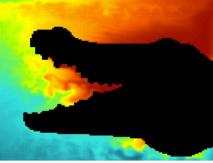
**CROCO – training 2024** 



### **CROCO Architecture**



Camille Mazoyer, IRD, camille.mazoyer@ird.fr

#### How does it work?

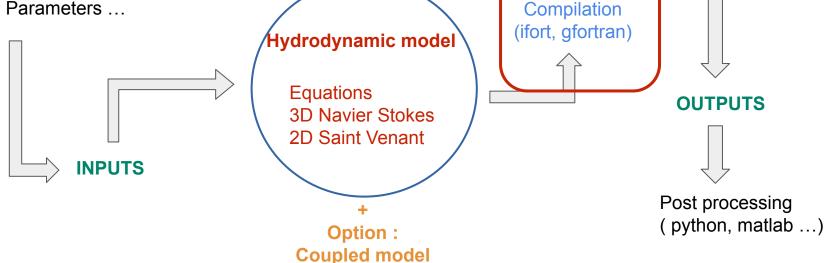


Execution

#### Préprocessing

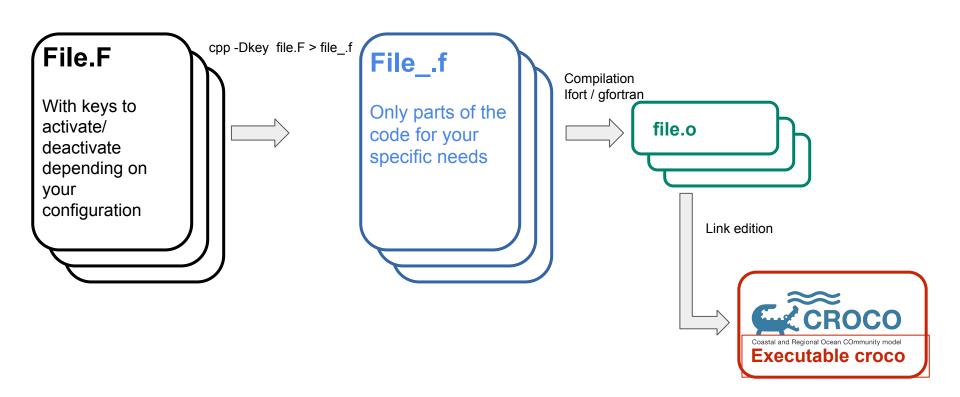
- Bathymetry
- Initial condition
- Boundary conditions
- Forcing (atmosphere, wave, rivers...)

Parameters ...



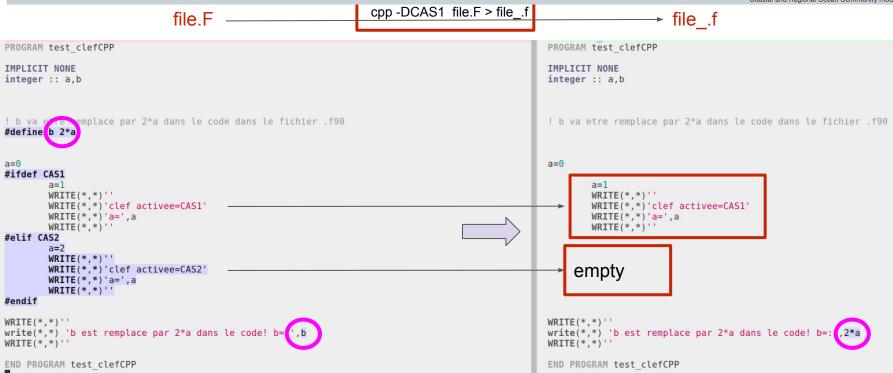
# Compilation in 2 steps





## Compilation with CPP keys





## Compilation with CPP keys



```
file.F file_.f
```

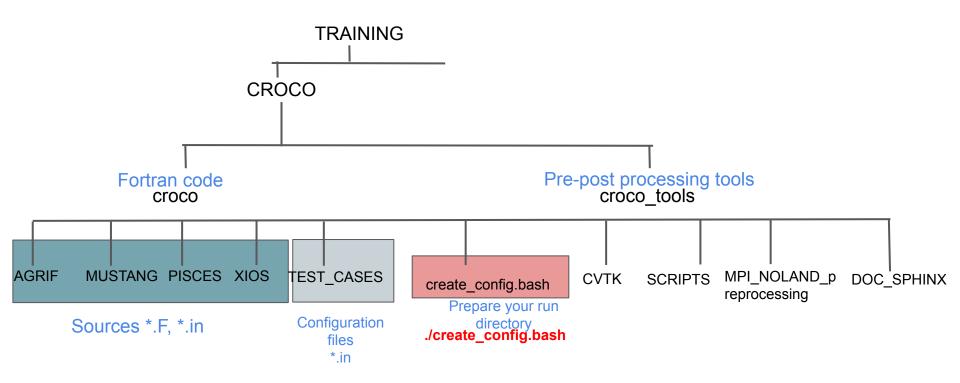
```
#include "cppdefs.h"
        subroutine step()
        implicit none
  #include "param.h"
  #include "scalars.h"
  #include "zoom.h"
23 #include "arid.h"
24 #include "coupling.h"
 #include "ocean3d.h"
6 #include "ocean2d.h"
  #include "mpi cpl.h"
28 #ifdef MUSTANG
29 # include "coupler define MUSTANG.h"
 #endif
  #ifdef AGRIF
        IF (agrif fixed().NE.sortedint(nbtimes)) return
        nbtimes = nbtimes + 1
38 #ifdef SOLVE3D
0 #if defined OA COUPLING || defined OW COUPLING
  !--Get phase of OASIS3
         if ( (iif==-1).and.(oasis time>=0).and.(nbstep3d
            print *, 'oasis time before get = ', oasis t
           call cpl prism get (oasis time)
  #ifdef AGRIF
         else if ((.not.agrif root()).and.(iif==-1).and.
                   (oasis time>=0).and.(nbtimes<=Agrif Max
                   (nbprttime<nbmaxprttime)) then
```

real zeta(GLOBAL\_2D\_ARRAY,4)
real ubar(GLOBAL\_2D\_ARRAY,4)
real vbar(GLOBAL\_2D\_ARRAY,4)
common /ocean\_zeta/zeta
common /ocean\_ubar/ubar
common /ocean vbar/vbar

Each include is paste in file\_.f

#### Architecture of CROCO sources





#### create \_config.bash

cp croco/create\_config.bash .

=> Edit create\_config.bash (e.g. with vi)

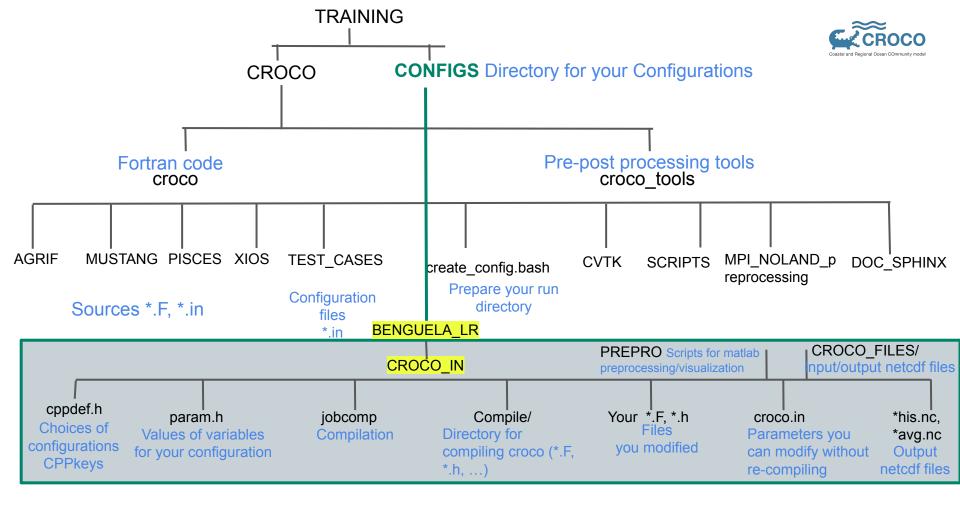
Note: 3 options of configuration architectures available:

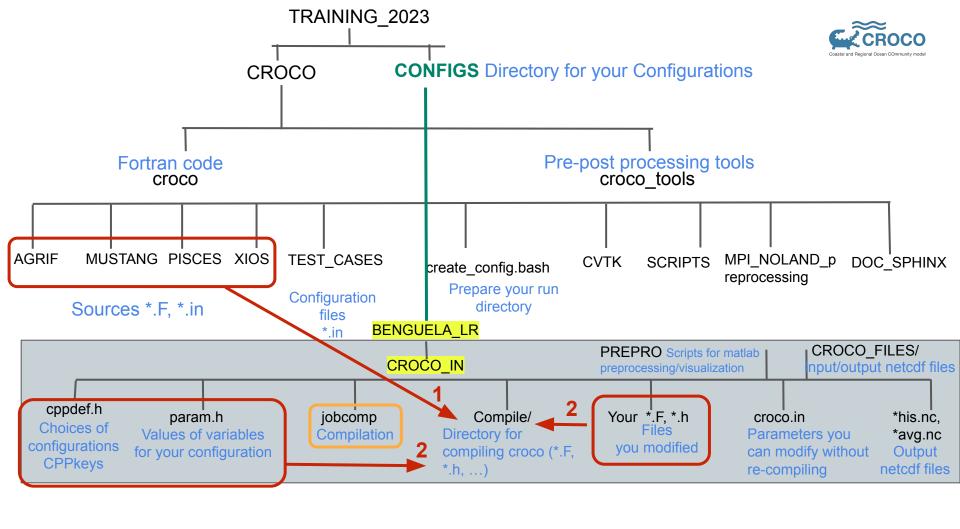
"all-dev": for dev of analytical tests
"all-prod": for production
climatological / interannual
simulations => provides additional
scripts

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

=> choose « all-dev »

```
BEGIN USER MODIFICATIONS
 Machine you are working on
  Known machines: Linux DATARMOR IRENE JEANZAY
MACHINE="DATARMOR"
  croco source directory
CROCO DIR=
          /home/userX/TRAINING/CROCO/croco
  croco tools directory
TOOLS_DIR= /home/userX/TRAINING/CROCO/croco tools
# Configuration name
MY CONFIG NAME=
                BENGUELA LR
 Home and Work configuration directories
                /home/userX/TRAINING/CONFIGS
MY CONFIG HOME=
MY CONFIG WORK=
                /home/userX/TRAINING/CONFIGS
 Options of you
## default option : all—dev for the usual ("all—in") architecture, for forced croco run and/or dev.
#options=( all-dev )
## example for production run architecture
options=
          (all-dev)
## example for production run architecture and coupling with external models :
#options=( all-prod-cpl )
```





### Execution, scripts



#### General architecture of the configuration folder:

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
submitjob.sh