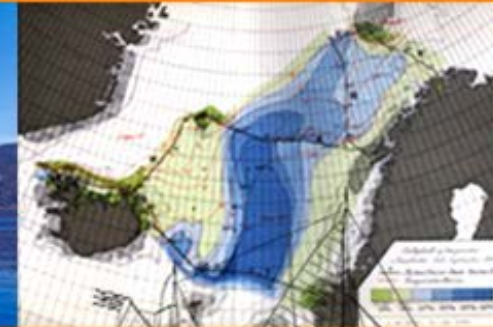
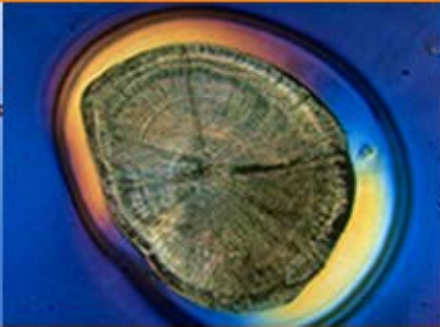


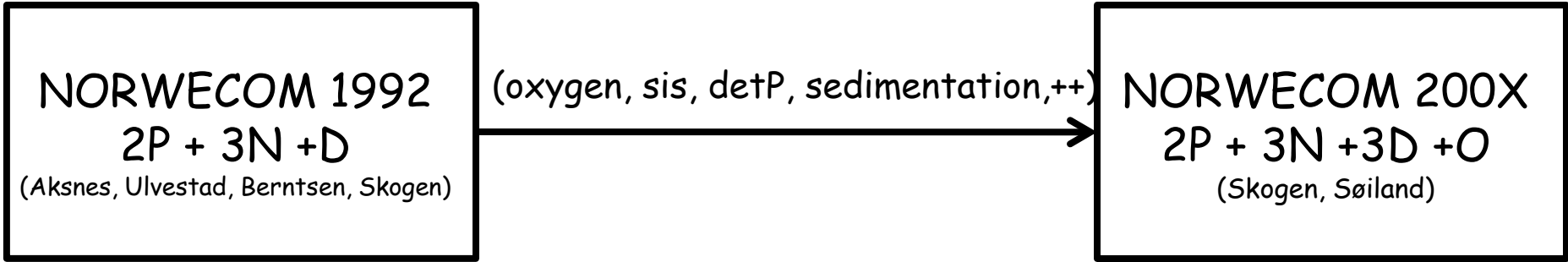


INSTITUTE OF MARINE RESEARCH



Towards a generic zooplankton IBM module in NORWECOM.E2E

Morten D. Skogen & Geir Huse



- Micro/meso-zoo
- Contaminants
- Calanus IBM
- FishIBM

NORWECOM 2010

Recoding, structuring, new modules

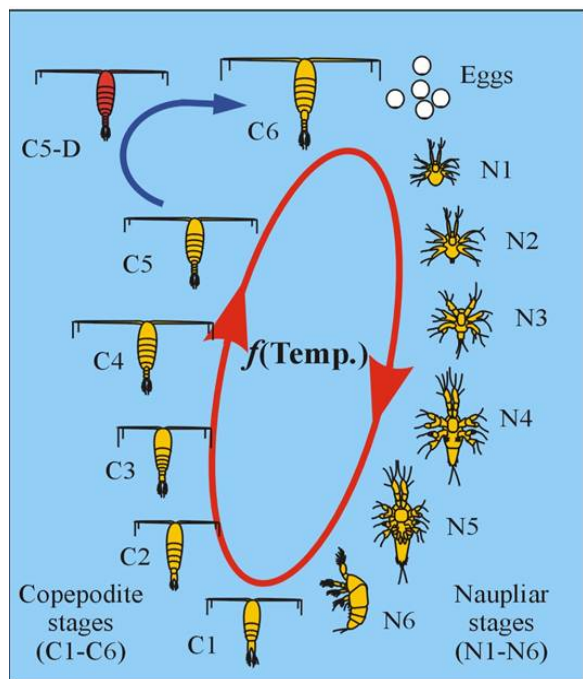
Online (POM Noth Sea)
(without IBMs)

Offline ROMS

Full Offline

NORWECOM.E2E

IBM Calanus module



From <http://pulse.unh.edu/>

Stage

Structural weight

Individual number

Fat content

Position

Depth



Feeding: functional response, type 2 (Campbell 2001)

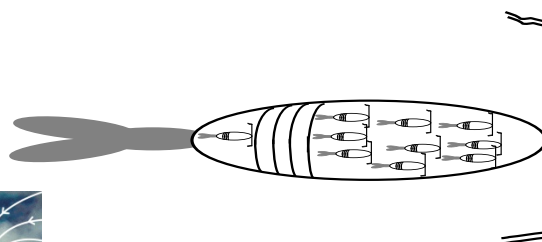
Growth: bioenergetics (Carlotti & Wolf 1998)

Mortality: predation, starvation, spawning

Reproduction: mature adults above weight and fat thres.

Vertical movement: ontogeny, dvm

Horizontal movement: by currents



Object oriented approach

Object oriented programming - is a programming paradigm that uses "objects"- data structures consisting of data fields and methods together with their interactions - to design applications and computer programs

1) define a datatype with all necessary information about a calanus (or fish or krill) superindividual and use it inside a data type that contains all information about a stock/species.....

and

2) define routines that operates on these data types

```
module zoo_generic
```

```
! Type with attributes and strategy vector ++
```

```
type superind
```

```
    logical  :: alive      ! True if alive, false if dead  
    integer  :: stage     ! 0 egg, 1-6 naupli, 7-13 copepod  
    integer  :: inumb     ! Number of individuals in superind.  
    real      :: x,y,z    ! Position (x,y,z)
```

```
    .....
```

```
end type superind
```

```
! Type for a whole population. Superind+ population specific parameters
```

```
type zoopop
```

```
    character(len=6) :: species ! Name of species  
    integer           :: pop    ! Pointer to last super ind.
```

```
    .....population specific parameters.....
```

```
    type(superind), dimension(:), allocatable :: zoopl
```

```
end type zoopop
```

type zooplankton

! Attribute vector

logical :: alive ! true if alive, false if dead
integer :: stage ! 0 is egg, 1-6 nauplia, 7-11 copepodits, 12 adult
real :: inumb ! Internal number in individuals
real :: sweight ! Structural weight in micrograms
real :: lstage ! Stage longevity
real :: fat ! Fat energy level in KJ
real :: moult ! Moulting cycle fraction (Egg,N2,N3) | cum. egg number (adult)
real :: maxegg ! Total number of eggs in a super individual
real :: xpos,ypos,zpos ! Position (x,y,z)
integer :: diapause ! 0 diapause,1 active,2 move down,3 move up
real :: growth ! Summing of daily growth
real :: egestion ! Summing of hourly egestion
integer :: mynumb ! Unique identifier
real :: ingestion ! Ingestion rate
real :: grate ! growth rate
integer :: numegg ! number of spawned eggs
real :: inumb_int ! Internal number in individuals at birth

! Strategy vector

integer :: wud ! Wake_Up_Day
real :: fsr ! FSR
integer :: afd ! Allocation_to_Fat_Day
integer :: owd ! Over_Wintering_Depth
integer :: vm1
real :: vm2

! Predation risk

real :: prisk(4) ! Predation risk. Species dependent. Level set in CALCPRISK

end type zooplankton

Population specific parameters *C.finmarchicus*:

```
real, dimension(0:13) :: st_wgt    ! Stage weight
real, dimension(0:13) :: mort      ! Mortality rates
real, dimension(0:13) :: a         ! Stage longevity parameter
real      :: stloli                ! Lower limit stage weight factor before starvation
real      :: strate                ! Starvation rate
integer   :: maxmoult              ! Maximum number of eggs to be spawned
integer   :: maxlstage             ! Maximum number of days in each stage
integer   :: mthresh              ! Mortality threshold, predation or MonteCarlo
real      :: cmat                  ! Structural weight for spawning
real      :: eggw                  ! Egg weight
real      :: cs                    ! Fat/eggw ratio for spawning
real      :: initw                 ! Initial weight at spawning
real      :: starvation            ! Rate of starvation at low stage_weight
real      :: meffect               ! Mutation effect
real      :: mprob                 ! Mutation probability
.....      :: .....
```

Fortran 90 modules:

- **zoo_generic**: typedefinitions and subroutines that operates on these (and are assumed general)
 - kill_zoopl, init_zoopl, copy_zoopl, spawn_zoopl, count_zoopl, popzip, phytomort, calmort, calrep, vertmov, combind
- **tools** (bad name..): useful routines that does not operate on the zooplankton types
 - val3d, getr, easyr, deriv, surlig
- **ladim**: Lagrangian Advection DIffusion Model (Ådlandsvik)
- **calanus_mod** (one module for each species): uses the other modules and (until now) the remaining species specific routines
 - calanus (*mainprogram* should be *identical* for all zooplankton)
calsta, initialise, calcprisk

First test - implementing a Krill module (Strand + Huse, IMR).....

- swimming behaviour (to be implemented in ladim)
- new/additional strategies:
 - night-depth
 - meters above bottom
 - first and last day for spawning
- additional parameters:
 - maxmig = max daily vertical migration
 - first copepod stage (krill=5, calfin=7)
 - stage for spawning (krill=12, calfin=13)
 - a few other minor details...
- **conclusion:** can easily adapt the generic module, perhaps with a few less elegant: `if(zoopl%species.eq.'krill')then`

Plan to extend model with:

- calanus helgolandicus, glacialis, hyperboreus
- generic fish-module

Remaining issues:

- general coupling between different IBMs (today only two-way hardcoded coupling between phyto and calfin IBM, and coupling between IBM fish model and IBM calfin implemented)
- order of predation - who to eat first - both between superindividuals and between stocks/species
-

Thank you for your attention



Bergen seen from Mt. Ulriken

