

# CROCO – training 2023

## TP tests cases



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## TUTORIALS

1. System requirements
2. Download
3. Contents & Architecture
4. Summary of essential steps

### 5. Test Cases

5.1. BASIN

5.2. Set up you own test case

6. Regional: Preparing your configuration

7. Regional: Preprocessing (Matlab)

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## 5. Test Cases

- [5.1. BASIN](#)
- [5.2. Set up you own test case](#)

[← Previous](#)

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Built with [Sphinx](#) using a [theme](#) provided by [Read the Docs](#).

# Create config directory for basin testcase

```
ssh -X userX@croco
```

```
mkdir TRAINING_2023
```

```
cd TRAINING_2023
```

```
cp -r
```

```
/home/COMMONDATA/codes/CROCO .
```

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

```
mkdir CONFIGS
```

```
vim create_config.bash
```

```
./create_config.bash
```

```
# BEGIN USER MODIFICATIONS
# Machine you are working on
# Known machines: Linux DATARMOR IRENE JEANZAY
# -----
MACHINE="DATARMOR"
# croco source directory
# -----
CROCO_DIR= /home/userX/TRAINING_2023/CROCO/croco
# croco_tools directory
# -----
TOOLS_DIR= /home/userX/TRAINING_2023/CROCO/croco_tools
# Configuration name
# -----
MY_CONFIG_NAME= BASIN
# Home and Work configuration directories
# -----
MY_CONFIG_HOME= /home/userX/TRAINING_2023/CONFIGS
MY_CONFIG_WORK= /home/userX/TRAINING_2023/CONFIGS
# Options of your configuration
# -----
## 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-prod )

## example for production run architecture and coupling with external models :
#options=( all-prod-cpl )
```

# TRAINING\_2023

CROCO

**CONFIGS** Directory for your Configurations

Fortran code  
croco

Pre-post processing tools  
croco\_tools

AGRIF

MUSTANG

PISCES

XIOS

TEST\_CASES

create\_config.bash

CVTK

SCRIPTS

MPI\_NOLAND\_p  
reprocessing

DOC\_SPHINX

Sources \*.F, \*.in

Configuration files  
\*.in

Prepare your run directory

**BASIN**

**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 TRAINING\_2023/CONFIGS/BASIN/CROCO\_IN

3. Edit `cppdefs.h` for using BASIN case

```
# define BASIN  
  
# undef REGIONAL
```

You can also explore the CPP options selected for BASIN case.

You can check the BASIN settings in `param.h`.

## Cd TRAINING\_2023/CONFIGS/BASIN

Jobcom should work automatically in Seolane : do not modify it

5. Compile the model:

- By using classical launch command (on individual computers):

```
./jobcomp > jobcomp.log
```

If compilation is successful, you should have a `croco` executable in your directory.

You will also find a `Compile` directory containing the model source files:

- `.F` files: original model source files that have been copied from `$croco/OCEAN`
- `_.f` files: pre-compiled files in which only parts defined by cpp-keys are kept
- `.o` object files

6. **Copy the namelist input file for BASIN case:**

```
Cp ~/TRAINING_2023/CROCO/croco/TEST_CASES/croco.in.Basin .
```

Eventually edit it.

7. **Run the model:**

```
./croco croco.in.Basin
```

If your run is successful you should obtain the following files:

```
basin_rst.nc # restart file  
basin_his.nc # instantaneous output file
```

8. **Have a look at the results:**

```
ncview basin_his.nc
```

## Cd TRAINING\_2023/CONFIGS/BASIN

### 9. Test: some questions:

- What is the size of the grid (see param.h)?
- What are the name of the horizontal directions?
- What is the spatial resolution in both horizontal directions?
- How many vertical levels do you have?
- How are the vertical levels distributed (look for the cpp key `NEW_S_COORD`)?
- What are the initial dynamical conditions (see both cppdefs.h and croco.in)?
- What do the air-sea exchanges look like?



## Cd TRAINING\_2023/CONFIGS/BASIN/CROCO\_IN

10. Re-run this case in parallel on 4 CPUs:

To run in parallel, your first need to edit `cppdefs.h`, `param.h`, and to recompile.

- Edit `cppdefs.h`:

```
# define MPI
```

- Edit `param.h`:

```
#ifdef MPI  
integer NP_XI, NP_ETA, NNODES  
parameter (NP_XI=2, NP_ETA=2, NNODES=NP_XI*NP_ETA)  
parameter (NPP=1)  
parameter (NSUB_X=1, NSUB_E=1)
```

### Note

MPI tiles should be at least 20x20 points.

- Recompile.

# Parallel compilation and execution

## Cd TRAINING\_2023/CONFIGS/BASIN/CROCO\_IN

- Run the model in parallel:
  - By using classical launch command (on individual computers):

```
mpirun -np NPROCS croco croco.in
```

where NPROCS is the number of CPUs you want to allocate. `mpirun -np NPROCS` is a typical mpi command, but it may be adjusted to your MPI compiler and machine settings.

- **OR** by using a batch script (e.g. PBS) to launch the model (in clusters), examples are provided:

```
cp ~/croco/croco_tools/job_croco_mpi.pbs .
```

Edit `job_croco_mpi.pbs` according to your MPI settings in `param.h` and launch the run:

```
qsub job_croco_mpi.pbs
```

### Warning

NPROCS needs to be consistent to what you indicated in `param.h` during compilation

# CPP key BASIN inside the source code

## Cd TRAINING\_2023/CONFIGS/BASIN/CROCO\_IN

grep BASIN Compile/\*

```
OCEAN/ana_grid.F:# if defined BASIN
OCEAN/ana_grid.F:# if defined BASIN
OCEAN/ana_grid.F:# if defined BASIN || defined EQUATOR || defined GRAV_ADJ \
OCEAN/ana_initial.F:#ifdef BASIN
OCEAN/ana_initial.F:#  ifdef BASIN
OCEAN/analytical.F:# ifdef BASIN
OCEAN/cppdefs_dev.h:#if defined BASIN || defined EQUATOR || defined GRAV_ADJ \
OCEAN/cppdefs.h:#undef BASIN /* Basin Example */
OCEAN/cppdefs.h:#elif defined BASIN
OCEAN/param.h:#if defined BASIN
OCEAN/read_inp.F:#if defined BASIN
```

# CPP key BASIN inside the source code

## Cd TRAINING\_2023/CONFIGS/BASIN/CROCO\_IN

grep -i BASIN Compile/\*

Key activation and clefs  
CPP liées  
`cppdef.h`

Parameters related to  
the key  
`param.h`

Grid  
`ana_grid.F`

Initialisation: U, V, ...  
`ana_initial.F`

Analytic functions or parameters: bathymetry,  
coriolis, ...  
`analytical.F`

Cppkey activated (hidden)  
`cppdef_dev.h`

What \*.in to read automatically  
`read_inp.F`

# CPP key BASIN inside the source code

## Cd TRAINING\_2023/CONFIGS/BASIN/CROCO\_IN

grep -i BASIN Compile/\*

Key activation and clefs  
CPP liées  
**cppdef.h**

Parameters related to  
the key  
**param.h**

Grid

**ana\_grid.F**

**ana\_grid.F**

```
!  
# if defined BASIN  
        depth=5000.  
        f0=1.E-4  
        beta=2.E-11  
# elif defined SINGLE_COLUMN  
...  
# if defined BASIN  
        Length_XI =3600.0e+3  
        Length_ETA=2800.0e+3  
# elif defined SINGLE_COLUMN  
...  
# if defined BASIN || defined EQ  
        || defined SO  
        || defined KH  
        || defined AC  
        || defined JE  
do j=JstrR,JendR  
do i=IstrR,IendR  
h(i,j)=depth  
enddo  
enddo
```

# CPP key BASIN inside the source code

## Cd TRAINING\_2023/CONFIGS/BASIN/CROCO\_IN

grep -i BASIN Compile/\*

Grid  
ana\_grid.F

Initialisation: U, V, ...  
ana\_initial.F

Key activation and clefs  
CPP liées  
cppdef.h

Parameters related to  
the key  
param.h

A  
c  
a  
Cp  
cp  
Wh  
rea

```
ana_initial.F
# ifdef TRACERS
# ifdef BASIN
  cff1=(44.690/39.382)**2
  cff2=cff1*(rho0*800./g)*(5.0e-5/((42.689/44.690)**2))
# ifdef TEMPERATURE
  do k=1,N
    do j=JR_RANGE
      do i=IR_RANGE
        t(i,j,k,1,itemp)=cff2*exp(z_r(i,j,k)/800.)
          *(0.6-0.4*tanh(z_r(i,j,k)/800.))
        t(i,j,k,2,itemp)=t(i,j,k,1,itemp)
      enddo
    enddo
  enddo
# endif /* TEMPERATURE */
# elif defined SINGLE_COLUMN && defined WILLIS_DEARDORFF
# ifdef TEMPERATURE
  do k=1,N
    do j=JR_RANGE
```

# CPP key BASIN inside the source code

## Cd TRAINING\_2023/CONFIGS/BASIN/CROCO\_IN

grep -i BASIN Compile/\*

Key activation and clefs  
CPP liées  
**cppdef.h**

Parameters related to  
the key  
**param.h**

```
analytical.F  
  
Stress du vent  
  
# ifdef BASIN  
  cff1=0.0001 * 0.5*(1.+tanh((time-6.*86400.)/(3.*86400.)))  
  cff2=2.*pi/el  
  do j=JstrR,JendR  
    do i=IstrR,IendR  
      sustr(i,j)=-cff1*cos(cff2*yr(i,j))  
    enddo  
  enddo
```

coriolis, ...

**analytical.F**

Cppkey activated (hidden)

**cppdef\_dev.h**

What \*.in to read automatically

**read\_inp.F**