

Adding Variables to ROMS

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Considerations

- Input/initialization
- Output?
- Is it gridded or not?
- Where does it logically belong?



Four Examples

- Scalar variable
- A 3-D tracer
- A static 2-D gridded field
- A time-dependent 2-D gridded field





Simplest Case

- A scalar (not gridded), common to all processes
- No need for I/O
- In ROMS/Modules/mod_scalars.F:

#ifdef ICE_MODEL

logical, dimension(Ngrids) :: Lice

#endif

#ifdef ICE_MODEL

Lice(ng)=.TRUE.

#endif







Where?

- Lice is a logical switch, similar to logicals Lbiology, Lfloats, and Lsediment
- Put it with the others, but keep the #ifdefs around so you can find the ICE_MODEL changes later





Second Case

- The ecosystem models have NBT tracers
- If using NEMURO, the value of NBT comes from nemuro_mod.h, included by mod_biology.F
- The biology files are all under ROMS/Nonlinear/Biology





NEMURO Iron

- Jerome Fiechter added iron to NEMURO, using three new tracers
- # ifdef IRON_LIMIT

NBT = 14

else

NBT = 11

endif





New Tracer ID (nemuro_mod.h)

! Set biological tracer indices. integer :: iopal # ifdef IRON_LIMIT integer :: iFeSp ! Small phyt Fe integer :: iFeLp ! Large phyt Fe integer :: iFeD_ ! Dissolved Fe # endif





Give them Values





Other Iron Changes

- nemuro_mod.h set up some rate constants
- nemuro_inp.h read in rate constants
- Add the rate constants to a copy of nemuro.in called nemuro_iron.in
- Add code to nemuro.h to use the rate constants and iron fields







Iron I/O Changes

- nemuro_def.h NetCDF definitions for writing iron vars (see the opal code)
- nemuro_wrt.h output of iron vars
- nemuro_var.h check for idTvar (iFeSp) and friends when reading varinfo.dat
- Add them to varinfo.dat
- Add iron fields to boundary and initial condition files (or ana_initial.h)





Finally...

- Adding passive tracers is even simpler, changing NPT in ocean.in
- To output them, change Hout(inert) in ocean.in
- To provide initial and boundary conditions, call the fields dye_01, dye_west_01, etc.





Third Case

- Spatially variable bottom friction
- 2-D field to be read in from the grid file, treated like the other grid variables
- I tried to add it live for a ROMS training in Maryland – and failed





Label Your Code

- Start by picking a cppdef: RDRG_GRID, also ANA_RDRG in the case of defining it analytically
- More later on the ANA_ business
- Add both RDRG_GRID and ANA_RDRG to checkdefs.F and cppdefs.h





Which Module?

I put it in mod_forces.F, with the other bottom drag variables:

#ifdef RDRG_GRID

real(r8), pointer :: rdrg_grid(:,:)

#endif

#ifdef RDRG_GRID

allocate(FORCES(ng)%rdrg_grid(LBi:UBi,LBj:UBj))

#endif

ifdef RDRG_GRID

```
FORCES(ng) % rdrg_grid(i,j) = IniVal
```

endif





mod_forces.F

- The three chunks go in three different parts of mod_forces
 - Define the data structure
 - Allocate the gridded variables
 - Initialize the gridded variables
- If adding a gridded variable, make sure to add it to a module with other gridded variables







Input via get_grid.F

 Because I put the variable in mod_forces, we need to "use" it:

#ifdef RDRG_GRID
 USE mod_forces

#endif

- We can check to see if rdrg_grid is in the file, then read it, just like the 'h' variable, including communication
- Doesn't need to be in varinfo.dat





Analytic function

- You might want to read the grid file first, then call ana_rdrg instead of reading the field
- In initial.F, after the get_grid block:





```
#ifdef ANA RDRG
!$OMP PARALLEL DO PRIVATE(thread, subs, tile)
SHARED(ng, numthreads)
      DO thread=0, numthreads-1
        subs=NtileX(ng)*NtileE(ng)/numthreads
        DO tile=subs*thread, subs*(thread+1)-1
          CALL ana rdrg (ng, TILE, iNLM)
        END DO
      END DO
!$OMP END PARALLEL DO
#endif
```





More ANA_RDRG

Add to analytical.F:

- # if defined ANA_RDRG
- # include <ana_rdrg.h>
- # endif

Copy ana_mask.h to create ana_rdrg.h then edit it



ana_rdrg.h

- Add "USE mod_forces"
- Put "FORCES(ng) % rdrg_grid" in argument list to ana_rdrg_tile, also variables it depends on
- Strip out "mask" scratch array and all the umask, vmask stuff





Passing to Tile Routine

CALL ana_rdrg_tile (ng, tile, model, & LBi, UBi, LBj, UBj, & IminS, ImaxS, JminS, JmaxS, & FORCES(ng) % rdrg_grid, & GRID(ng) % h) • Change where called and declaration: real(r8), intent(out) :: rdrg_grid(LBi:,LBj:)

real(r8), intent(out) :: h(LBi:,LBj:)



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Set rdrg_grid

#ifdef NEP5 DO j=JstrR, JendR DO i=IstrR,IendR rdrg grid(i,j)=... END DO END DO #else ana rdrg.h: no values provided for rdrg grid. #endif





Communication

#if defined EW PERIODIC || defined NS PERIODIC CALL exchange r2d tile (ng, tile, & LBi, UBi, LBj, UBj, & & rdrg grid) & #endif #ifdef DISTRIBUTE CALL mp exchange2d (ng, tile, model, 1, & LBi, UBi, LBj, UBj, & & NghostPoints, EWperiodic, & & NSperiodic, rdrg grid) δ #endif







ANANAME

Ana_mask.h has:

ANANAME(15) = $_{\rm FILE}$

• ANANAME is defined in mod_ncparam.F:

character (len=256), dimension(37) :: ANANAME

- Then used in def_info.F and close_io.F
- "Magic number" of 37 in all three places





More ANANAME

 We can change all three values to 38 and set it in ana_rdrg as:

ANANAME(38) = ___FILE__

- Then next week/year Hernan adds ana_jellyfish and takes over 38
- There's got to be a better way...





Using rdrg_grid

- Used in set_vbc.F
- Need to pass it to set_vbc_tile routine – copy how bustr is passed
- Search on rdrg and add option:

#endif





Output

- It could be written to all output files along with the other grid variables unless NO_WRITE_GRID
- Put in def_info.F and wrt_info.F
- Still don't need to add to varinfo.dat



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Fourth Case

- Bio sediment variables for putting ammonia back into the water from falling detritus
- Code used in NEMURO for Bering Sea (from Enrique Curchitser)
- Time dependent want to store averages, write to stations, etc.





NEMURO_SED1

- We also tried a NEMURO_SED2, but it behaved badly
- Create two new variables in mod_ocean.F, PONsed and OPALsed
- 2-D so declare, allocate, initialize just like Hsbl
- This time we need to add it to varinfo.dat





Averages

- Add avgPONsed and avgOPALsed to mod_averages, in three places as usual
- The allocate and initialize are protected by tests on
 - Aout(idPONsed, ng)
 - Aout(idOPALsed)
- Aout is array of average output choices







Aout and Sout

 To properly set Aout(idPONsed), we need to add code to inp_par, matching code for Aout(idHsbl), Sout(idHsbl)





Set_avg.F

- Again, mimic the code for Hsbl to accumulate PONsed values into avgPONsed and avgOPALsed
 - Initialize at start of averaging duration
 - Add values until end of averaging duration
 - Scale by number of timesteps at end of averaging duration





I/O

- Add code to def_avg.F and wrt_avg.F to get averages output
- Add code to def_rst.F, wrt_rst.F and get_state.F to read it on restart
- Add code to def_station.F and wrt_station.F
- Add code to def_his.F and wrt_his.F but we neglected to check for Hout (idPONsed) in inp_par.F
- We also left it out of ana_biology.h





Last bits

- Don't forget to actually use and timestep your new variables!
- We need to add idOPALsed and idPONsed to mod_ncparam.F:
 - integer :: idOPALsed ! opal in sediment
 - integer :: idPONsed ! PON in sediment
- Files nemuro.diffs and drag.diffs are at:

http://www.arsc.edu/~kate/ROMS/HK

