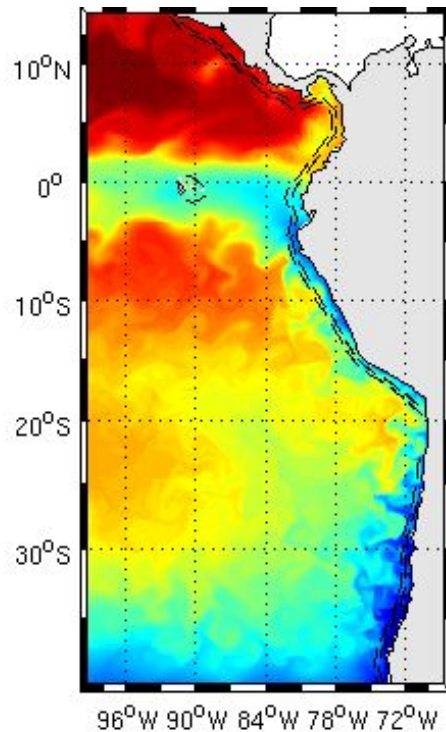
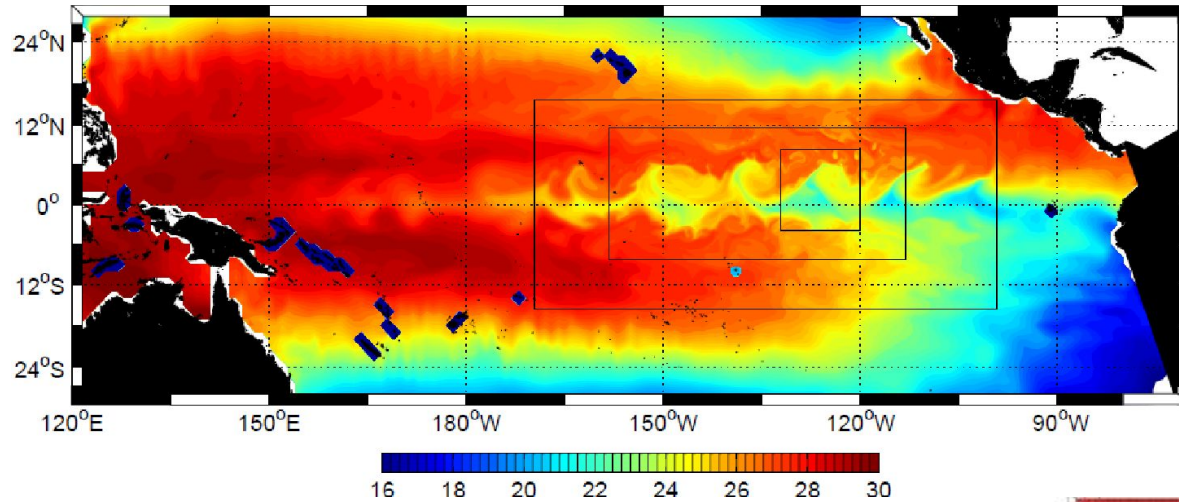
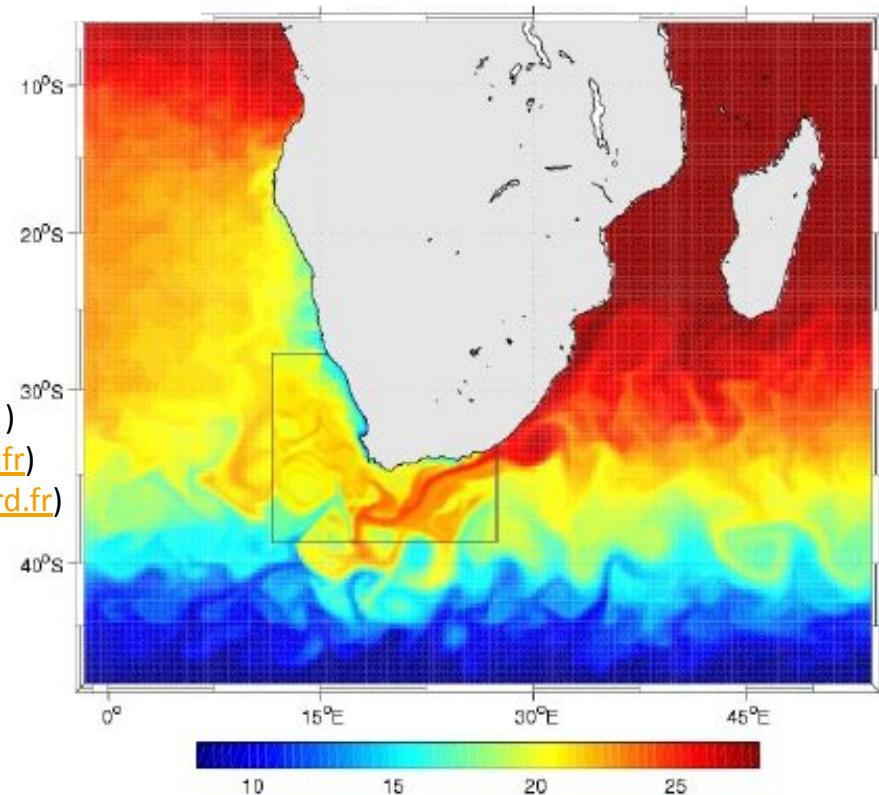


Introduction to CROCO / CROCO_TOOLS ocean modeling system



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- 1) Climatology simulation (simulation with no restart)
- 2) Inter annual simulation

Strategy to build a configuration

```
cd $HOME/TRAINING_2023
```

```
cp CROCO/croco/create_config.bash .
```

=> Edit create_config.bash
(e.g. with vi)

Note : 3 options of configuration
architectures available :

"all-dev": for dev of analytical tests

"all-prod": for production

climatological / interannual

**simulations => provides additional
scripts**

"all-prod-cpl" : for coupled simulations
(ww3, wrf)=> provides additional
scripts

=> **choose « all-prod »**

```
# croco source directory
#
CROCO_DIR=/home/user30/TRAINING_2023/CROCO/croco

# croco_tools directory
TOOLS_DIR=/home/user30/TRAINING_2023/CROCO/croco_tools

# Configuration name
# -----
MY_CONFIG_NAME=BENGUELA_LR

# Home and Work configuration directories
# -----
MY_CONFIG_HOME=/home/user30/TRAINING_2023/CONFIGS
MY_CONFIG_WORK=/home/user30/TRAINING_2023/CONFIGS

# Options of your configuration
# -----
## default option : all-dev for the usual ("all-in") archi
options=( all-prod )
```

Strategy to build a configuration

Run the create_config script:

```
./create_config.bash
```

=> It will create a BENGUELA_LR configuration in your CONFIGS directory

```
cd CONFIGS/BENGUELA_LR
```

```
ls -l
```

```
create_config.bash.bck  
CROCO_FILES  
CROCO_IN  
DATA  
myenv_mypath.sh  
myjob.sh  
myname1ist.sh  
PREPRO  
README_coupling_tools  
run_croco.bash  
run_croco_forecast.bash  
run_croco_inter.bash  
SCRIPTS_TOOLBOX  
submitjob.sh
```

Strategy to build a configuration

```
cd $HOME/TRAINING_2023/CONFIGS/BENGUELA_LR
```

General architecture of the configuration folder:

create_config.bash.bck ----- Backup of create_config script
myenv_mypath.sh ----- Environment file

PREPRO ----- Directory for preprocessing

CROCO_IN ----- Directory for CROCO compilation and settings

CROCO_FILES ----- Directory for CROCO inputs and outputs files

SCRATCH ----- Directory where the run is executed

run_croco.bash ----- Script for launching climatological runs

run_croco_inter.bash ----- Script for launching interannual runs

run_croco_forecast.bash ----- Script for launching forecast runs

mynamelist.sh
myjob.sh
submitjob.sh
SCRIPTS_TOOLBOX ----- Scripts for setting and launching simulation
with the coupling toolbox

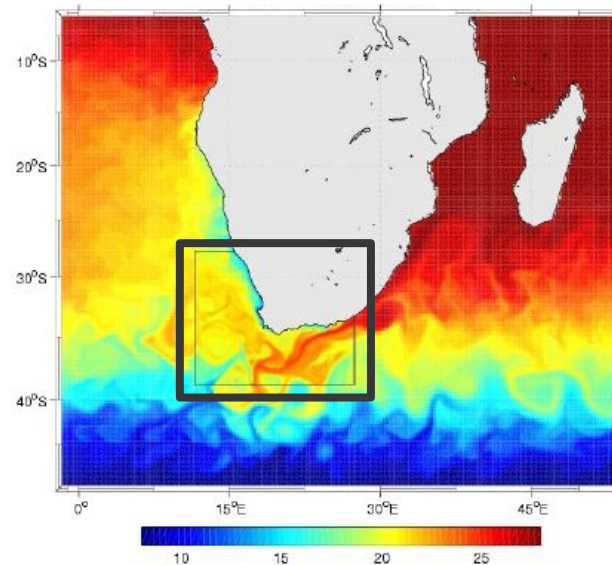
```
create_config.bash.bck
CROCO_FILES
CROCO_IN
DATA
myenv_mypath.sh
myjob.sh
mynamelist.sh
PREPRO
README_coupling_tools
run_croco.bash
run_croco_forecast.bash
run_croco_inter.bash
SCRIPTS_TOOLBOX
```

```
cd $HOME/TRAINING_2023/CONFIGS/BENGUELA_LR
```

```
cd PREPRO/CROCO
```

```
ls -l
```

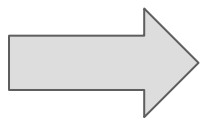
```
croctools_param.m  
find_childgrid_inparentgrid.m  
job_prepro_matlab.pbs  
make_grid_from_WRF.m  
oct_start.m  
prepro_cfsr.m  
prepro_soda.m  
README_nest_cpl  
README_preprocess_croco  
start.m  
town.dat
```



A regional realistic configuration , atmospheric forcing only
no tides
realistic bathymetry and stratification
horizontal resolution $dl = 1/3^\circ$ (33 km)
vertical resolution : $N=32$

```
theta_s = 7.;  
theta_b = 2.;  
hc      = 200.;  
vtransform = 2.; % s-coordinate type (1: old- ; 2: new- coordinates)  
           % ! take care to define NEW_S_COORD cpp-key in cppdefs.h
```

=> better discretization on surface layers



https://croco-ocean.gitlabpages.inria.fr/croco_doc/tutos/tutos.04.config.html

Croco Doc

Search docs

MODEL DOC

- 1. Governing Equations
- 2. Model variables
- 3. Grid and Coordinates
- 4. Numerics
- 5. Parametrizations
- 6. Parallelisation
- 7. Atmospheric Surface Boundary Layer
- 8. Open boundaries conditions
- 9. Rivers
- 10. Tides
- 11. Nesting Capabilities
- 12. Sediment and Biology models

TUTORIALS

- 1. System requirements
- 2. Download
- 3. Contents & Architecture
- 4. Summary of essential steps
- 5. Test Cases
- 6. Regional: Preparing your configuration**
- 7. Regional: Preprocessing (Matlab)
- 8. Compiling
- 9. Running the model
- 10. Increasing the resolution:

Docs » <no title> » 6. Regional: Preparing your configuration

6. Regional: Preparing your configuration

To prepare your configuration working directory, you can use the script `create_config.bash` provided in CROCO sources:

```
cp ~/croco/croco/create_config.bash ~/CONFIGS/.
```

Edit your paths and settings in `create_config.bash`:

```
#####  
# BEGIN USER MODIFICATIONS  
  
# Machine you are working on  
# Known machines: Linux DATARMOR IRENE JEANZAY  
# -----  
MACHINE="Linux"  
  
# CROCO parent directory  
# (where croco_tools directory and croco source directory can be found)  
# -----  
CROCO_DIR=~/.croco/croco  
TOOLS_DIR=~/.croco/croco_tools  
  
# Configuration name  
# -----  
MY_CONFIG_NAME=BENGUELA_LR  
  
# Home and work configuration directories
```


Documentation :

https://croco-ocean.gitlabpages.inria.fr/croco_doc/tutos/tutos.04.config.html

7. Regional: Preprocessing (Matlab)

CROCO preprocessing tools have been developed under Matlab software by IRD researchers (former Roms_tools). Note: These tools have been made to build easily regional configurations using climatological data. To use interannual data, some facilities are available (NCEP, CFSR, QuickScat data for atmospheric forcing, SODA and ECCO for lateral boundaries). However, to use other data, you will need to adapt the scripts. All utilities/toolbox requested for matlab crocotools programs are provided within the UTILITIES directory, or can be downloaded here: <http://www.croco-ocean.org/download/utilities/>

- 7.1. Contents of the `croco_tools`
- 7.2. Philosophy of the `croco_tools`
- 7.3. Climatological pre-processing
- 7.4. Interannual pre-processing

← Previous

Next →

`crocotools_param.m` is separated into several sections:

1 - Configuration parameters	used by <code>make_grid.m</code> (and others..)
2 - Generic file and directory names	need to match your work architecture
3 - Surface forcing parameters	used by <code>make_forcing.m</code> and by <code>make_bulk.m</code>
4 - Open boundaries and initial conditions parameters	used by <code>make_clim.m</code> , <code>make_biol.m</code> , <code>make_bry.m</code> <code>make_OGCM.m</code> and <code>make_OGCM_frcst.m</code>
5 - Parameters for tidal forcing	used by <code>make_tides.m</code>
6 - Reference date and simulation times	used for <code>make_tides</code> , <code>make_CFSR</code> (or <code>make_NCEP</code>), <code>make_OGC</code>
7 - Parameters for Interannual forcing	SODA, ECCO, CFSR, NCEP, ...
8 - Parameters for the forecast system	used by <code>make_forecast.m</code>
9 - Parameters for the diagnostic tools	used by scripts in <code>Diagnostic_tools</code>

Building the grid

Launch matlab session

`cd $HOME/TRAINING_2023/CONFIGS/BENGUELA_LR/PREPRO/CROCO`

`matlab -nodesktop`

Section 1 in crocotools_param.m

```
%
% 1 - Configuration parameters
% used by make_grid.m (and others..)
%
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
is octave=exist('octave_config_info');
%
% CROCO title names and directories
%
CROCO_title = 'Benguela Model';
CROCO_config = 'Benguela_LR';
%
% Grid dimensions:
%
lonmin = 8; % Minimum longitude [degree east]
lonmax = 22; % Maximum longitude [degree east]
latmin = -38; % Minimum latitude [degree north]
latmax = -26; % Maximum latitude [degree north]
%
% Grid resolution [degree]
%
dl = 1/3;
%
% Number of vertical Levels (! should be the same in param.h !)
%
N = 32;
```

```
%
% Vertical grid parameters (! should be the same in croco.in !)
%
theta_s = 7.;
theta_b = 2.;
hc = 200.;
vtransform = 2.; % s-coordinate type (1: old- ; 2: new- coordinates)
% ! take care to define NEW_S_COORD cpp-key in cppdefs.h
%
% Topography: choice of filter
%
topo_smooth = 1; % 1: old ; 2: new filter (better but slower)
%
% Minimum depth at the shore [m] (depends on the resolution,
% rule of thumb: dl=1, hmin=300, dl=1/4, hmin=150, ...)
% This affect the filtering since it works on grad(h)/h.
%
hmin = 75;    caution: if h<hmin, then h=hmin !!!
%
% Maximum depth at the shore [m] (to prevent the generation
% of too big walls along the coast)
%
hmax_coast = 500;
....
```

```
129 %
130 makeplot = 1; % 1: create graphics after each preprocessing step
131 %
```

CROCO

CONFIGS Directory for your Configurations

Fortran code
croco

Pre-post processing tools
croco_tools

AGRIF

MUSTANG

PISCES

XIOS

TEST_CASES

create_config.bash

CVT
K

SCRIPTS

MPI_NOLAND
_preprocessing

DOC_SPHINX

Sources *.F,
*.in

Configuration files
*.in

Prepare your run
directory

BENGUELA_LR

CROCO_IN

PREPRO Scripts for matlab
preprocessing/visualization

CROCO_FILES/
Input/output netcdf files

cppdef.h

Choices of
configurations
CPPkeys

param.h

Values of variables
for your
configuration

jobcomp

Compilation

Compile/

Directory for
compiling croco
(*.F, *.h, ...)

Your *.F, *.h
Files
you modified

croco.in

Parameters you
can modify
without
re-compiling

*his.nc,
*avg.nc

Output
netcdf files

Building the grid

make_grid.m , create the input file croco_grd.nc

Making the grid:
 /home1/datawork/gcambon/TRAINING_2019/CONFIGS/Run_BENGUELA_LR/CROCO_FILES/croco_grd.nc

Title: Benguela Model
 Resolution: 1/3 deg

Do you want to use interactive grid maker ?
 (e.g., for grid rotation or parameter adjustments) : y,[n] n

Create the grid file...

LLm = 41
 MMm = 42

Fill the grid file...

Compute the metrics...

Min dx=29.1913 km - Max dx=33.3244 km
 Min dy=29.2434 km - Max dy=33.1967 km

Fill the grid file...

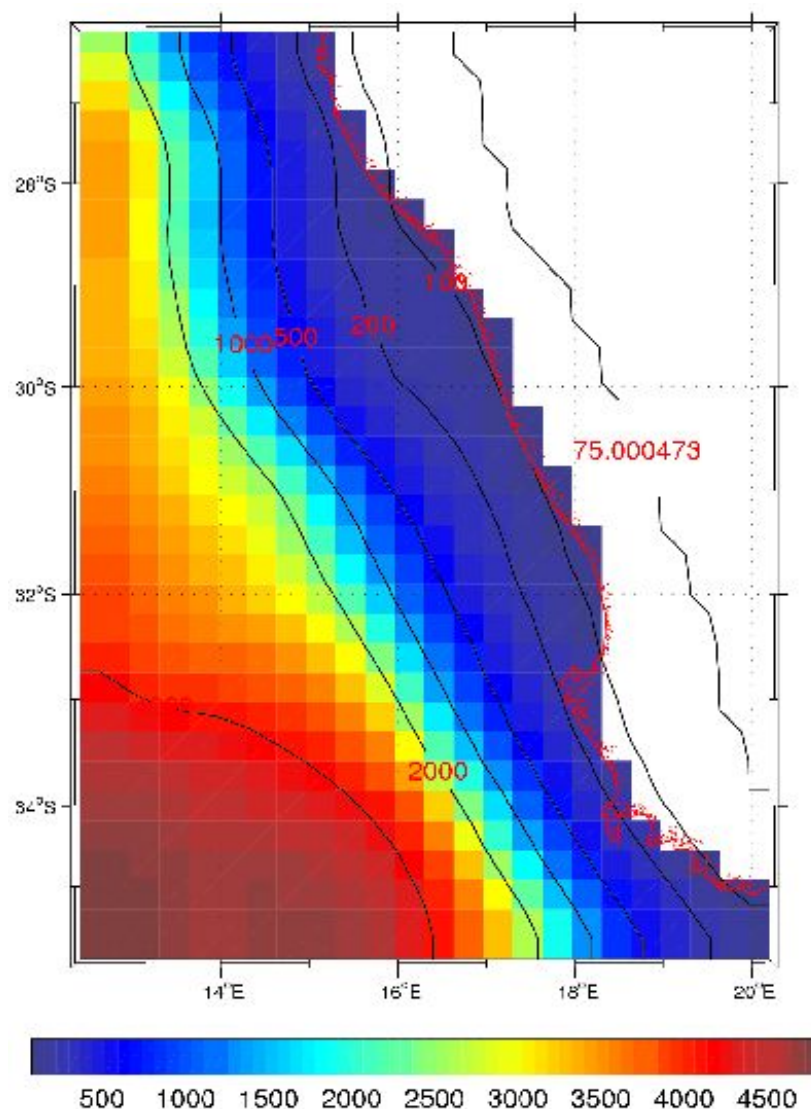
Add topography...

CROCO resolution : 31.3 km
 Topography data resolution : 3.42 km
 Topography resolution halved 4 times
 New topography resolution : 54.6 km

Processing coastline_l.mat ...

Do you want to use editmask ? y,[n]

...



Building the grid

Making the grid:
 /home1/datawork/gcambon/TRAINING_2019/CONFIGS/Run_BENGUELA_LR/CROCO_FILES/croco_grd.nc

Title: Benguela Model
 Resolution: 1/3 deg

Do you want to use interactive grid maker ?
 (e.g., for grid rotation or parameter adjustments) : y,[n] n

Create the grid file...

LLm = 41
 MMm = 42

same as in param.h !!!!

Fill the grid file...

Compute the metrics...

Min dx=29.1913 km - Max dx=33.3244 km

Min dy=29.2434 km - Max dy=33.1967 km

Fill the grid file...

Add topography...

CROCO resolution : 31.3 km

Topography data resolution : 3.42 km

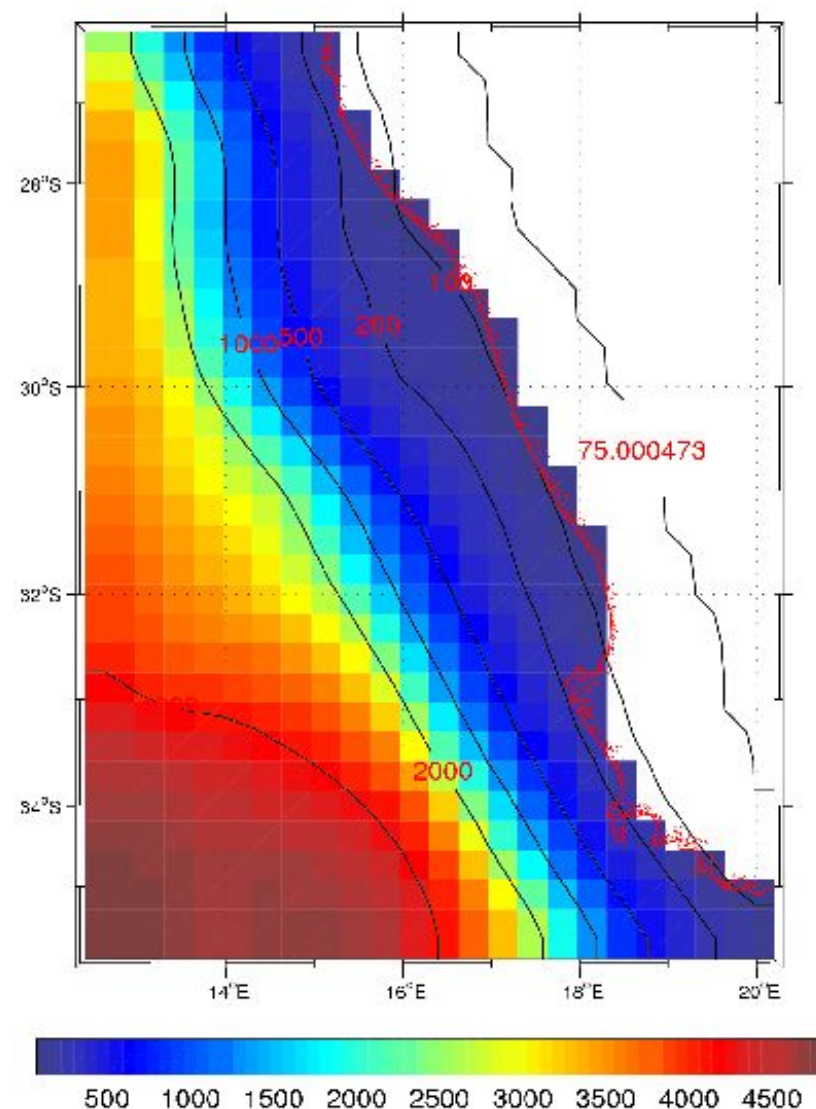
Topography resolution halved 4 times

New topography resolution : 54.6 km

Processing coastline_l.mat ...

Do you want to use editmask ? y,[n]

...



Build the atmospheric surface boundary forcing

make_forcing .m **create the input file croco_frc.nc**

Section 3 crocotools_param.m

```

%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
% COADS directory (for climatology runs)
%
coads_dir=[DATADIR,'COADS05/'];
%
% COADS time (for climatology runs)
%
coads_time=(15:30:345); % days: middle of each month
coads_cycle=360; % repetition of a typical year of 360 days
%
%coads_time=(15.2188:30.4375:350.0313); % year of 365.25 days in case
%coads_cycle=365.25; % interannual QSCAT winds
% % are used with clim. heat flux
%
% Pathfinder SST data used by pathfinder_sst.m
%
pathfinder_sst_name=[DATADIR,...
'SST_pathfinder/climato_pathfinder.nc'];
%
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%

```

Benguela Model

Read in the grid...

Create the forcing file...

Getting tau_x for time index 1

Getting tau_y for time index 1

Getting tau_x for time index 2

Getting tau_y for time index 2

Getting tau_x for time index 3

Getting tau_y for time index 3

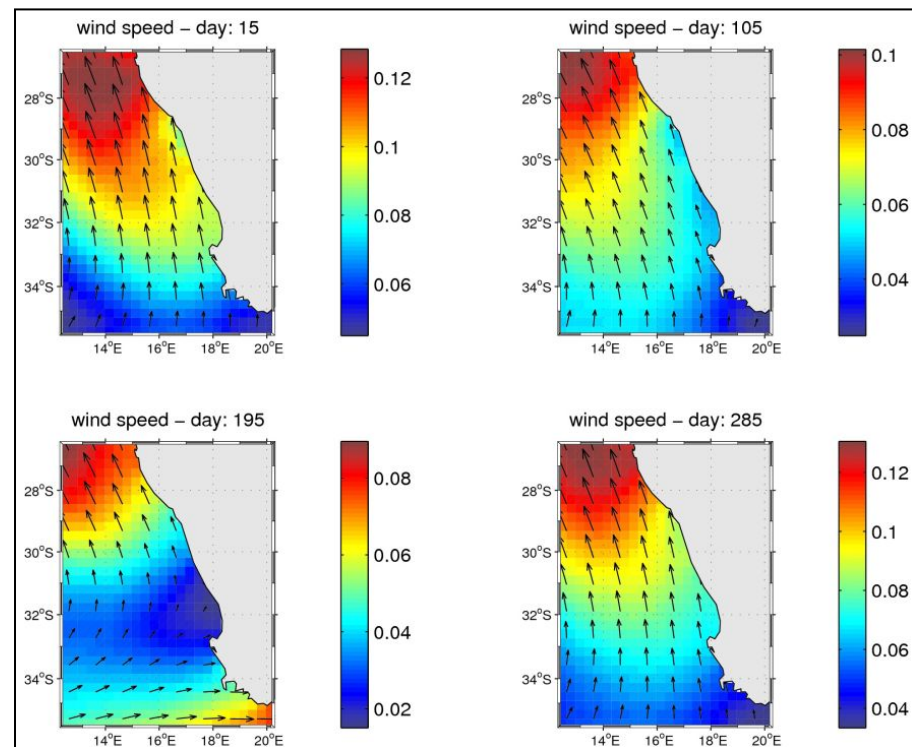
...

Getting shortrad for time index 9

Getting shortrad for time index 10

Getting shortrad for time index 11

Getting shortrad for time index 12



Build the atmospheric forcing: 2 options are available:

- create a forcing file with wind stress (zonal and meridional components), surface net heat flux, surface freshwater flux (E-P), solar shortwave radiation, SST, SSS, surface net sensitivity to SST (used for heat flux correction dQ_{dSST} for nudging towards model SST and model SSS)
- or create a bulk file which will be read during the run to perform bulk parameterization of the fluxes using COAMPS or Fairall 2003 formulation. This bulk file contains: surface air temperature, relative humidity, precipitation rate, wind speed at 10m, net outgoing longwave radiation, downward longwave radiation, shortwave radiation, surface wind speed (zonal and meridional components). It also contains surface wind stress (zonal and meridional components), but it is not requested and used in the model (except for specific debugging work). The bulk formulation computes its own wind stress.

```
make_bulk
```

or:

```
make_forcing
```

The settings relative to surface forcing are in section 3 of `crocoools_param.m`. In the case of climatological forcing, the variables are cycled. You can see that here, for the sake of simplicity, we are running the model on a repeating climatological year of 360 days.

A few figures illustrate the wind stress vectors and norm at 4 different periods of the year.

Note

`make_bulk` creates a forcing file that will be used with the cpp key `BULK_FLUX`, while `make_forcing` creates a forcing file containing wind stress directly and will be used when `undefined BULK_FLUX`. This second option is relevant if your atmospheric forcing comes from an atmospheric model with sufficient output frequency, or/and if you are comparing forced and coupled runs. Otherwise it is suggested to use `make_bulk`.

Build the initial/open boundary oceanic conditions

Section 4 in crocotools_param.m

```

%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
Open boundaries switches (! should be consistent with cppdefs.h !)
%
obc = [1 1 1 1]; % open boundaries (1=open , [S E N W])
%
% Level of reference for geostrophy calculation
%
zref = -1000;
%
% initial/boundary data options (1 = process)
% (used in make_clim, make_biol, make_bry,
% make_OGCM.m and make_OGCM_frcst.m)
%
makeini = 1; % initial data
makeclim = 1; % climatological data (for boundaries and nudging layers)
makebry = 1; % lateral boundary data
makenpzd = 0; % initial and boundary data for NChIPZD and N2ChIPZD2
models
makebioebus= 0; % initial and boundary data for BioEBUS model
makepisces = 0; % initial and boundary data for PISCES model
%
%
makeoa = 1; % oa data (intermediate file)
makeZbry = 1; % boundary data in Z coordinate (intermediate file)
insitu2pot = 1; % transform in-situ temperature to potential temperature
%
% Day of initialisation for climatology experiments (=0 : 1st january 0h)
%
tini=0;
%

```

```

%
% Pisces biogeochemical seasonal climatology
%
woapisces_dir = [DATADIR,'WOAPISCES/']; % only compatible with woa_dir
%
% Surface chlorophyll seasonal climatology (SeaWifs)
%
chla_dir=[DATADIR,'SeaWifs/'];
%
% Runoff monthly seasonal climatology (Dai and Trenberth)
%
global_clim_riverdir=[DATADIR,'RUNOFF_DAI/'];
global_clim_rivername=[global_clim_riverdir,'Dai_Trenberth_runoff_global
_clim.nc'];
%
% Set times and cycles for the boundary conditions:
% monthly climatology
%
woa_time=(15:30:345); % days: middle of each month
woa_cycle=360; % repetition of a typical year of 360 days
%

```

Build the initial oceanic conditions

> **make_ini.m** **create croco_ini.nc**

Benguela Model

Making initial file:

```
/home1/datawork/gcambon/TRAINING_2019/CONFIGS/Run_BENGUELA_LR/CROCO_FILES/croco_ini.nc
```

Title: Climatology

Creating the file :

```
/home1/datawork/gcambon/TRAINING_2019/CONFIGS/Run_BENGUELA_LR/CROCO_FILES/croco_ini.nc
VTRANSFORM = 2
```

Interpolations / extrapolations

Temperature...

Ext tracers: ro = 0 km - default value = NaN

```
ext_tracers_ini: time index: 1 of total: 12
ext_tracers_ini: horizontal interpolation of seasonal data
ext_tracers_ini: vertical interpolation
```

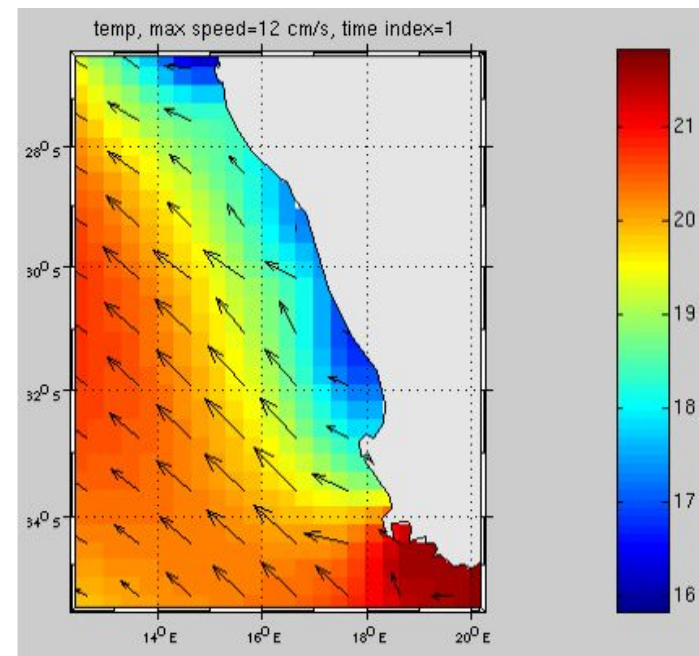
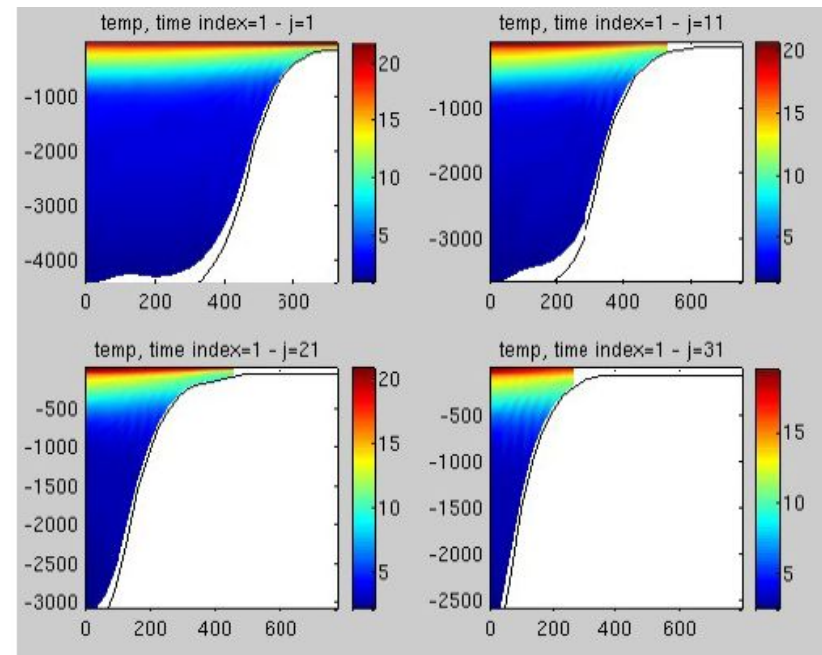
Salinity...

Ext tracers: ro = 0 km - default value = NaN

```
ext_tracers_ini: time index: 1 of total: 12
ext_tracers_ini: horizontal interpolation of seasonal data
ext_tracers_ini: vertical interpolation
```

Compute potential temperature from in-situ...

```
getpot: Time index: 1 of total: 1
```



> make_clim.m create croco_clm.nc

```
Creating the file :  
/home1/datawork/gcambon/TRAINING_2019/CONFIGS/Run_BENGUELA  
_LR/CROCO_FILES/croco_clm.nc
```

```
VTRANSFORM = 2
```

```
NetCDF_File:
```

```
'/home1/datawork/gcambon/TRAINING_2019/CONFIGS/Run_BENGUELA  
_LR/CROCO_FILES/croco_clm.nc'
```

```
nDimensions: 20
```

```
nVariables: 31
```

```
nGlobalAttributes: 0
```

```
RecordDimension: ''
```

```
nRecords: 0
```

```
Permission: 'clobber'
```

```
DefineMode: 'define'
```

```
FillMode: 'fill'
```

```
MaxNameLen: 0
```

```
Create the OA file...
```

```
Creating the file :
```

```
/home1/datawork/gcambon/TRAINING_2019/CONFIGS/Run_BENGUELA  
_LR/CROCO_FILES/croco_oa.nc
```

```
Horizontal extrapolations
```

```
Temperature...
```

```
Ext tracers: Roa = 0 km - default value = NaN
```

```
Ext tracers: horizontal interpolation of the annual data
```

```
Ext tracers: horizontal interpolation of the seasonal data
```

```
time index: 1 of total: 12
```

```
...
```

```
time index: 12 of total: 12
```

```
Salinity...
```

```
Ext tracers: Roa = 0 km - default value = NaN
```

```
Ext tracers: horizontal interpolation of the annual data
```

```
Ext tracers: horizontal interpolation of the seasonal data
```

```
time index: 1 of total: 12
```

```
...
```

```
Vertical interpolations
```

```
Temperature...
```

```
Time index: 1 of total: 12
```

```
...
```

```
Compute potential temperature from in-situ...
```

```
getpot: Time index: 1 of total: 12
```

```
...
```

```
Compute geostrophic currents
```

```
time index: 1 of total: 12
```

```
Flux correction : -7.2065
```

```
Mask: 16 iterations
```

```
...
```

> make_bry.m create croco_bry.nc

Create the boundary file...

Creating the file :

/home1/datawork/gcambon/TRAINING_2019/CONFIGS/Run_BENGUEL
A_LR/CROCO_FILES/croco_bry.nc

VTRANSFORM = 2

nc =

NetCDF_File:

'/home1/datawork/gcambon/TRAINING_2019/CONFIGS/Run_BENGUEL
A_LR/CROCO_FILES/croco_bry.nc'

nDimensions: 21

nVariables: 10

nGlobalAttributes: 0

RecordDimension: ''

nRecords: 0

Permission: 'lobber'

DefineMode: 'define'

FillMode: 'fill'

MaxNameLen: 0

Create the boundary Z-file...

Creating the file :

/home1/datawork/gcambon/TRAINING_2019/CONFIGS/Run_BENGUEL
A_LR/CROCO_FILES/croco_bry_Z.nc

Horizontal extrapolations

Processing southern boundary...

Temperature...

Ext tracers: horizontal interpolation of the annual data

Ext tracers: horizontal interpolation of the seasonal data

time index: 1 of total: 12

Summary for pre-processing input files [climato]

```
cd $HOME/TRAINING/CONFIGS/BENGUELA_LR/PREPRO/CROCO
```

matlab &

>> **start** : Add all the needed matlab path of the system

>> **make_grid**

=> creates CROCO_FILES/croco_grd.nc

- Horizontal grid : position of the grid points, size of the grid cells
- Bottom topography
- Land mask

>> **make_forcing**

=> create CROCO_FILES/croco_frc.nc

- Surface forcing : wind stress, surface heat flux, surface freshwater flux

>> **make_ini**

=> creates CROCO_FILES/croco_ini.nc

initial conditions : T, S, currents , SSH

>> **make_clim** (or **make_bry**)

=> creates CROCO_FILES/croco_clm.nc (or .croco_bry.nc)

- Lateral oceanic boundary conditions T, S, currents , SSH

make_forcing

to generate an atmospheric forcing file



make_clim , make_bry

- u= geostrophic currents+ eckman transport to generate an OBC forcing file
- T, S from COADS dataset

CROCO

CONFIGS Directory for your Configurations

Fortran code
croco

Pre-post processing tools
croco_tools

- AGRIF
- MUSTANG
- PISCES
- XIOS
- TEST_CASES
- create_config.bash
- CVT K
- SCRIPTS
- MPI_NOLAND_preprocessing
- DOC_SPHINX

Sources *.F,
*.in

Configuration files
*.in

Prepare your run directory

BENGUELA_LR

CROCO_IN

PREPRO Scripts for matlab preprocessing/visualization

CROCO_FILES/
Input/output netcdf files

cppdef.h
Choices of configurations
CPPkeys

param.h
Values of variables for your configuration

jobcomp
Compilation

Compile/
Directory for compiling croco
(*F, *.h, ...)

Your *.F, *.h
Files
you modified

croco.in
Parameters you can modify without re-compiling

*his.nc,
*avg.nc
Output netcdf files

Configuring the model CROCO

```
cd ~/TRAINING_2023/CONFIGS/BENGUELA_LR/CROCO_IN
```

```

#if defined REGIONAL
/*
!-----
!-----
!
! Applications */
!-----
! BASIC OPTIONS
!-----
!
!
*/
/* Configuration Name */
# define BENGUELA_LR
/* Parallelization */
# undef OPENMP
# define MPI
# undef MPI_NOLAND
/* I/O server */
# undef XIOS
# undef XIOS2
/* Non-hydrostatic option */
# undef NBQ
/* Nesting */
# undef AGRIF
# undef AGRIF_2WAY
/* OA and OW Coupling via OASIS (MPI) */
# undef OA_COUPLING
# undef OW_COUPLING
/* Wave-current interactions */
# undef MRL_WCI
/* Open Boundary Conditions */
# undef TIDES
# define OBC_EAST
# define OBC_WEST
# define OBC_NORTH
# define OBC_SOUTH

!-----
!-----
!
! Grid configuration */
!-----
!
!
! Model dynamics */
# define SOLVE3D
# define UV_COR
# define UV_ADV

/* dedicated croco.log file */
# undef LOGFILE
/* Calendar */
# undef USE_CALENDAR
/*!
!-----
! PRE-SELECTED OPTIONS
!
! ADVANCED OPTIONS ARE IN CPPDEFS_DEV.H
!-----
*/
/* Parallelization */
# ifdef MPI
# undef PARALLEL_FILES
# endif
# undef NC4PAR
# undef AUTOTILING
/* Non-hydrostatic options */
# ifdef NBQ
# define W_HADV_TVD
# define W_VADV_TVD
# endif

```

cppdefs.h file (1/2)

Configuring the model CROCO

```

/* Equation of State */
# define SALINITY
# define NONLIN_EOS
    /* Lateral Momentum Advection (default UP3) */
# define UV_HADV_UP3
# undef UV_HADV_UP5
# undef UV_HADV_WENO5
# undef UV_HADV_TVD
    /* Lateral Explicit Momentum Mixing */
# undef UV_VIS2
# ifdef UV_VIS2
# define UV_VIS_SMAGO
# endif
    /* Vertical Momentum Advection */
# define UV_VADV_SPLINES
# undef UV_VADV_WENO5
# undef UV_VADV_TVD
    /* Lateral Tracer Advection (default UP3) */
# undef TS_HADV_UP3
# define TS_HADV_RSUP3
# undef TS_HADV_UP5
# undef TS_HADV_WENO5
    /* Lateral Explicit Tracer Mixing */
# undef TS_DIF2
# undef TS_DIF4
# undef TS_MIX_S
    /* Vertical Tracer Advection */
# undef TS_VADV_SPLINES
# define TS_VADV_AKIMA
# undef TS_VADV_WENO5
    /* Sponge layers for UV and TS */
# define SPONGE

    /* Semi-implicit Vertical Tracer/Mom
    Advection */
# undef VADV_ADAPT_IMP
    /* Bottom friction in fast 3D step
    */
# undef BSTRESS_FAST
    /* Vertical Mixing */
# undef BODYFORCE
# undef BVF_MIXING
# define LMD_MIXING
# undef GLS_MIXING
# ifdef LMD_MIXING
# define LMD_SKPP
# define LMD_BKPP
# define LMD_RIMIX
# define LMD_CONVEC
# undef LMD_DDMIX
# define LMD_NONLOCAL
# undef MLCONVEC
# endif
    /* Surface Forcing */
# undef BULK_FLUX
# ifdef BULK_FLUX
# define BULK_FAIRALL
# define BULK_LW
# define BULK_EP
# define BULK_SMFLUX
# undef SST_SKIN
# undef ANA_DIURNAL_SW
# undef ONLINE# ifdef ONLINE
# undef AROME
# undef ERA_ECMWF
# endif

# undef READ_PATM
# ifdef READ_PATM
# define OBC_PATM
# endif
# else
# define QCORRECTION
# define SFLX_CORR
# undef SFLX_CORR_COEF
# define ANA_DIURNAL_SW
# endif
# undef SMFLUX_CFB
# undef SEA_ICE_NOFLUX
    /* Wave-current interactions */
# ifdef OW_COUPLING
# define MRL_WCI
# define BBL
# endif
# ifdef MRL_WCI
# ifndef OW_COUPLING
# define WAVE_OFFLINE
# undef WKB_WWAVE
# endif
# undef WAVE_ROLLER
# define WAVE_STREAMING
# define WAVE_FRICTION
# define WAVE_RAMP
# ifdef WKB_WWAVE
# undef WKB_OBC_NORTH
# undef WKB_OBC_SOUTH
# undef WKB_OBC_WEST
# undef WKB_OBC_EAST
# endif
# endif

```

cppdefs.h file (2/2)

Configuring the model CROCO

Array dimensions and parameters (param.h)

```

!-----
! Dimensions of Physical Grid and array dimensions
!-----
!
! LLm,MMm  Number of the internal points of the PHYSICAL grid.
!           in the XI- and ETA-directions [physical side boundary
!           points and peroodic ghost points (if any) are excluded].
!
! Lm,Mm    Number of the internal points [see above] of array
!           covering a Message Passing subdomain. In the case when
!           no Message Passing partitioning is used, these two are
!           the same as LLm,MMm.
!
! N        Number of vertical levels.
!
integer LLm,Lm,MMm,Mm,N, LLm0,MMm0
#if defined AGRIF
integer LLmm2, MMmm2
#endif
#if defined BASIN
parameter (LLm0=60, MMm0=50, N=10)
#elif defined CANYON_A
parameter (LLm0=65, MMm0=48, N=16)
#elif defined CANYON_B
parameter (LLm0=66, MMm0=48, N=16)
#elif defined EQUATOR
parameter (LLm0=40, MMm0=32, N=32) ! 100 km resolution

```

```

#elif defined KH_INST
....
# elif defined BENGUELA_LR
parameter (LLm0=41, MMm0=42,
N=32) ! BENGUELA_LR
# elif defined BENGUELA_HR
parameter (LLm0=83, MMm0=85,
N=32) ! BENGUELA_HR
# elif defined BENGUELA_VHR
parameter (LLm0=167, MMm0=170,
N=32) ! BENGUELA_VHR
# elif defined MENOR
parameter (LLm0=1059, MMm0=447,
N=40) ! MENOR
# elif defined SEINE
parameter (LLm0=411, MMm0=181,
N=20) ! SEINE
# else
parameter (LLm0=94, MMm0=81,
N=40)
# endif
#else
parameter (LLm0=xx, MMm0=xx, N=xx)
#endif
...
!
...

```

Domain decomposition for parallelization : param.h

```

!
!-----
! MPI related variables
!-----
!
integer Lmmpi,Mmmpi,iminmpi,imaxmpi,jminmpi,jmaxmpi
common /comm_setup_mpi1/ Lmmpi,Mmmpi
common /comm_setup_mpi2/ iminmpi,imaxmpi,jminmpi,jmaxmpi
!
! Domain subdivision parameters
! =====
!
! NPP      Maximum allowed number of parallel threads;
! NSUB_X,NSUB_E Number of SHARED memory subdomains in XI- and
!                      ETA-directions;
! NNODES   Total number of MPI processes (nodes);
! NP_XI, NP_ETA Number of MPI subdomains in XI- and ETA-directions;
!
integer NSUB_X, NSUB_E, NPP

#ifdef MPI
integer NP_XI, NP_ETA, NNODES
parameter (NP_XI=1, NP_ETA=2,
NNODES=NP_XI*NP_ETA)
parameter (NPP=1)
parameter (NSUB_X=1, NSUB_E=1)
#elif defined OPENMP
parameter (NPP=4)
# ifdef AUTOTILING
common/distrib/NSUB_X, NSUB_E
# else
parameter (NSUB_X=1, NSUB_E=NPP)
# endif
#else
parameter (NPP=1)
# ifdef AUTOTILING
common/distrib/NSUB_X, NSUB_E
# else
parameter (NSUB_X=1, NSUB_E=NPP)
# endif
#endif
....

```

CROCO

CONFIGS Directory for your Configurations

Fortran code
croco

Pre-post processing tools
croco_tools

AGRIF

MUSTANG

PISCES

XIOS

TEST_CASES

create_config.bash

CVT
K

SCRIPTS

MPI_NOLAND_preprocessing

DOC_SPHINX

Sources *.F,
*.in

Configuration files
*.in

Prepare your run directory

BENGUELA_LR

CROCO_IN

PREPRO Scripts for matlab preprocessing/visualization

CROCO_FILES/
Input/output netcdf files

cppdef.h

Choices of configurations
CPPkeys

param.h

Values of variables for your configuration

jobcomp
Compilation

Compile/
Directory for compiling croco
(*.F, *.h, ...)

Your *.F, *.h
Files
you modified

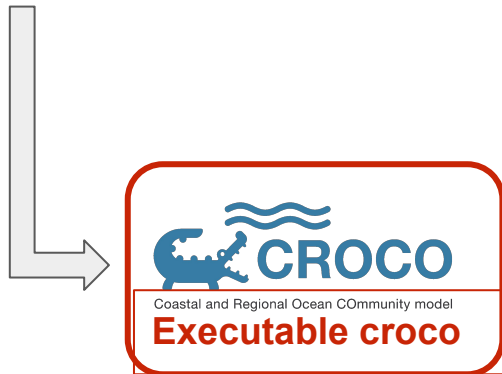
croco.in

Parameters you can modify without re-compiling

*his.nc,
*avg.nc
Output netcdf files

```
cd $HOME/TRAINING_2023/CONFIGS/BENGUELA_LR/CROCO_IN
```

```
./jobcomp.bash
```



CROCO

CONFIGS Directory for your Configurations

Fortran code
croco

Pre-post processing tools
croco_tools

AGRIF

MUSTANG

PISCES

XIOS

TEST_CASES

create_config.bash

CVT
K

SCRIPTS

MPI_NOLAND
_preprocessing

DOC_SPHINX

Sources *.F,
*.in

Configuration files
*.in

Prepare your run
directory

BENGUELA_LR

CROCO_IN

PREPRO Scripts for matlab
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for your
configuration

jobcomp

Compilation

Compile/

Directory for
compiling croco
(*.F, *.h, ...)

Your *.F, *.h
Files
you modified

croco.in

Parameters you
can modify
without
re-compiling

*his.nc,
*avg.nc

Output
netcdf files

Copy an example of croco.in suitable for Regional configuration:

```
cd /home/userX/TRAINING_2023
cd CONFIGS/BENGUELA_LR/CROCO_IN
cp ~/TRAINING_2023/CROCO/croco/OCEAN/croco.in .
```

```
grid: filename
    ../CROCO_FILES/croco_grd.nc
forcing: filename
    ../CROCO_FILES/croco_frc.nc
bulk_forcing: filename
    ../CROCO_FILES/croco_blk.nc
climatology: filename
    ../CROCO_FILES/croco_clm.nc
boundary: filename
    ../CROCO_FILES/croco_bry.nc
initial: NRREC / filename
    1
    ../CROCO_FILES/croco_ini.nc
restart: NRST, NRPFRST / filename
    720 -1
    ../CROCO_FILES/croco_rst.nc
history: LDEFHIS, NWRT, NRPFHIS / filename
    T 72 0
    ../CROCO_FILES/croco_his.nc
averages: NTSAVG, NAVG, NRPF AVG / filename
    1 72 0
    ../CROCO_FILES/croco_avg.nc
```

change CROCO_FILES to ../CROCO_FILES everywhere



Documentation :

https://croco-ocean.gitlabpages.inria.fr/croco_doc/tutos/tutos.08.compil.html

variables or diagnostics in output netcdf files

```

title:
  BENGUELA TEST MODEL
time_stepping: NTIMES dt[sec] NDTFAST NINFO
  8640 1200 60 1
time_stepping_nbq: NDTNBQ CSOUND_NBQ VISC2_NBQ
  1 1000 0.01
S-coord: THETA_S, THETA_B, Hc(m)
  7.0d0 2.0d0 200.0d0
run_start_date:
  01/04/2014 00:00:00
run_end_date:
  01/01/2016 00:00:00
output_time_steps: DT_HIS(H), DT_AVG(H), DT_RST(H)
  1 6 12
grid: filename
  CROCO_FILES/croco_grd.nc
forcing: filename
  CROCO_FILES/croco_frc.nc
bulk_forcing: filename
  CROCO_FILES/croco_blk.nc
climatology: filename
  CROCO_FILES/croco_clm.nc
boundary: filename
  CROCO_FILES/croco_bry.nc
initial: NRREC filename
  1
  CROCO_FILES/croco_ini.nc
restart: NRST, NRPFRST / filename
  2160 -1
  CROCO_FILES/croco_rst.nc
history: LDEFHIS, NWRT, NRPFHIS / filename
  T 72 0
  CROCO_FILES/croco_his.nc
averages: NTSAVG, NAVG, NRPAVG / filename
  1 72 0
  CROCO_FILES/croco_avg.nc
    
```

same as in the crocotools_param.m

```

primary_history_fields: zeta UBAR VBAR U V wrtT(1:NT)
  T T T T T 30*T
auxiliary_history_fields: rho Omega W Akv Akt Aks Visc3d Diff3d HBL HBBL
  Bostr Wstr Ustr Vstr Shfl Swfl rsw rlw lat sen HEL
  F F T F T F F F T T T T T T T 10*T
gls_history_fields: TKE GLS Lscale
  T T T
primary_averages: zeta UBAR VBAR U V wrtT(1:NT)
  T T T T T 30*T
auxiliary_averages: rho Omega W Akv Akt Aks Visc3d Diff3d HBL HBBL Bostr
  Wstr Ustr Vstr Shfl Swfl rsw rlw lat sen HEL
  F T T F T F F F T T T T T T T 10*T
gls_averages: TKE GLS Lscale
  T T T
rho0:
  1025.d0
lateral_visc: VISC2, VISC4 [m^2/sec for all]
  0. 0.
tracer_diff2: TNU2(1:NT) [m^2/sec for all]
  30*0.d0
tracer_diff4: TNU4(1:NT) [m^4/sec for all]
  30*0.d11
vertical_mixing: Akv_bak, Akt_bak [m^2/sec]
  0.d0 30*0.d0
bottom_drag: RDRG [m/s], RDRG2, Zob [m], Cdb_min, Cdb_max
  3.0d-04 0.d-3 0.d-3 1.d-4 1.d-1
gamma2:
  1.d0
sponge: X_SPONGE [m], V_SPONGE [m^2/sec]
  XXX XXX
nudg_cof: TauT_in, TauT_out, TauM_in, TauM_out [days for all]
  1. 360. 3. 360.
    
```

mpirun -np 2 ./croco croco.in

Southern Benguela

480 ntimes Total number of timesteps for 3D equations.

5400.00 dt Timestep [sec] for 3D equations

60 ndtfast Number of 2D timesteps within each 3D step.

1 ninfo Number of timesteps between runtime diagnostics.

...

Activated C-preprocessing Options:

...

Spherical grid detected

hmin	hmax	grdmin	grdmax	Cu_min	Cu_max
75.000000	4803.032721	.301836927E+05	.331215714E+05	0.12176008	0.91533005

volume=9.523986093261087500000E+14 open_cross=6.104836888312444686890E+09

...

MAIN: started time-stepping.

STEP	time[DAY]	KINETIC_ENRG	POTEN_ENRG	TOTAL_ENRG	NET_VOLUME	trd
0	0.00000	0.000000000E+00	2.1475858E+01	2.1475858E+01	9.5239861E+14	0
1	0.06250	1.306369099E-04	2.1476230E+01	2.1476361E+01	9.5239208E+14	0

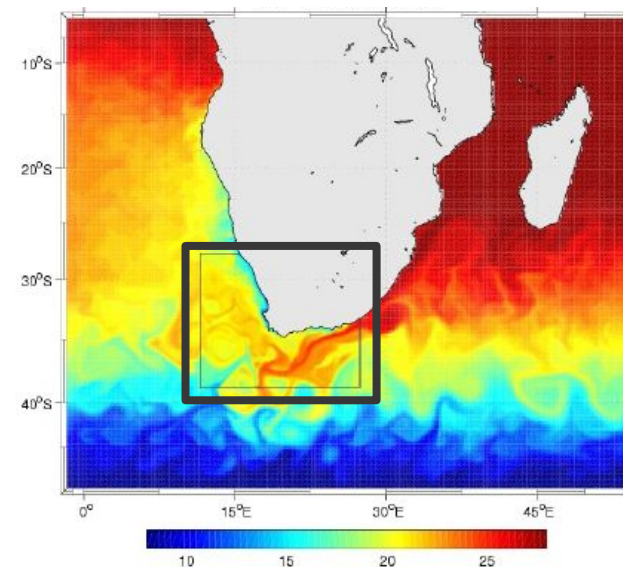
...

Courant number:
evaluation of the CFL criterion:
 $dx/dt >$ fastest waves (here gravity waves).
Cu_max < 1 !!!

- 1) Crocotools presentation for preprocessing
- 2) Climatology simulation (simulation with no restart)
- 3) **Inter annual simulation**

More realistic regional configuration (BENGULA_INTER)

- Realistic Bathymetry
- Initial conditions taken from OGCM model (mercator)
- Open boundaries OGCM forcing (monthly mercator outputs)
- Atmospheric forcing from ERA5 atmospheric model (hourly outputs) —> but the interpolation will be done online in the CROCO model
- No tides forcing



MAIN STEPS

- **Preprocessing :**
 - * Grid generation (make_grid)
 - * initial file and boundary files (make_OGCM_mercator)
 - * **the forcing file will be done online in CROCO**
- ⇒ **but before we have to adapt the croctools_param.m**
- **CROCO configuration**
 - * Files needed to be adapted: param.h, cppdefs.h, croco.in
 - * **Bash RUN : croco_inter.bash**
- **Visualisation : lunch croco_gui** in PREPRO directory

inter annual simulation

- 1) Create a configuration with create_config
- 2) Modify crocotools_param.m file in your configuration (in PREPRO/CROCO)

```

CROCO_title = 'Benguela Model';
CROCO_config = 'Benguela_LR';
%
% Grid dimensions:
%
lonmin = 8; % Minimum longitude [degree east]
lonmax = 22; % Maximum longitude [degree east]
latmin = -38; % Minimum latitude [degree north]
latmax = -26; % Maximum latitude [degree north]
%
% Grid resolution [degree]
%
dl = 1/3;
%
% Number of vertical Levels (! should be the same in par

```

```

makeini = 1;
makeclim = 0;
makebry = 1;
makenpzd = 0;
makebioebus = 0;
makepisces = 0;
makequota = 0;

```

```

129 %
130 makeplot = 1; % 1: create graphics after each preprocessing step
131 %

```

```

332 Ymin = 2013; % first forcing year
333 Ymax = 2013; % last forcing year
334 Mmin = 1; % first forcing month
335 Mmax = 3; % last forcing month

```

```

%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
%
% 7 - Parameters for Interannual forcing (SODA, ECCO, CFSR, NCEP, ...)
%
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
%
Download_data = 0; % Get data from OPENDAP sites
level = 0; % AGRIF level; 0 = parent grid
%

```

```
%-----  
% Options for make_ERA5  
%-----  
%  
%ERA5_dir = [FORC_DATA_DIR,'ERA5 ',CROCO_config,'/']; %  
ERA5_dir = '/home/COMMONDATA/data_tutos/ERA5_GLOB/';  
My_ERA5_dir = [FORC_DATA_DIR,'ERA5_native_',CROCO_config,'/']; %  
%  
itolap era5 = 2; %
```

ligne 412

```
423 OGCM = 'mercator'; % Select OGCM: SODA, ECCO, mercator  
424 %  
425 OGCM_dir = [FORC_DATA_DIR,OGCM,'_',CROCO_config,'/']; % OGCM data dir.  
426 %  
427 OGCM_dir = '/home/COMMONDATA/data_tutos/MERCATOR_GLOB/';  
428 %
```

ERA5_dir= '/home/COMMONDATA/data_tutos/ERA5_GLOB/';
OGCM='mercator';
OGCM_dir='/home/COMMONDATA/data_tutos/MERCATOR_GLOB';

inter annual simulation

```
cd $HOME/TRAINING_2023/CONFIGS/XXX/PREPRO/CROCO
```

matlab &

```
>> start : Add all the needed matlab path of the system
```

```
>> make_grid
```

```
=> creates CROCO_FILES/croco_grd.nc
```

- Horizontal grid : position of the grid points, size of the grid cells
- Bottom topography
- Land mask

```
>> make_forcing
```

```
=> create CROCO_FILES/croco_frc.nc
```

- ~~• Surface forcing : wind stress, surface heat flux, surface freshwater flux~~

```
>> make_ini
```

```
=> creates CROCO_FILES/croco_ini.nc
```

```
initial conditions : T, S, currents , SSH
```

```
>> make_OGCM_mercator
```

```
=> creates CROCO_FILES/croco_clm.nc (or .croco_bry.nc)
```

- Lateral oceanic boundary conditions T, S, currents , SSH

make_era5 will create atmospheric forcings BUT it takes time

IDEA => use interpolation of atmospheric forcings online during CROCO execution

CROCO_IN/cppdefs.h et param.h

```
72 # undef OPENMP  
73 # define MPI  
74
```

activate **MPI** parallelisation cpp option

```
153 # define BULK_FLUX  
154 # ifdef BULK_FLUX  
155 # undef BULK_ECUMEV0  
156 # undef BULK_ECUMEV6  
157 # undef BULK_WASP  
158 # define BULK_GUSTINESS  
159 # define BULK_LW  
160 # undef SST_SKIN  
161 # undef ANA_DIURNAL_SW  
162 # define ONLINE  
163 # ifdef ONLINE  
164 # undef AROME  
165 # define ERA_ECMWF  
166 # endif  
167 # undef READ_PATM  
168 # ifdef READ_PATM  
169 # define OBC_PATM
```

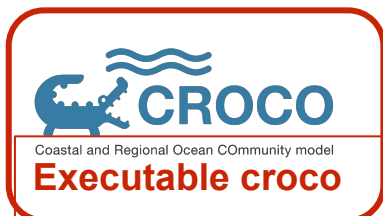
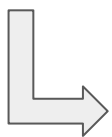
activate **BULK_FLUX** , **ONLINE** , **ERA_EMWF** cpp options for online interpolation of ERA5 atmospheric forcing

MODIFY

CROCO_IN/**param.h** : modify size grid if needed
modify number of procs => 2 procs

COMPILE

./jobcomp.bash



MODIFY

croco_inter.in

```
online:  byear  bmonth recordsperday byearend bmonthend / data path
         NYONLINE NMONLINE 24          2013  4
         /home/COMMONDATA/data_tutos/ERA5_GLOB/
```

byear : beginning year = NYONLINE

bmonth : beginning month = NMONLINE

recordsperday : records per day = 24

byearend: ending year = 2013

bmonthend : ending month = 4

—> see run_croco_inter.bash


```
30 NBPROCS=2
31
32 # command for running the mode : ./ for sequ
33 # WARNING: for mpi run command, it is needed
34 #RUNCMD='./'
35 RUNCMD="mpirun -np $NBPROCS "
36 #RUNCMD="$MPI_LAUNCH "
```

```
55 # Atmospheric surface forcing dataset u
56 ATMOS_BULK=ERA5
57 # Atmospheric surface forcing dataset u
58 ATMOS_FRC=QSCAT
59 # Oceanic boundary and initial dataset
60 OGCM=mercator
61 # Dynoff dataset (Dye and Trochardt
```

BULK_FILES=0 !!!!!

```
74 # Start and End year
75 NY_START=2013
76 NY_END=2013
77 # Start and End month
78 NM_START=1
79 NM_END=3
80 # Set month format at 1 or 2 dig
```

./run_croco_inter.bash

cd ~/TRAINING_2023/CONFIGS/XXX

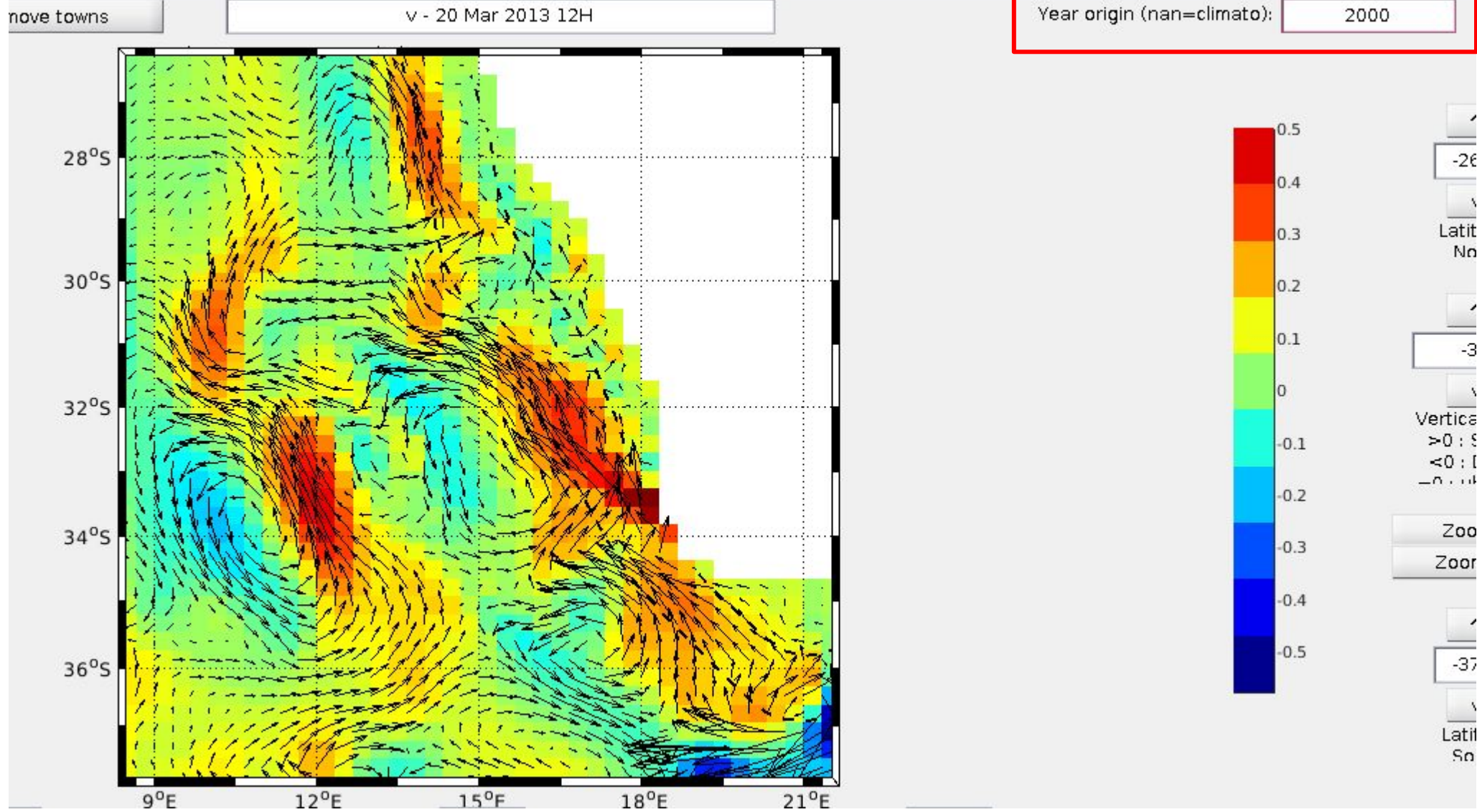
run the script for inter annual: **./run_croco_inter.bash**

```
user30@croco:~/TRAINING_2023/CONFIGS/CONFIG_INTER$ bash run_croco_inter.bash
Getting croco from /home/user30/TRAINING_2023/CONFIGS/CONFIG_INTER/CROCO_IN
Getting AGRIF_FixedGrids.in from /home/user30/TRAINING_2023/CONFIGS/CONFIG_INTER/CROCO_IN
Getting croco_grd.nc from /home/user30/TRAINING_2023/CONFIGS/CONFIG_INTER/CROCO_FILES
Getting croco_inter.in from /home/user30/TRAINING_2023/CONFIGS/CONFIG_INTER/CROCO_IN
Getting croco_ini_mercator_Y2013M01.nc from /home/user30/TRAINING_2023/CONFIGS/CONFIG_INTER/CROCO_FILES
Computing YEAR 2013 MONTH 01
Getting croco_blk_ERA5_Y2013M01.nc from /home/user30/TRAINING_2023/CONFIGS/CONFIG_INTER/CROCO_FILES
Getting croco_bry_mercator_Y2013M01.nc from /home/user30/TRAINING_2023/CONFIGS/CONFIG_INTER/CROCO_FILES
YEAR = 2013 MONTH = 1 DAYS = 31 DT = 3600 NTIMES = 0
No exact restart
set NUMRECINI = 1

Writing in croco_inter.in
USING DT          = 3600
USING NFAST       = 60
USING NUMTIMES    = 744
USING NUMAVG      = 72
USING NUMHIS      = 744
USING NUMRST      = 744
USING NUMRECINI   = 1

Computing for Y2013M01
sam. 09 sept. 2023 21:43:22 UTC
sam. 09 sept. 2023 21:43:48 UTC
Test croco_Y2013M01.out
All good
XXXXXXXXXX
Computing YEAR 2013 MONTH 02
```

OUTPUTS are saved in **SCRATCH**
directory



In many studies, there is a need for long simulations : spin-up , statistical equilibrium.

For regional models, 10 years appears to be a reasonable model duration.

In this case, it is easier to do several simulations of 1 month, using restart at the end of each month to start the next month.

run_croco.bash for climatological forcing or **run_croco_inter.bash** for interannual forcing do it **automatically**

- It get the grid, the initial file and the boundary file
- In case of a interannual forcing, we have realistic forcing, segmented month by month, from various atmospheric/oceanic reanalysis [for example : croco_ini_SODA_Y2004M1.nc, croco_frc_CFSR_Y2004M1.nc or croco_bry_ECCO_Y2004M1.nc]
- It runs the model for 1 month
- It store the files in a specific for in the directory SCRATCH
 - For climatological forcing : roms_avg_Y4M3.nc (i.e march of year 4)
 - For interannual forcing : roms_avg_Y2004M3.nc (i.e march of year 2004)
- It replace the initial file by the restart file (croco_rst.nc) which has been generated at the end of the month.
- It relaunch the model for the next month

Documentation :

https://croco-ocean.gitlabpages.inria.fr/croco_doc/tutos/tutos.10.run.inter.html

11. Running with interannual forcing

11.1. Run after classical interannual pre-processing

Before running you should prepare your interannual inputs files following the Interannual Preprocessing tutorial.

To run a plurimonth simulation, we provide the following scripts in `~/croco/croco/SCRIPTS/Plurimonths_scripts` :

- `run_croco.bash` : Plurimonth run with climatological forcing

Documentation :

https://croco-ocean.gitlabpages.inria.fr/croco_doc/tutos/tutos.10.run.inter.html

11. Running with interannual forcing

11.1. Run after classical interannual pre-processing

Before running you should prepare your interannual inputs files following the Interannual Preprocessing tutorial.

To run a plurimonth simulation, we provide the following scripts in `~/croco/croco/SCRIPTS/Plurimonths_scripts` :

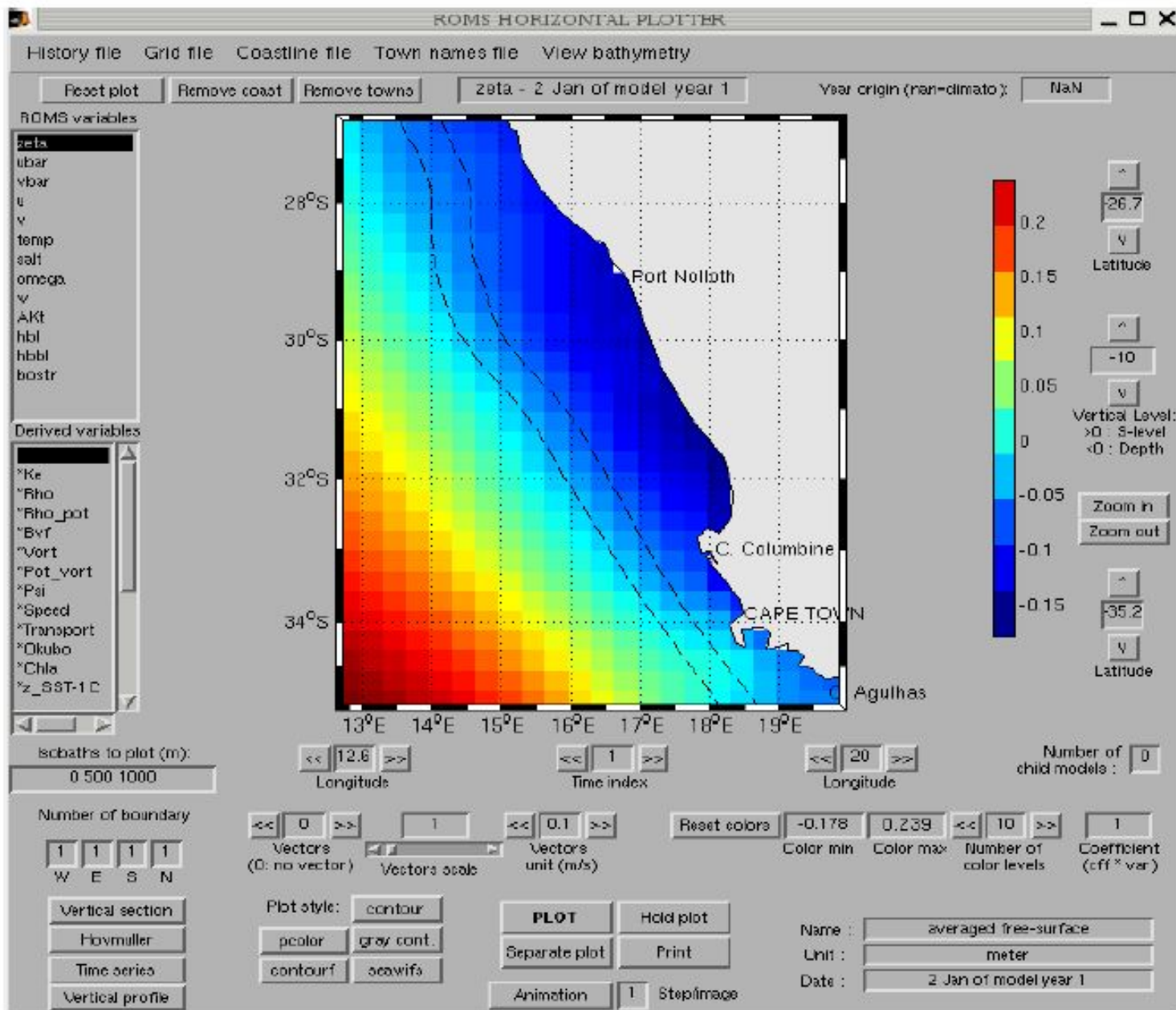
- `run_croco_inter.bash` : Plurimonth run with interannual forcing

Documentation :

https://croco-ocean.gitlabpages.inria.fr/croco_doc/tutos/tutos.14.visu.matlab.html

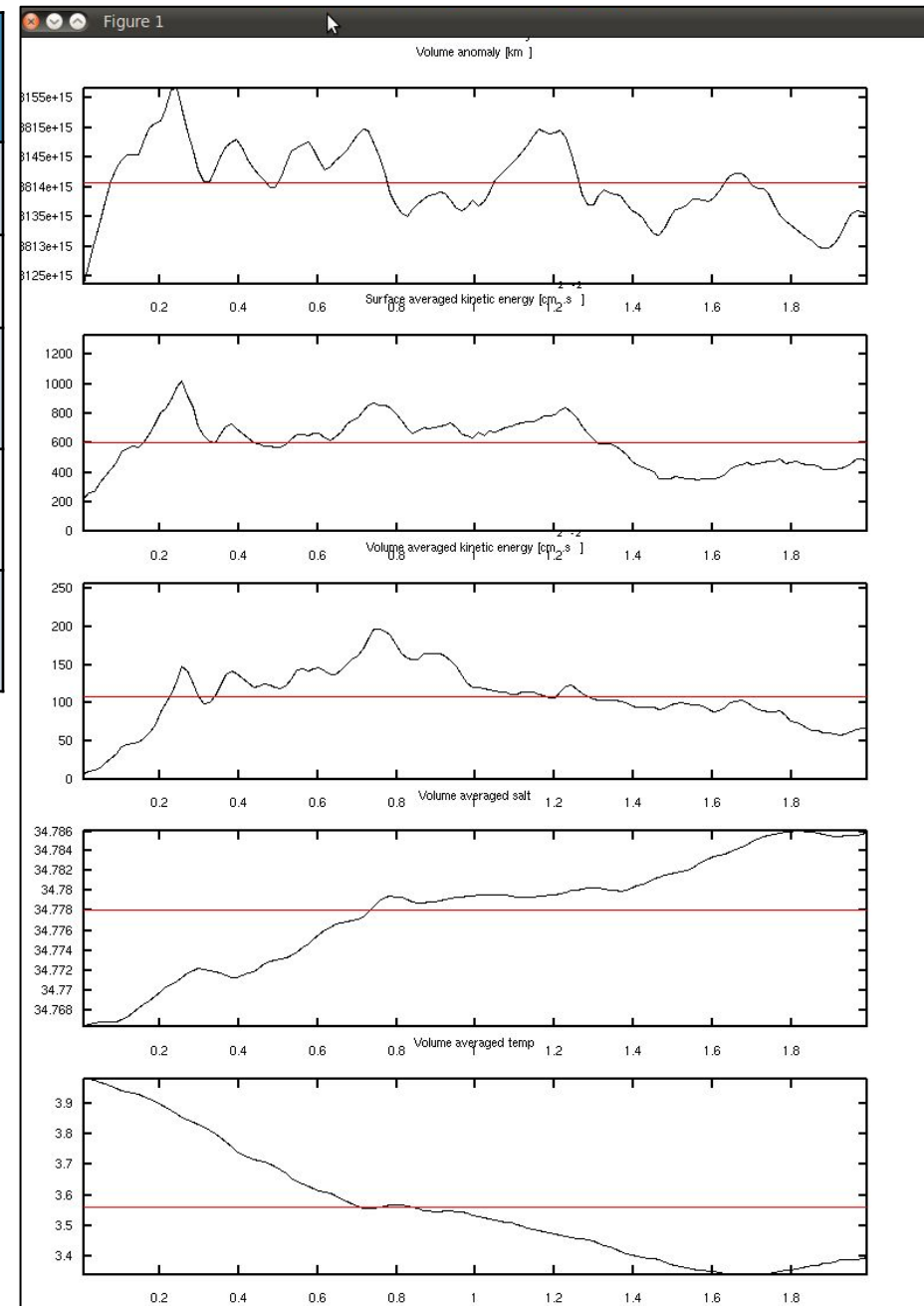
\$matlab

>> croco_gui



Scripts for the simulation analysis in croco_tools/Diagnostics_tools

croco_diags	Get volume and surface averaged quantities
plot_diags	Plot averaged quantities
get_Mmean	Get monthly mean climatology
get_Smean	Get seasonal and annual mean climatology
get_Meddy	Get seasonal and annual variance climatology
get_Seddy	Get seasonal and annual variance climatology



Documentation :

https://croco-ocean.gitlabpages.inria.fr/croco_doc/tutos/tutos.13.tides.html

The tides are imposed at open boundaries using the characteristic open boundary conditions (define **OBC_M2CHARACT** cpp-keys)

- ξ_{tides} , \bar{u}_{tides} , \bar{v}_{tides} : ssh and depth averaged zonal and meridian currents are added at the open boundaries
- ξ_{tides} , \bar{u}_{tides} , \bar{v}_{tides} are computed from the tidal harmonics given by some tidal model, in our case TPXO9 (0.25° resolution, 10 tidal components : M2, N2,S2,K2, K1, O1,P1, Q1, Lm, Mm)
- The global tidal model gives harmonics constants for all the principal tidal waves. These constants permits to compute at every time t, $\xi_{tides}^N(t)$, $\bar{u}_{tides}^N(t)$, and $\bar{v}_{tides}^N(t)$ of the tidal wave component N.

You need :

- Choose the number of tidal wave component you want
- Interpolate on the grid the different harmonic constants
- Possibility to add the generator potential

Adding Tides

matlab >> **make_tides**

```

Start date for nodal correction : 1-Jan-2000
Reading CROCO grid parameters ...
Tidal components : M2 S2 N2 K2 K1 O1 P1 Q1 Mf Mm
Processing tide : 1 of 10
  ssh...
Getting ssh_r for time index 1
Getting ssh_i for time index 1
  u...
Getting u_r for time index 1
Getting u_i for time index 1
  v...
Getting v_r for time index 1
Getting v_i for time index 1
Convert to tidal ellipse parameters...
Process equilibrium tidal potential...
Process tidal loading and self-attraction potential...
Get total tidal potential...
Processing tide : 2 of 10
  ssh...
Getting ssh_r for time index 2
Getting ssh_i for time index 2
  u...
Getting u_r for time index 2
Getting u_i for time index 2
  v...
Getting v_r for time index 2
Getting v_i for time index 2
Convert to tidal ellipse parameters...
Process equilibrium tidal potential...
Process tidal loading and self-attraction potential...
Get total tidal potential...

```

The tidal forcings are added in the croco_frc.nc file.

To define the tides, in cppdefs.h

```

# ifdef TIDES
#  define SSH_TIDES
#  define UV_TIDES
#  define POT_TIDES
#  undef TIDES_MAS
#  ifndef UV_TIDES
#   define OBC_REDUCED_PHYSICS
#  endif
#  define TIDERAMP
# endif

```

Tidal forcing parameter & temporal parameters

```
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%  
%  
% 5 - Parameters for tidal forcing  
%  
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%  
%  
% TPXO file name (TPXO6 or TPXO7)  
%  
tidename=[DATADIR,'TPXO7/TPXO7.nc'];  
%  
% Self-Attraction and Loading GOT99.2 file name  
%  
sal_tides=1;  
salname=[DATADIR,'GOT99.2/GOT99_SAL.nc'];  
%  
% Number of tides component to process  
%  
Ntides=10;  
%  
% Chose order from the rank in the TPXO file :  
% "M2 S2 N2 K2 K1 O1 P1 Q1 Mf Mm"  
% " 1 2 3 4 5 6 7 8 9 10"  
%  
tidalrank=[1 2 3 4 5 6 7 8 9 10];  
%  
% Compare with tidegauge observations  
%  
lon0 = 18.37; % Example:  
lat0 = -33.91; % Cape Town location  
Z0 = 1; % Mean depth of tide gauge  
%  
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
```

For real time runs, with tides forcing , a procedure correct phases and amplitudes (nodal corrections) (Egbert and Erofeeva (2002))

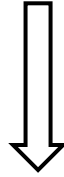
In CROCO , several biogeochemical model : 1D vertical equation (subroutine biology.F)

▪ Nitrogen -based model of increasing complexity:

▪ **NPZD**

▪ **N2PZDD2**

▪ **BIOBUS**



▪ Not only Nitrogen-based (including Iron, etc.): **PISCES**

For initial and lateral boundary condition :

- Seasonal climatology of nitrate provided from WOA (2005 or 2009)
- Seasonal climatology of surface chlorophyll provided from SeasWifs

Choose the BGC model you want (section Applications in cppdefs.h)

```
# define BIOLOGY
...
/* Choice of Biology models */
# ifdef BIOLOGY
# undef PISCES
# undef BIO_NChIPZD
# undef BIO_N2ChIPZD2
# define BIO_BioEBUS
        /* Biology options */
# ifdef PISCES
# undef DIURNAL_INPUT_SRFLX
# define key_pisces
# endif
# ifdef BIO_NChIPZD
# define OXYGEN
# endif
# ifdef BIO_BioEBUS
# define NITROUS_OXIDE
# endif
```

Adding a biological forcings

croco_tools :

- `make_biol` : for climatology file
- `make_bry_npzd` :
 - `_bioebus` :
 - `_pisces` : for boundary files

crocotools_param.m

```
%
% Pisces biogeochemical seasonal climatology
%
woapisces_dir = [DATADIR,'WOAPISCES/']; % only
compatible with woa_dir
%
% Surface chlorophyll seasonal climatology (SeaWifs)
%
chla_dir=[DATADIR,'SeaWifs/'];
%
```

> make_biol

```
Add_no3: creating variables
and attributes for the OA file
write no3time
```

```
Add_no3: creating variables
and attributes for the
Climatology file
```

```
Ext tracers: Roa = 0 km -
default value = NaN
```

```
Ext tracers: horizontal
interpolation of the annual
data
```

```
Ext tracers: horizontal
interpolation of the seasonal
data
```

```
time index: 1 of total: 4
```

```
time index: 2 of total: 4
```

```
time index: 3 of total: 4
```

```
time index: 4 of total: 4
```

...

Documentation :

https://croco-ocean.gitlabpages.inria.fr/croco_doc/tutos/tutos.12.rivers.html

- Crocotools: make_runoff

Section Applications in cppdefs.h

```
/* Point Sources - Rivers */  
# undef PSOURCE  
# define PSOURCE_NCFILE  
# ifdef PSOURCE_NCFILE  
#   define PSOURCE_NCFILE_TS  
# endif
```

croctools_param.m

```
%  
% Runoff monthly seasonal climatology (Dai and Trenberth)  
%  
global_clim_riverdir=[DATADIR,'RUNOFF_DAI/'];  
global_clim_rivename=[global_clim_riverdir,'Dai_Trenberth_runoff_global_clim.nc'];  
%  
% Set times and cycles for the boundary conditions:  
% monthly climatology  
%%  
% Set times and cycles for runoff conditions:  
% monthly climatology  
%  
qbar_time=[15:30:365];  
qbar_cycle=360;  
%  
% Tracer runoff concentration processing flag  
% pource_ts = 1 => Runoff tracers concentration processing is activated.  
%           It needs the climatology file created with make_clim.m  
% psource_ts = 0 => No Runoff tracers concentration processing  
%           It reads analytical values in croco.in  
%           or use default value defined in analytical.F  
%  
psource_ts=0;  
%
```


Adding a river forcing

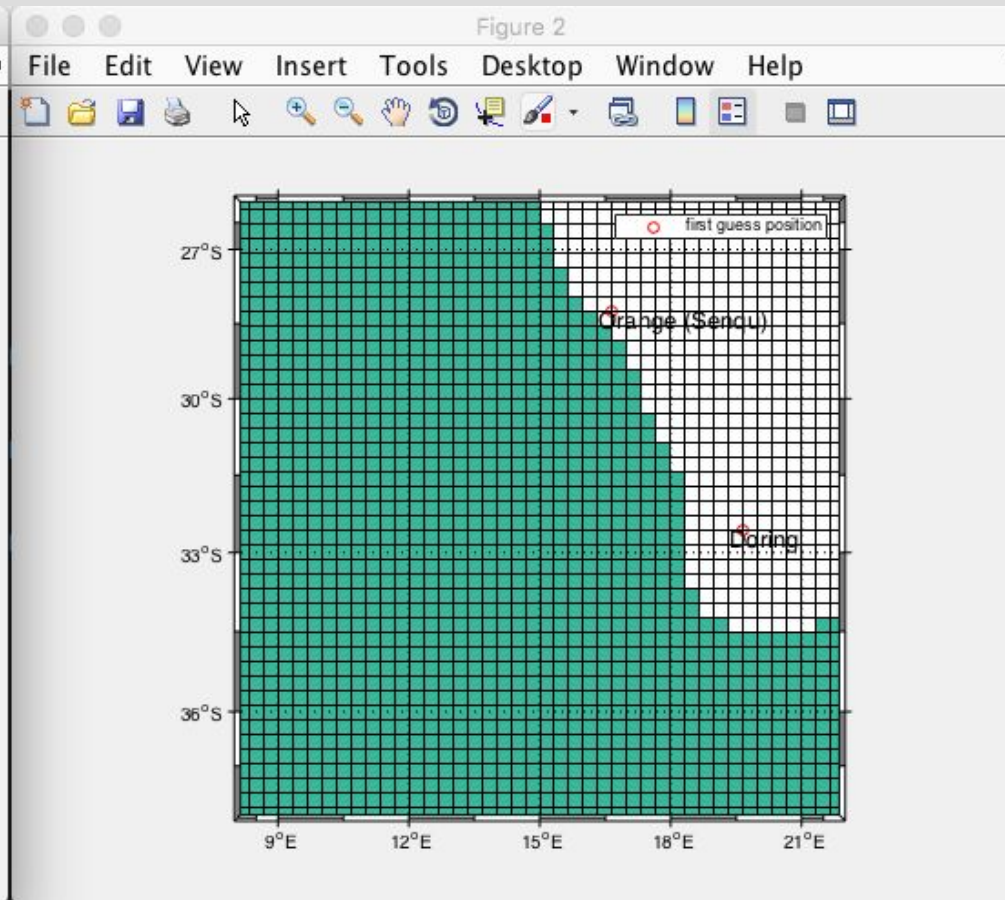
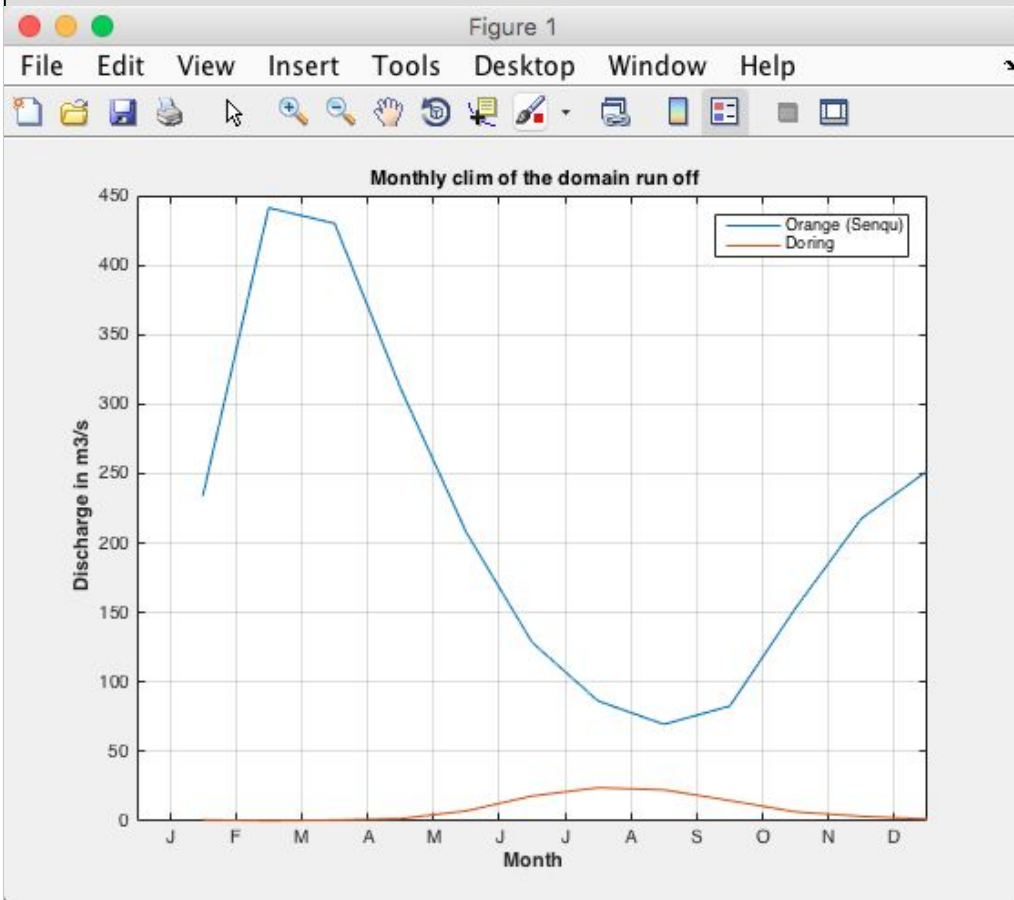
croco_tools : make_runoff.m

- > In make_runoff (line 29)
- mkdir: /Users/gcambon/DATA/for_asi18_sess2/CONFIGS/BENGUELA_LR/CROCO_FILES/: File exists
- Create runoff forcing from Dai and Trenberth's global monthly climatological run-off dataset
- Reading the global monthly climatological run-off dataset...
- There are 2 rivers in the domain :
- Domain contains rivers :
- 1 - Orange (Senqu) flowing in ocean ATL
- 2 - Doring flowing in ocean ATL
- First guess:
- =====
- - Process river #1: Orange (Senqu)
- Position is approximetly J=35 and I=26
- lon src in grid (rho point) ~16.3333
- lat src in grid (rho point) ~-28.5604
- - Process river #2: Doring
- Position is approximetly J=20 and I=35
- lon src in grid (rho point) ~19.3333
- lat src in grid (rho point) ~-32.8514
- ~~Do you want to use river (Yes[1], No[0]) ? Orange (Senqu)~~

Adding a river forcing

croco_tools : make_runoff.m

- > In make_runoff (line 29)
- mkdir: /Users/gcambon/DATA/for_asi18_sess2/CONFIGS/BENGUELA_LR/CROCO_FILES/: File



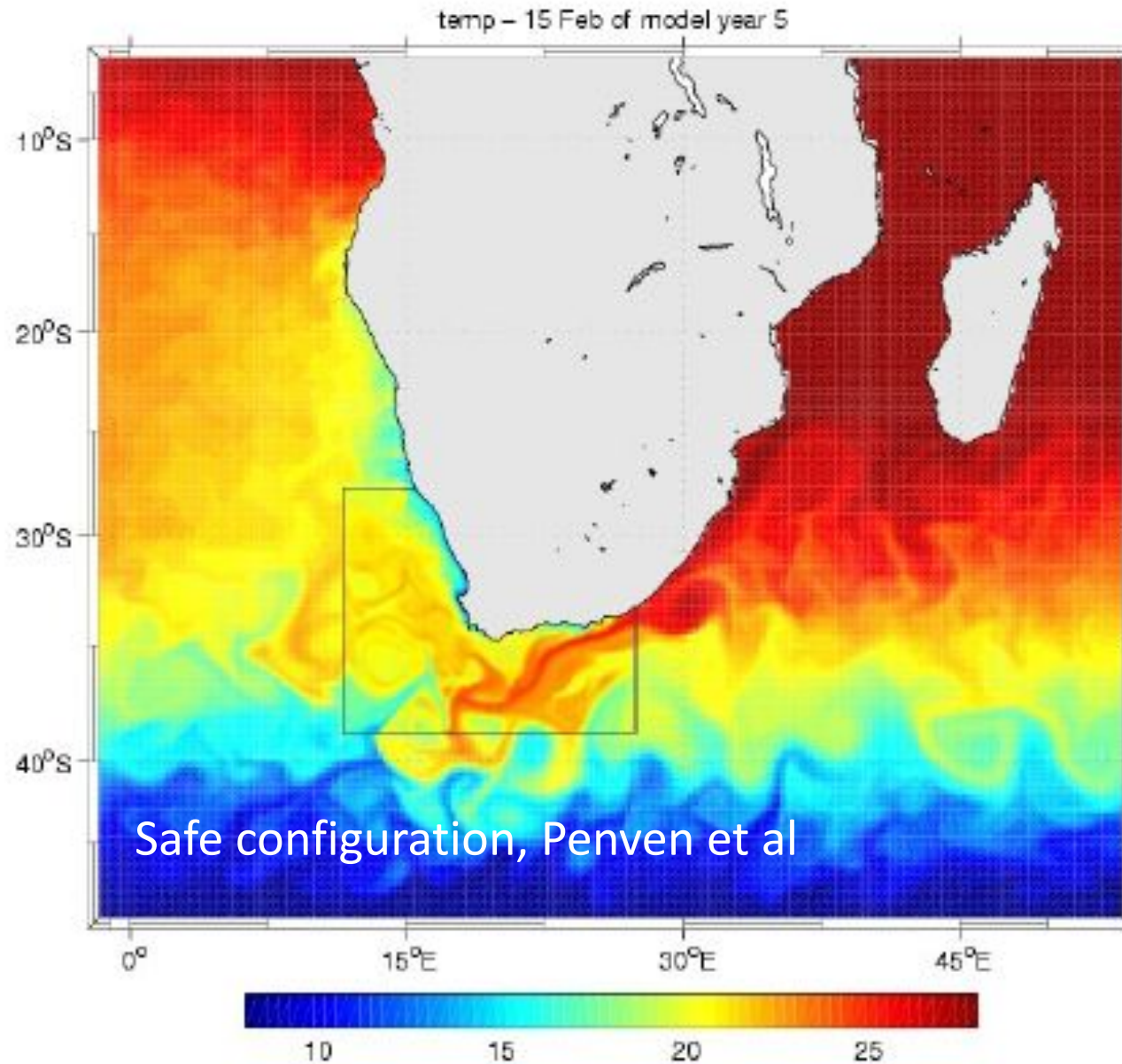
- - Process river #2: Doring
- Position is approximetly J=20 and I=35
- lon src in grid (rho point) ~19.3333
- lat src in grid (rho point) ~-32.8514
- Do you want to use river (Yes[1], No[0]) ? Orange (Senqu)

Adding an AGRIF nest (online nesting)

Documentation :

https://croco-ocean.gitlabpages.inria.fr/croco_doc/tutos/tutos.11.nesting.html

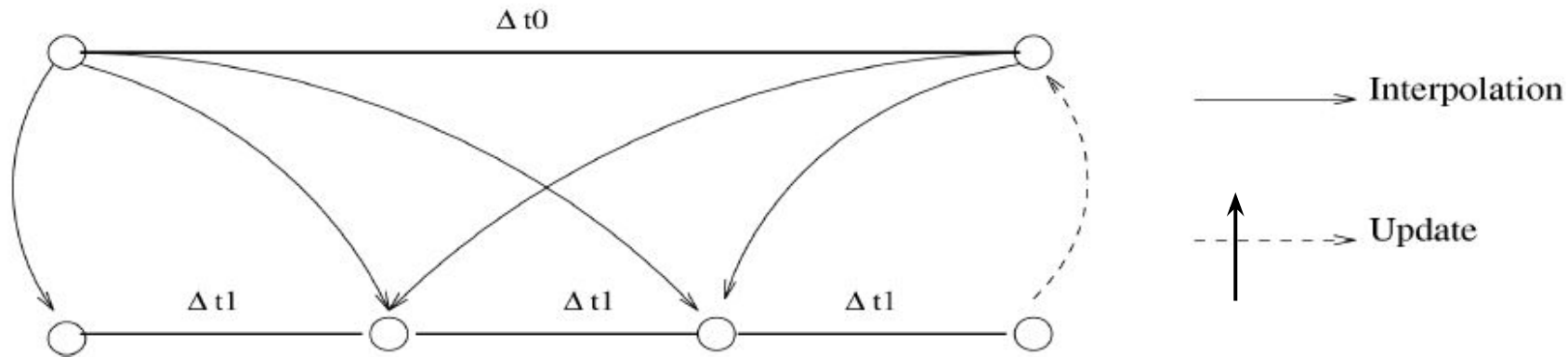
Example :



Nesting capability added to ROMS

- AGRIF package
 - Adaptive Mesh Refinement
 - Manage arbitrary number of fixed grid and embedding level

Temporal coupling between a parent and a child grid for a refinement factor of 3 :



Needs to run an embedded model : Surface forcing and initial conditions datas files.

AGRIF names the different datas files as :

Parent file names : XXX.nc → First child file names : XXX.nc.1
second child file names : XXX.nc.2

...

Adding an AGRIF nest (online nesting)

With matlab

In the benguela test case, for the parent grid file, select in the entrance window of NestGUI and click 'open'

Follow the steps :

1- Tune the child domain

1- Define the child domain :

Size of the child grid

2- Create the child grid file :

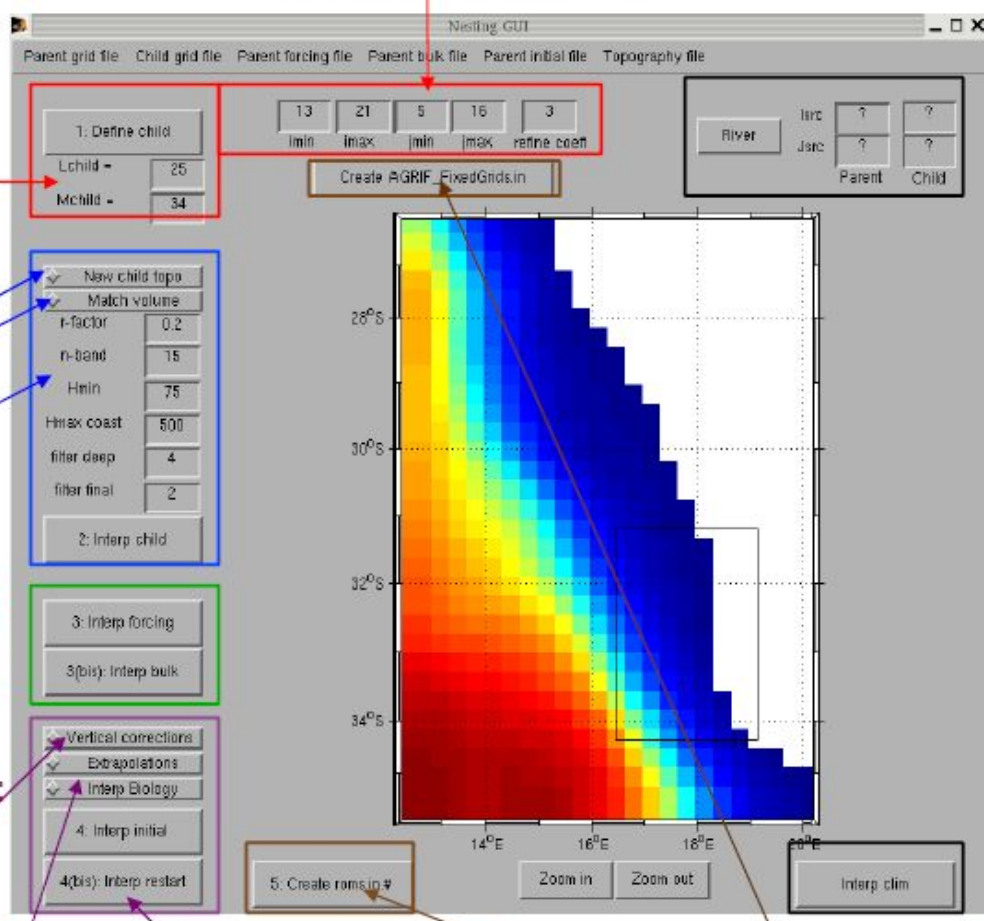
What topography file?
Child grid volume
Parameters to change
-->

3- Create the surface forcing file:

Select
--> roms_frc.nc.*. or roms_blk.nc.*.

4- Create the initial condition file:

Select roms_ini.nc
If different topography
Interpolate parent biological variables
--> roms_ini.nc.*.



Locate river on the coast

Generate boundary condition to test the child model alone

4- Select roms_rst.nc
--> roms_rst.nc.*.

5- Generate roms.in.*.
Create AGRIF_fixedGrids.in

Adding an AGRIF nest (online nesting)

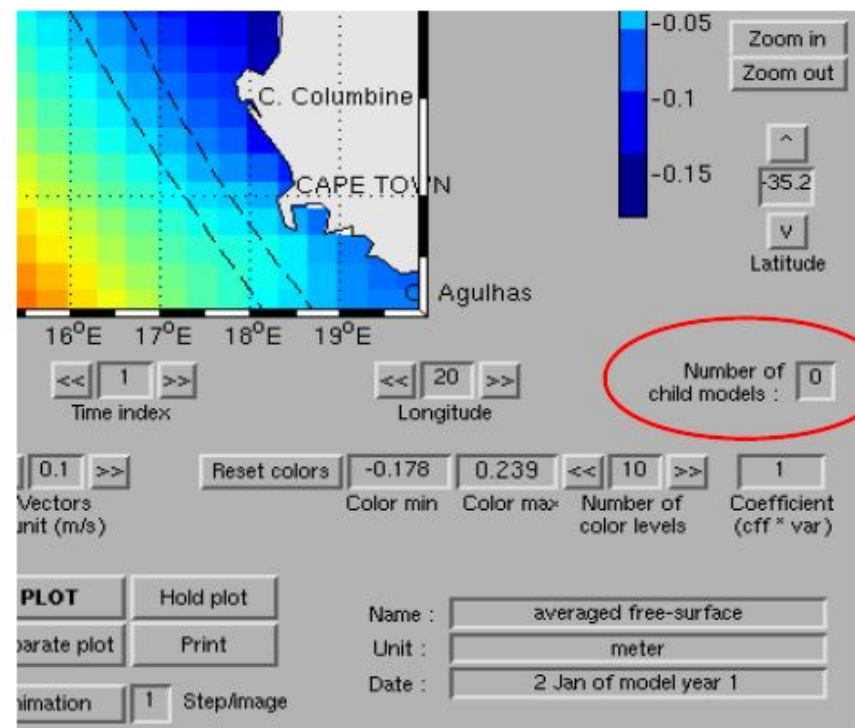
- To run a simulation with nesting, define the CPP keys and compile (./jobcomp)
 - AGRIF
 - AGRIF_2W
- Position of the different grid in AGRIF_FixedGrids.in file

```

1
23 37 12 29 3 3 3 3
0
# number of children per parent
# imin imax jmin jmax spacerefx spacerefy timerefx timerefy
# [all coordinates are relative to each parent grid!]
    
```

- Namelist relative to the different nest level croco.in.1, croco.in.2 etc ...

- Visualization (in Matlab) :
 - >>matlab
 - >>croco_gui



Adding an AGRIF nest (online nesting)

The file Agrif_FixedGrids.in define the position of the nested grid

```

1
23 37 12 29 3 3 3 3
0
# number of children per parent
# imin imax jmin jmax spacerefx spacerefy timerefx timerefy
# [all coordinates are relative to each parent grid!]

```

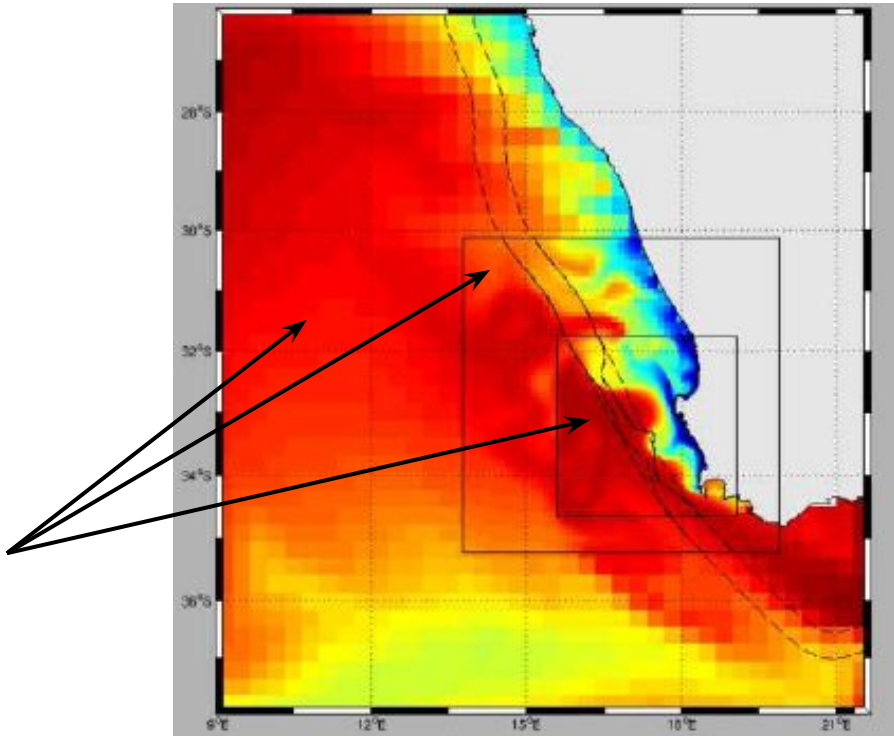
2 grids : #0 and #1
 #1 is embedded in #0

```

1
23 37 12 29 3 3 3 3
1
12 28 15 33 3 3 3 3
0
# number of children per parent
# imin imax jmin jmax spacerefx spacerefy timerefx timerefy
# [all coordinates are relative to each parent grid!]

```

3 grids : #0,#1 and #2
 #1 embedded in #0 ;
 #2 is embedded in the #1



Needs to run an embedded model :

Surface forcing and initial conditions datas files.

For grid #xx :

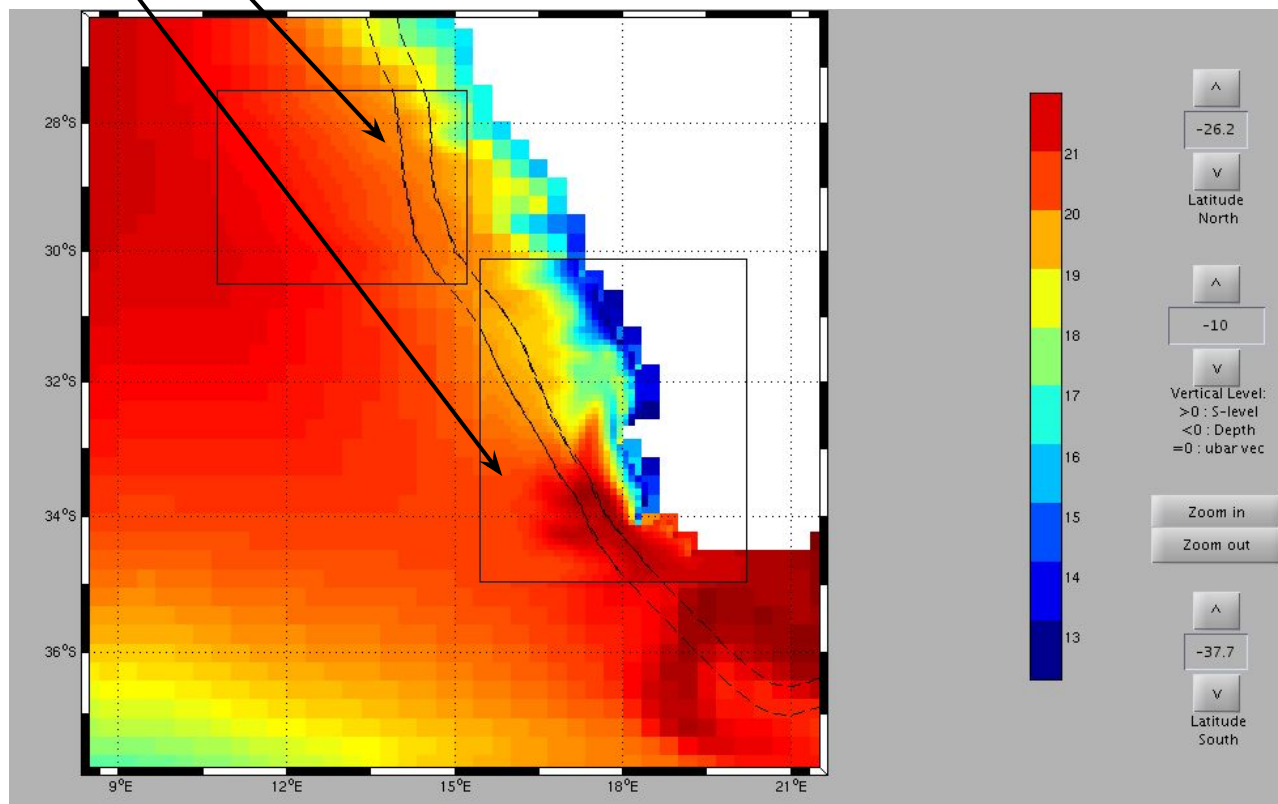
- croco_grd.nc.xx
- croco_blk.nc.xx
- croco.in.xx
- croco_frc.nc.xx
- croco.ini.nc.xx

Adding an AGRIF nest (online nesting)

```

2
23 37 12 29 3 3 3 3
9 22 28 38 3 3 3 3
0
0
#number of children per parent
# ...
    
```

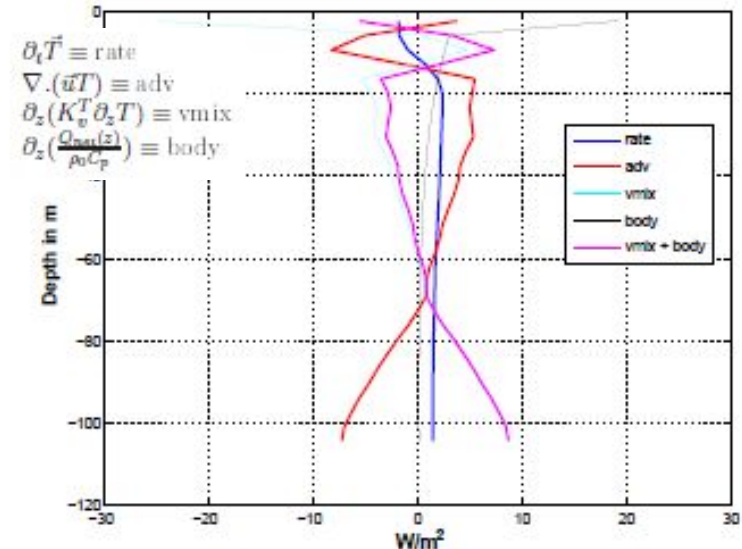
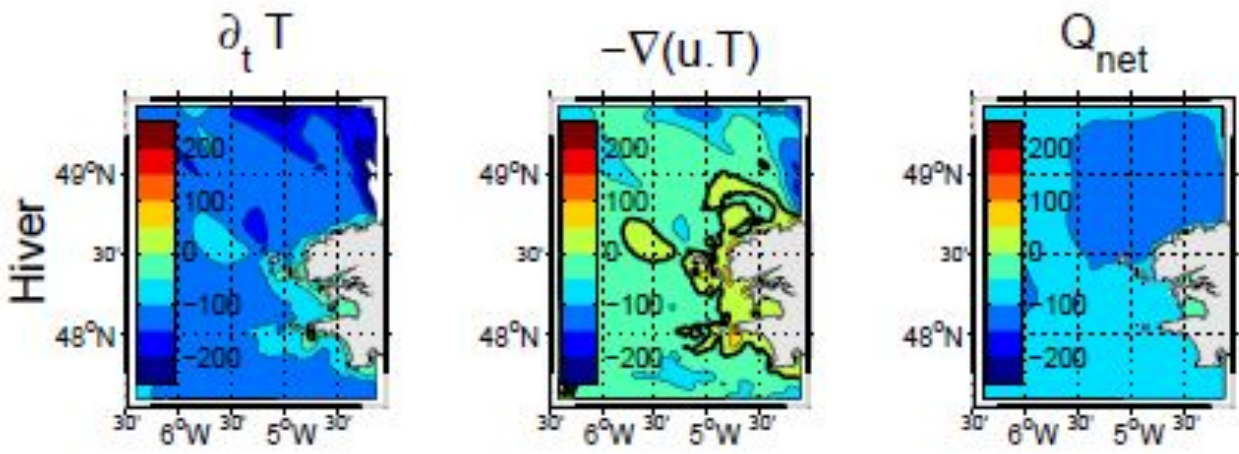
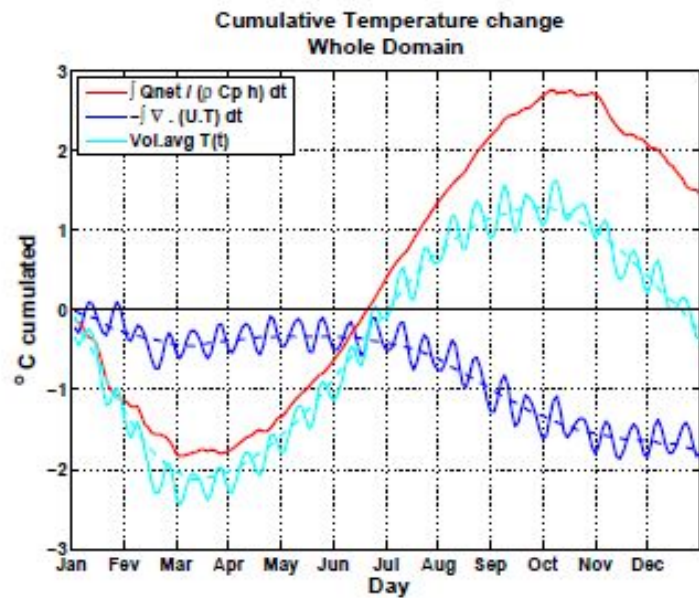
3 grids : #0,#1 and #2
 •#1 embedded in #0 ;
 •#2 is embedded in #0 :
 independent grids



Online Diagnostics : Tracer equations terms

$$\begin{aligned} \partial_t T + \nabla \cdot (\vec{u}T) &= F^T + D^T \\ &= K_h^T \Delta T + \partial_z (K_v^T \partial_z T) + \frac{1}{\rho_0 C_p} \partial_z (Q_{net}(z)) \end{aligned}$$

- $\partial_t \vec{T}$: = Time rate (rate)
- $\nabla \cdot (\vec{u}T)$: = Advection (adv)
- $\partial_z (K_v^T \partial_z T)$: Vert. mixing (vmix)
- $K_h^T \Delta \vec{T}$: Hori. mixing (hmix)
- $\partial_z \left(\frac{Q_{net}(z)}{\rho_0 C_p} \right)$ Solar heating forcing (body)



$\partial_t \vec{T} \equiv \text{rate}$
 $\nabla \cdot (\vec{u}T) \equiv \text{adv}$
 $\partial_z (K_v^T \partial_z T) \equiv \text{vmix}$
 $\partial_z \left(\frac{Q_{net}(z)}{\rho_0 C_p} \right) \equiv \text{body}$

Getting the terms of the equations, stored in netCDF files

- Momentum equation terms : define CPP keys DIAGNOSTICS_TS
- Tracer equation terms : define CPP keys DIAGNOSTICS_UV
- Biological fluxes terms : define CPP keys DIAGNOSTICS_BIO

In croco.in :

```
diagnostics: ldefdia nwrtdia nrpfdia /filename
             T    72    0
             CROCO_FILES/croco_dia.nc
diag_avg: ldefdia_avg ntsdia_avg nwrtdia_avg nrpfdia_avg /filename
          T     1    72    0
          CROCO_FILES/croco_dia_avg.nc
diag3D_history_fields: diag_tracers3D(1:NT)
                     30*T
diag2D_history_fields: diag_tracers2D(1:NT)
                     30*T
diag3D_average_fields: diag_tracers3D_avg(1:NT)
                      30*T
diag2D_average_fields: diag_tracers2D_avg(1:NT)
                      30*T
```

```
diagnosticsM: ldefdiaM nwrtdiaM nrpfdiaM /filename
              T    72    0
              CROCO_FILES/croco_diaM.nc
diagM_avg: ldefdiaM_avg ntsdiaM_avg nwrtdiaM_avg nrpfdiaM_avg
           /filename
           T     1    72    0
           CROCO_FILES/croco_diaM_avg.nc
diagM_history_fields: diag_momentum(1:2)
                    TT
diagM_average_fields: diag_momentum_avg(1:2)
                    TT
diagnostics_bio: ldefdiabio nwrtdiabio nrpfdiabio /filename
                 T    72    0
                 CROCO_FILES/croco_diabio.nc
diagbio_avg: ldefdiabio_avg ntsdiabio_avg nwrtdiabio_avg
            nrpfdiabio_avg /filename
            T     1    72    0
            CROCO_FILES/croco_diabio_avg.nc
```

Some precisions on advection

$$\nabla \cdot (\vec{u}T) \quad u\partial_x T \neq \partial_x(u.T)$$

only

$$\partial_x(uT) + \partial_y(v.T) + \partial_z(w.T) = u\partial_x T + v\partial_y T + w\partial_z T$$

CPPKEYS: **DIAGNOSTICS_TS**

Storage:

- croco_diaM.nc and croco_diaM_avg.nc files.
- Can choose in croco.in the writing frequency and the terms to store
- By default : the different momentum terms with advection terms in “flux forms” :

$$T_{xadv} = \partial_x(u.T)$$

- With cppkey **DIAGNOSTICS_TS_ADV** : compute the “advective” forms :

$$T_{xadv-II} = u.\partial_x(T)$$

- with cppkey **DIAGNOSTICS_TS_MLD** : integration of the different terms over the mixed layer HBL, computed by the KPP model

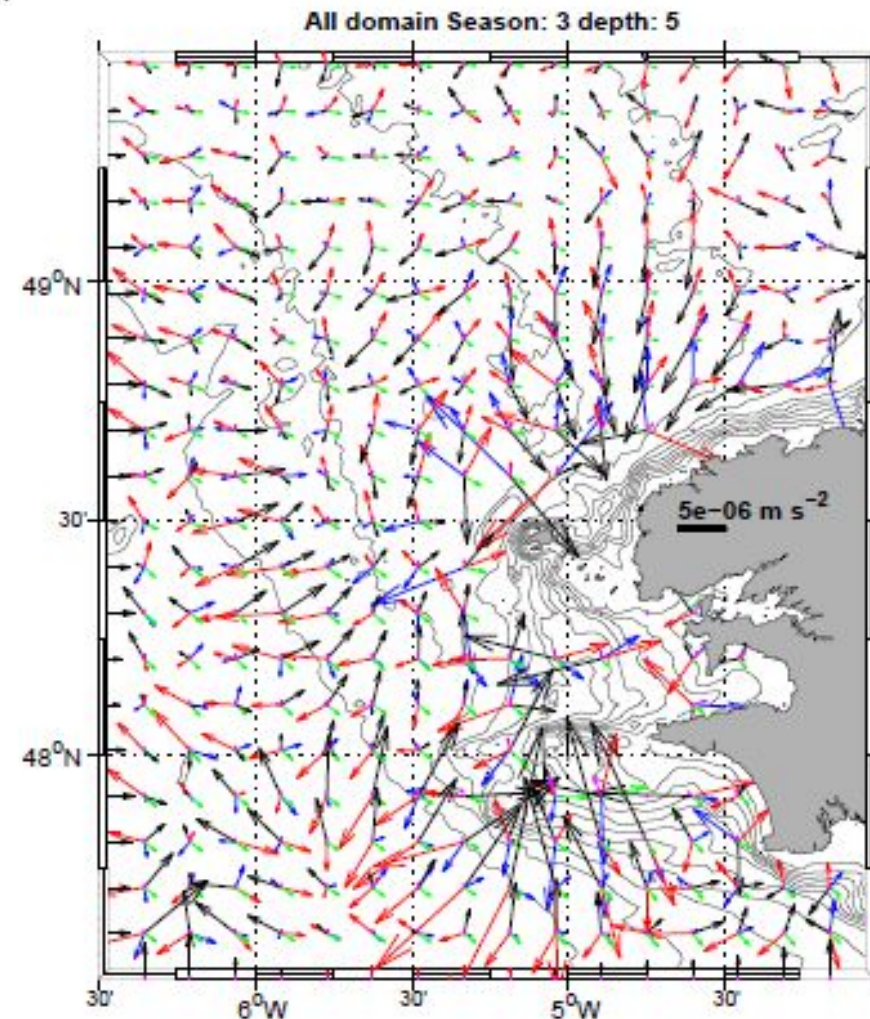
$$\frac{\partial u}{\partial t} + \vec{v} \cdot \nabla u - fv = -\frac{\partial \phi}{\partial x} - \frac{\partial}{\partial z} \left(\overline{u'w'} - \nu \frac{\partial u}{\partial z} \right) + \mathcal{F}_u + \mathcal{D}_u$$

$$\frac{\partial v}{\partial t} + \vec{v} \cdot \nabla v + fu = -\frac{\partial \phi}{\partial y} - \frac{\partial}{\partial z} \left(\overline{v'w'} - \nu \frac{\partial v}{\partial z} \right) + \mathcal{F}_v + \mathcal{D}_v$$

- Rate change term : $\partial_t \vec{u}$
- Coriolis term : $\begin{pmatrix} -fv \\ +fu \end{pmatrix}$
- Advection term : $\vec{u} \cdot \nabla \vec{u}$
- Pressure gradient term : $-\nabla P / \rho_0$
- Vertical mixing term : $\partial_z (K_v \partial_z \vec{u})$
- Horizontal mixing term : $K_H \Delta \vec{u}$

Termes d'accélération en $m \cdot s^{-2}$:

- Advection : $\vec{u} \cdot \nabla \vec{u}$
- Coriolis : $f\vec{u}$
- Mélange vertical : $\partial_z (K_v \partial_z \vec{u})$
- Gradient de pression : $-\partial P / \rho_0$
- Tendance temporelle : $\partial_t \vec{u}$



$$\begin{aligned}
 \vec{u} \cdot \nabla \vec{u} &\equiv \begin{pmatrix} u \partial_x u + v \partial_y u + w \partial_z u \\ u \partial_x v + v \partial_y v + w \partial_z v \end{pmatrix} = \begin{pmatrix} \partial_x (u.u) + \partial_y (u.v) + \partial_z (u.w) \\ \partial_x (u.v) + \partial_y (v.v) + \partial_z (v.w) \end{pmatrix} \\
 &= \begin{pmatrix} u_{xadv} + u_{yadv} + u_{vadv} \\ v_{xadv} + v_{yadv} + v_{vadv} \end{pmatrix}
 \end{aligned}$$

} Formulation
flux

BUT take care :

$$u \partial_x u \neq \partial_x (u.u)$$

only

$$\left(\partial_x (u.u) + \partial_y (u.v) + \partial_z (u.w) \right) = \left(u \partial_x u + v \partial_y u + w \partial_z u \right)$$

CPPKEYS: DIAGNOSTICS_UV