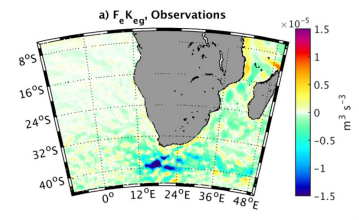
CROCO-WRF Coupling

First copy the croco code:

cp –r /home/student33/lustre/AirSea/croco .

In the CROCO directory, we will create a new configuration

vi create\_config.bash 

🡪 Edit :

MACHINE="WCHPC"

MY\_CONFIG\_NAME=Run\_coupled

MY\_CONFIG\_HOME=/home/studentXX/lustre/AirSea/

MY\_CONFIG\_WORK=/home/studentXX/lustre/AirSea/

options=( all-prod-cpl )

🡪 It will create new folders with (almost) all the files you need to run a coupled simulations

CROCO-WRF Coupling

Now we go to our directory:

cd /home/studentXX/lustre/AirSea/

We have several files to copy

1) Machines files (PBS header, etc):

cp –r /mnt/lustre/users/student33/WCHPC /mnt/lustre/users/studentXX/Run\_Coupled/SCRIPTS\_TOOLBOX/MACHINE/

2) Slight change in a WRF namelist file:

cd /mnt/lustre/users/studentXX/Run\_Coupled/WRF\_IN

Edit namelist.input.base.complete and remove the following line :

**max\_cpldom = <max\_cpldom>,**

3) Now we have to copy the input files (BRY, INI, etc)

• Go to WRF\_FILES : ln –fs /mnt/lustre/users/student33/Data/wrf\* .

• Go to CROCO\_FILES: ln –fs /mnt/lustre/users/student33/Data/croco\*nc .

🡪 Now we have all the files we need to coupled WRF and CROCO over the Benguela Upwelling System ☺ !

CROCO-WRF Coupling

Link the WRF model:

cd /mnt/lustre/users/studentXX

mkdir WRF\_Compil

cd WRF\_Compil

ln –fs /home/apps/chpc/earth/WRF-4.2.1\_cc\_intel2020/WRF/\* .

rm run

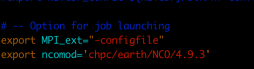
cp –r /home/apps/chpc/earth/WRF-4.2.1\_cc\_intel2020/WRF/run .

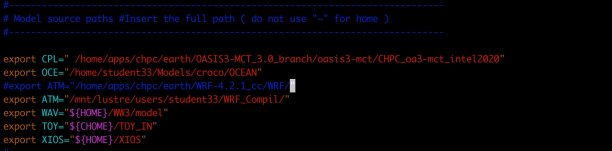
Go to /mnt/lustre/users/studentXX/Run\_Coupled and have a look on the available files We have four main files :

myenv\_mypath.sh -🡪 to configure our environment (eg where is MPI, NetCDF, etc) myjob.sh 🡪 Define the period we want to run

mynamelist.sh 🡪 What kind of simulation we want to do, here a WRF-CROCO simulation submitjob.sh 🡪 To submit the job

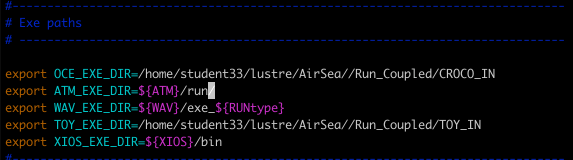
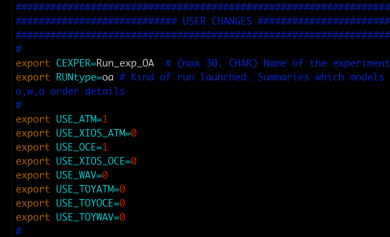
CROCO-WRF Coupling

Edit myenv\_mypath.sh following:



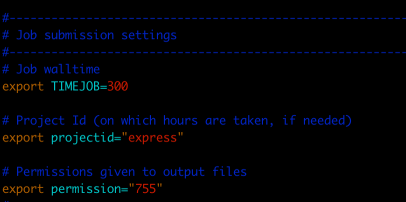
CROCO-WRF Coupling

Edit mynamelist.sh following:



CROCO-WRF Coupling

Edit myjob.sh following:

MPI\_LAUNCH=mpirun 

# Which Computer?

CROCO-WRF Coupling

Compiling CROCO

cd CROCO\_IN

Which files do we have to edit ?

You have to define

**# define OA\_COUPLING and MPI, undef CLIM and define BRY**

🡪 **Jobcomp**

🡪 **mv croco croco.oa**

🡪 **cd ..**

🡪 **./submitjob.sh**

🡪 The simulation should be running in the rundir directory