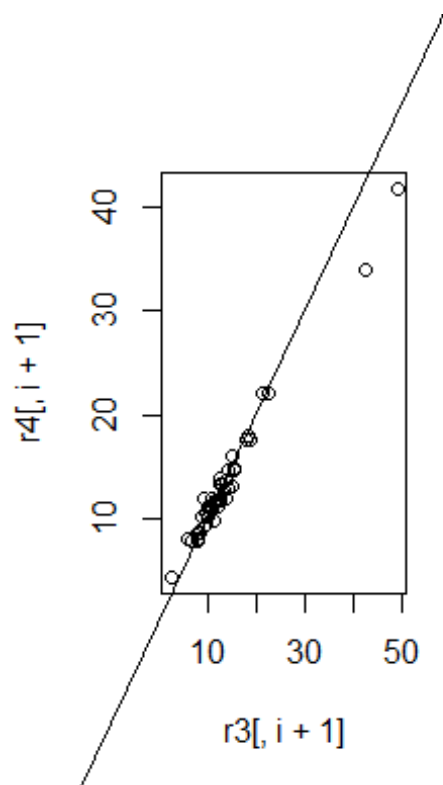
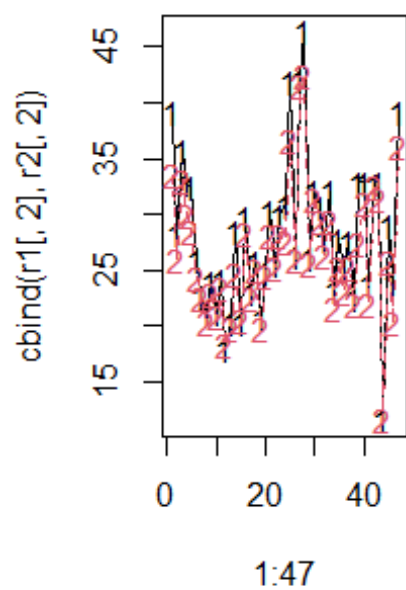
sbv <- ddply(bv, .(Year, Qs), summarize, # seasona
  l biovolume, weighted by EPU area
  meanBv=weighted.mean(Biovolume, na.rm=TRUE))

sbv.w <- ddply(bv, .(Year, Qs), summarize, # seasona
  l biovolume, weighted by EPU area
  meanBv=weighted.mean(Biovolume, areaKm2, na.rm=TRUE))

r3 <- reshape(sbv, direction="wide", idvar="Year", timevar="Qs")
r4 <- reshape(sbv.w, direction="wide", idvar="Year", timevar="Qs")

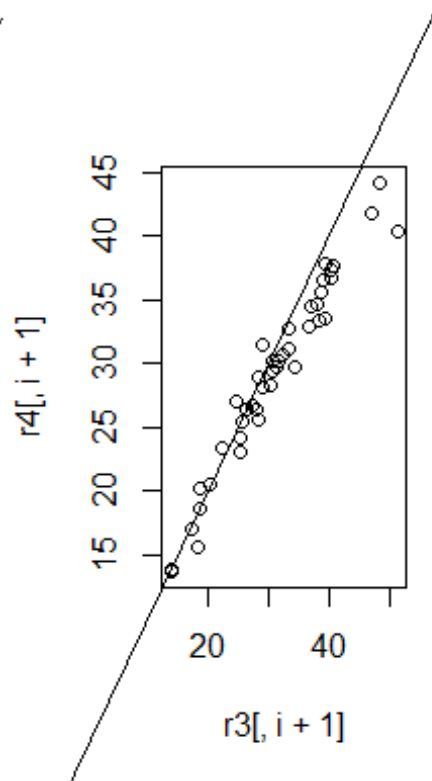
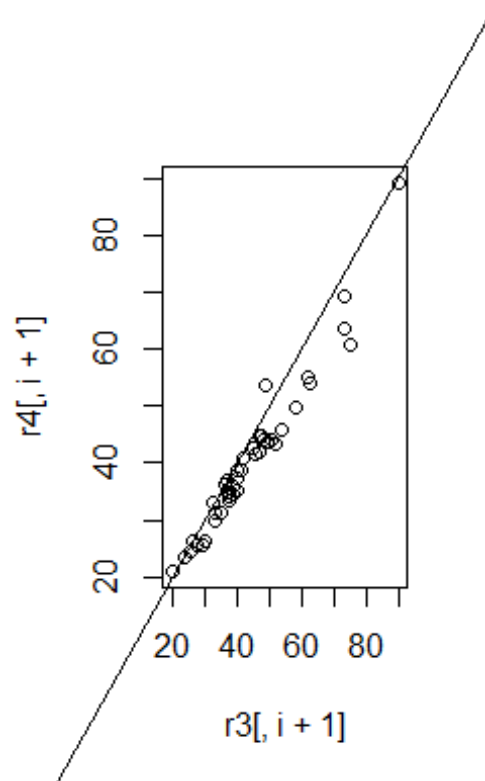
op <- par(ask=FALSE)
for(i in 1:4) {
  plot(r3[,i+1], r4[,i+1]); abline(lm(y ~ x, data = data.frame(x = 1:10, y = 1:10)))
  print(cor(r3[,i+1], r4[,i+1]))
}

```



```
## [1] NA
```

```
## [1] NA
```

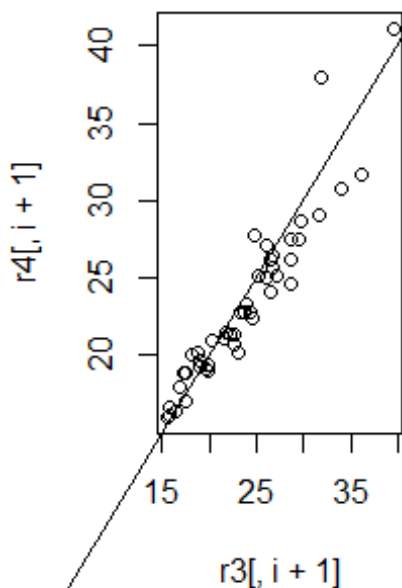


```
## [1] NA
```

```
## [1] NA
```

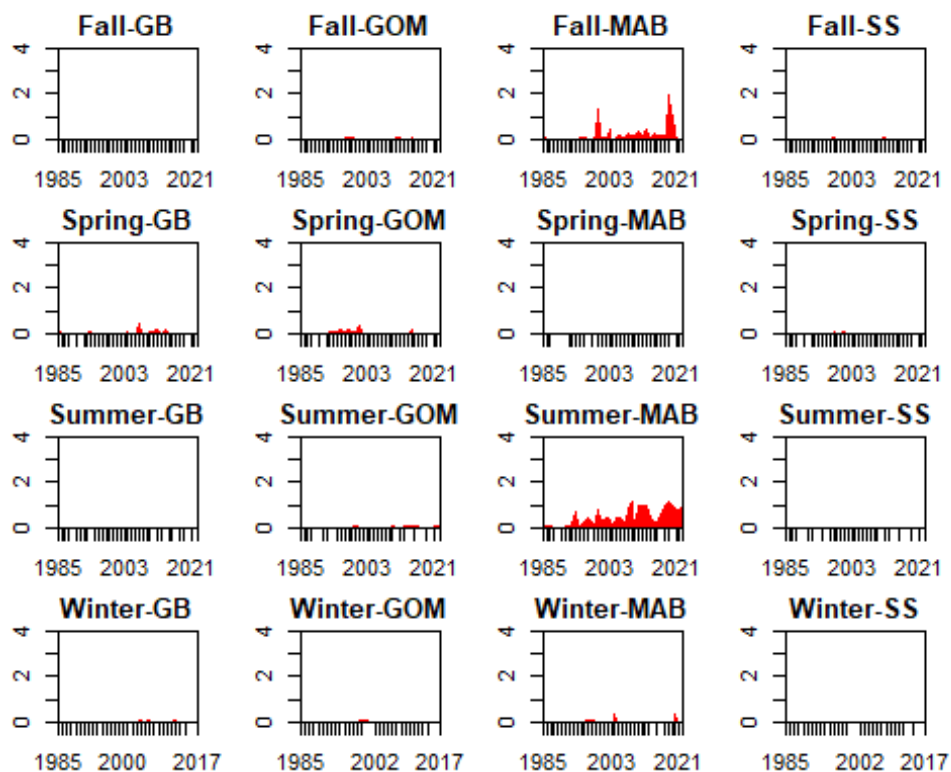
Miscellany:

```
plot(1,1,type="n",xlab="",ylab="",axes=FALSE)
legend("center",legend=colnames(a)[-c(1:3)],fill=rainbow(7),cex=2)
```



Biovolume
Small copepods
Large copepods
Gelatinous zoopl
Chaetognatha
Micronekton
Pteropod
Omit

```
op <- par(mfrow=c(4,4),mar=c(2,2,2,2))
lapply(split(a[a$Year>1984,], paste(a[a$Year>1984,"Season"],a[a$Year>1984,"EPU"],sep="-")
), function(x) {
#   stackpoly(x$Year,x[,-c(1:3)]/rowSums(x[,-c(1:3)]))
x <- x[-which(is.na(x$`Small copepods`)),]
stackpoly(x$Year,(x[,-c(1:3)]/1e6),stack=TRUE,col=rainbow(7),
  main=paste(x$Season[1],x$EPU[1],sep="-"),
#   ylab="Abund (x million) per 10^m2",
  staxx=FALSE,axis4=FALSE,ylim=c(0,4))
#   legend("topright",legend=colnames(a)[-c(1:3)],fill=rainbow(7))
})
)
```



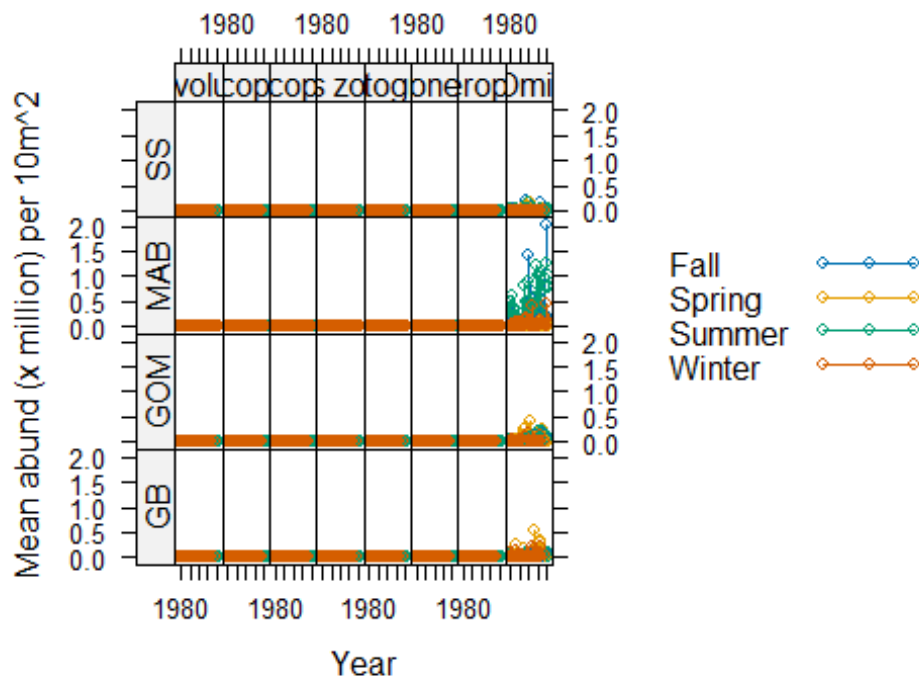
```
## `$Fall-GB`
## NULL
##
## `$Fall-GOM`
## NULL
##
## `$Fall-MAB`
## NULL
##
## `$Fall-SS`
## NULL
##
## `$Spring-GB`
## NULL
##
## `$Spring-GOM`
## NULL
##
## `$Spring-MAB`
## NULL
##
## `$Spring-SS`
## NULL
##
## `$Summer-GB`
## NULL
##
## `$Summer-GOM`
## NULL
##
## `$Summer-MAB`
## NULL
```

```
##
## `$Summer-SS`
## NULL
##
## `$Winter-GB`
## NULL
##
## `$Winter-GOM`
## NULL
##
## `$Winter-MAB`
## NULL
##
## `$Winter-SS`
## NULL
```

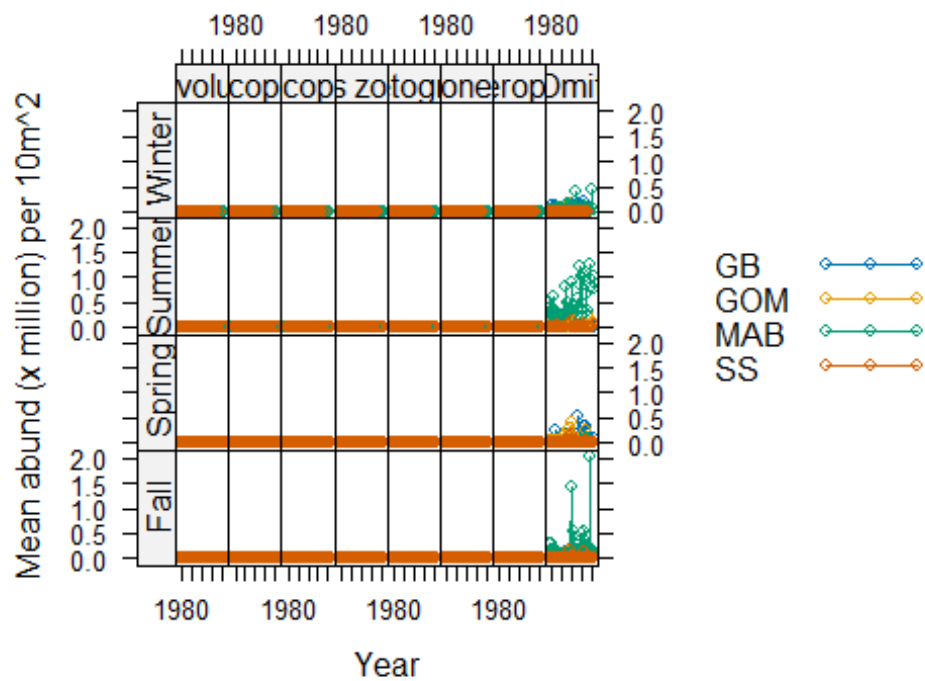
```
par(op)
```

```
A <- melt(a,id=c("Year","Season","EPU"))
```

```
useOuterStrips(xyplot((value/1e6)~Year | variable*EPU, groups=Season, data=A,type="o",auto.key=TRUE,ylab="Mean abund (x million) per 10m^2"))
```

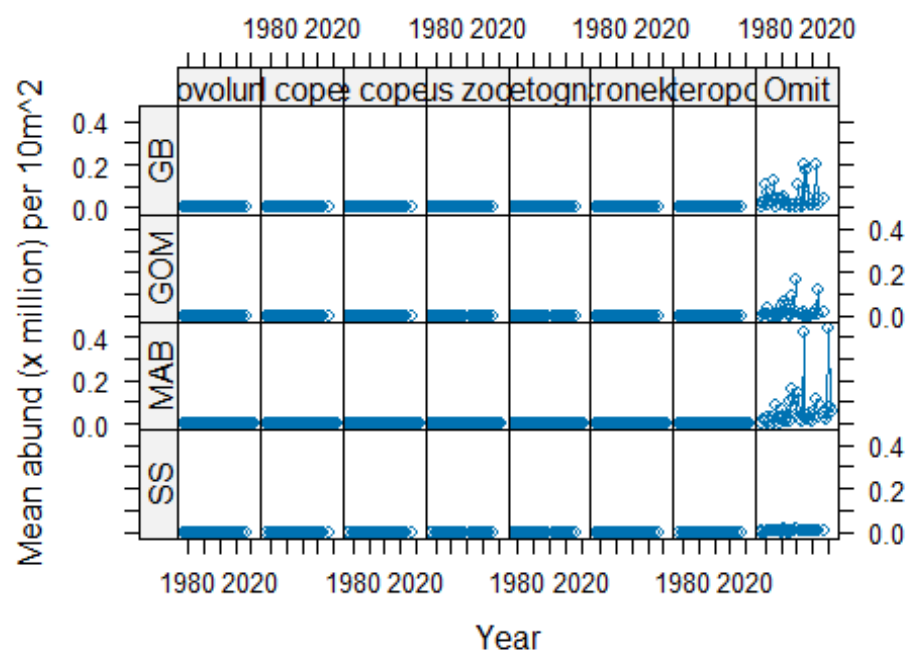


```
="Omit",]
useOuterStrips(xyplot((value/1e6)~Year | variable*Season, groups=EPU, data=A,type="o",auto.key=TRUE,ylab="Mean abund (x million) per 10m^2"))
```

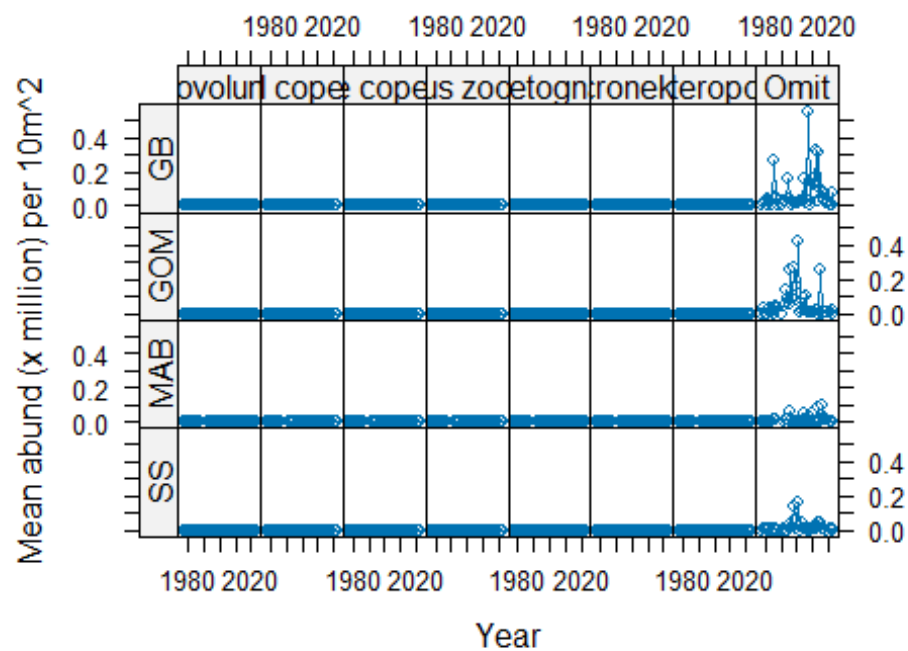


```
for(i in c("Winter","Spring","Summer","Fall")) {
  print(useOuterStrips(xyplot((value/1e6)~Year | variable*EPU, data=A[A$Season==i,],
    type="o",auto.key=TRUE,as.table=TRUE,main=i,ylab="Mean abund (x million) per 10m^2"
  )))
}
```

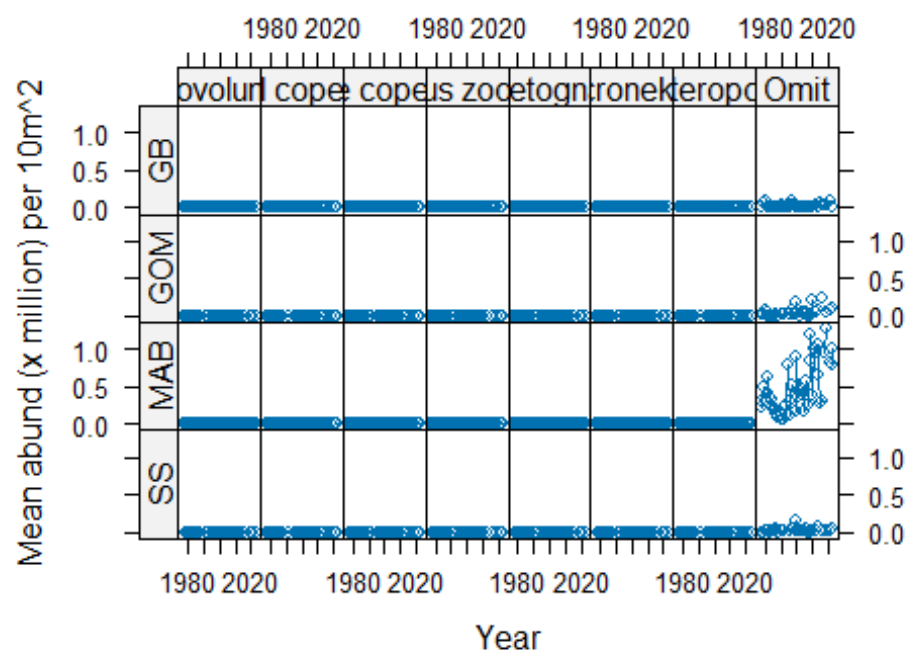
Winter



Spring



Summer



Fall

